

# *DROSOPHILA*

*Information Service*

# 47

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Prepared at the  
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*Thompson*

DROSOPHILA INFORMATION SERVICE

JULY 1971

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PASADENA, CALIFORNIA: CALIFORNIA INSTITUTE OF TECHNOLOGY  
Division of Biology

Note: The symbol, \*, is used for cross indexing and signifies that the mutant is carried in a stock whose number is shown at the right.

Wild Stocks

- 1 Canton-S
- \* Florida . . . . . 830
- 2 Hikone A-S (strong amylase of Kikkawa)
- 3 Hikone A-W (weak amylase of Kikkawa)
- 4 Lausanne-S
- 5 Oregon-R-C
- 6 Swedish-c
- 7 Urbana-S

Chromosome 1

- \* ac<sub>3</sub><sup>a</sup> . . . . . 174
- 8 ac<sup>w</sup>
- 9 amx/FM3, y<sup>3ld</sup> sc<sup>8</sup> dm B 1
- 10 amx lz<sup>g</sup> v/y f:=
- \* amx . . . . . (see lz<sup>K</sup>)
- 11 Ax
- \* bb<sup>G3</sup> . . . . . 26, 63, etc.
- \* bb<sup>1</sup> . . . . . 133
- \* bb<sup>N</sup> . . . . . 738, 780
- \* bb<sup>poi</sup> . . . . . 167
- \* bb<sup>poi</sup> . . . . . (see bb<sup>G3</sup>)
- 12 B
- 13 B<sub>3</sub>Bx<sup>r</sup> car/y f:=
- \* B<sub>1</sub><sup>i</sup> . . . . . 42
- \* B<sub>1</sub><sup>B</sup> . . . . . 45
- \* BB<sub>36b</sub> . . . . . 43
- \* BB<sub>36b</sub> . . . . . 44
- 14 Bg B/In(1)AM
- 15 bi ct<sup>g</sup>
- 16 bo v
- 17 br
- 18 br<sub>3</sub><sup>w</sup> ec rb t<sup>4</sup>/FM1, y<sup>3ld</sup> sc<sup>8</sup> w<sup>a</sup> lz<sup>s</sup> B
- \* br . . . . . 664
- 19 Bx<sub>2</sub>
- 20 Bx<sub>3</sub>
- 21 Bx<sub>J</sub>
- 22 Bx<sub>r</sub>
- \* Bx<sub>r</sub><sup>49k</sup> . . . . . 13
- \* Bx<sub>r</sub> . . . . . 144
- 23 car
- \* cho<sub>2</sub> . . . . . 101
- \* cho<sub>2</sub> . . . . . 187

- 24 cm
- 25 cm ct<sup>6</sup>
- \* Co<sup>53</sup> . . . . . 184
- 26 cs<sub>6</sub><sup>y w bb</sup>
- \* ct<sup>K</sup> . . . . . 15, 25, etc.
- \* ct<sup>n</sup> . . . . . 175
- 27 ct<sup>n</sup> oc/FM1, y<sup>3ld</sup> sc<sup>8</sup> w<sup>a</sup> lz<sup>s</sup> B
- \* cu-X . . . . . 787
- \* cv . . . . . 102, 103, etc.
- 28 cx
- 29 cx<sup>tg</sup> oc/FM1, y<sup>3ld</sup> sc<sup>8</sup> w<sup>a</sup> lz<sup>s</sup> B
- 30 dm/y f:=
- 31 dor/y f:=
- 32 dor/FM6, y<sup>3ld</sup> sc<sup>8</sup> dm B (nub/+); see also 764
- 33 dow/FM6, y<sup>3ld</sup> sc<sup>8</sup> dm B
- \* dx<sup>st</sup> . . . . . 38, 125
- \* dx<sup>st</sup> . . . . . 664, 665
- 34 dy
- \* e(bx)<sub>2</sub> . . . . . (= en<sup>2</sup>-bx) . . 788
- \* e(bx)<sub>x</sub> . . . . . (= en<sup>x</sup>-bx) . . 678
- \* e(S)<sub>x</sub> . . . . . (= en<sup>x</sup>-S) . . 666
- 35 Eag
- 36 ec
- 37 ec ct<sup>6</sup> s car/FM6, y<sup>3ld</sup> sc<sup>8</sup> dm B
- 38 ec dx
- \* eq . . . . . 102, 737
- 39 Ext/FM6, y<sup>3ld</sup> sc<sup>8</sup> dm B
- 40 f
- 41 f B<sub>3</sub>/y f:=
- 42 f B<sub>3</sub>/y f:=
- 43 f BB<sub>36b</sub>/y f:=
- 44 f BB<sub>36b</sub><sup>i</sup>/y f:=
- 45 f B<sub>1</sub><sup>i</sup>/y f:=
- 46 f fu/y f:=
- \* f<sub>3</sub> . . . . . 167
- \* f<sub>36a</sub> . . . . . 157
- 47 f B15
- \* f B27 . . . . . 774
- \* f . . . . . 775
- 48 fa<sub>n</sub>
- \* fa<sub>n</sub> . . . . . 782
- \* flp . . . . . (see flw)
- 49 flw
- 50 fo

51	fx/y f:=	
*	fu . . . . .	46
*	fw <sub>34e</sub> . . . . .	146, 170
*	fw <sub>2</sub> . . . . .	194
52	g <sub>2</sub> . . . . .	
53	g <sub>2</sub> pl/FM3, y <sup>3ld</sup> sc <sup>8</sup> dm B l	
54	g <sub>4</sub> ty/y f:=	
*	g <sub>2</sub> . . . . .	667, 733
55	gg <sub>3</sub> /FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B	
56	gg <sub>a</sub> . . . . .	
57	gt <sub>w</sub> . . . . .	
58	Hk <sub>2</sub> . . . . .	
*	Hw <sub>49c</sub> . . . . .	114, 733, etc.
59	Hw <sub>3</sub> /FM1, y <sup>3ld</sup> sc <sup>8</sup> w <sup>a</sup> lz <sup>s</sup> B	
60	if . . . . .	
61	kz . . . . .	
*	l(1)7 . . . . .	(see dor <sup>1</sup> )
62	l(1)Jl sc <sup>Jl</sup> /l(1)Jl sc <sup>Jl</sup> /Dp(l;f)24	
63	lh B car bb/y f:=	
64	lz/FM3, y <sup>3ld</sup> sc <sup>8</sup> dm B l	
65	lz <sub>3</sub> /y f:=	
66	lz <sub>34</sub> /y f:=	
67	lz <sub>36</sub> /y f:=	
68	lz <sub>37</sub> /y f:=	
69	lz <sub>48f</sub> . . . . .	
*	lz <sub>50e</sub> /y f:=	
*	lz <sub>BS</sub> . . . . .	766
70	lz <sub>D</sub> lz <sub>46</sub> lz <sub>g</sub> ras v/y f:=	
*	lz <sub>g</sub> . . . . .	667
*	lz <sub>K</sub> . . . . .	10, 180, etc.
71	lz <sub>s</sub> . . . . .	
*	lz <sub>y4</sub> . . . . .	18, 27, etc.
*	lz <sub>y4</sub> . . . . .	120, 766
72	m <sub>2</sub> . . . . .	
*	m <sub>D</sub> . . . . .	667, 733
73	m/FM3, y <sup>3ld</sup> sc <sup>8</sup> dm B l	
74	M(1)n/FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B	
75	M(1)o <sub>f</sub> /FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B	
76	M(1)o <sub>Sp</sub> /FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B	
*	M(1)Sp . . . . .	(see M(1)o <sup>Sp</sup> )
77	mal/y f:=	
*	mal <sub>bz</sub> . . . . .	183
78	na/y f:=	
79	ny f/FM1, y <sup>3ld</sup> sc <sup>8</sup> w <sup>a</sup> lz <sup>s</sup> B (ri)	
80	oc ptg/In(1)ClB	
*	od <sub>o</sub> . . . . .	(see os <sup>o</sup> )
81	os <sub>s</sub> . . . . .	
82	os <sub>3</sub> . . . . .	
83	pa sn/FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B	
84	peb v . . . . .	
*	pl . . . . .	53
*	pn . . . . .	109, 176, etc.
85	pn <sub>2</sub> . . . . .	
*	pn <sub>3</sub> . . . . .	105
*	ptg <sub>2</sub> . . . . .	679
86	ptg <sub>3</sub> . . . . .	
*	ptg <sub>4</sub> . . . . .	80
*	ptg <sub>9</sub> . . . . .	141
87	r <sub>39k</sub> /y f:=	
88	r <sub>39k</sub> f B/In(1)AM	
89	ras <sub>2</sub> dy . . . . .	
90	ras <sub>3</sub> . . . . .	
91	ras <sub>4</sub> m . . . . .	
*	ras <sub>4</sub> . . . . .	70
*	ras <sub>v</sub> . . . . .	845
92	rb . . . . .	
93	rb cx . . . . .	
94	rg <sub>2</sub> /FM1, y <sup>3ld</sup> sc <sup>8</sup> w <sup>a</sup> lz <sup>s</sup> B	
95	rst/FM1, y <sup>3ld</sup> sc <sup>8</sup> w <sup>a</sup> lz <sup>s</sup> B	
96	rux/FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B	
97	rux <sub>2</sub> . . . . .	
98	s . . . . .	
99	sbr . . . . .	
100	sc . . . . .	
101	sc cho t . . . . .	
102	sc cv v eq (sc reverted)	
103	sc cv v f . . . . .	
104	sc ec <sub>3</sub> cv <sub>2rv</sub> v <sub>2g</sub> f/FM3, y <sup>3ld</sup> sc <sup>8</sup> dm B l	
105	sc pn <sub>g</sub> Bx <sub>2</sub> . . . . .	(g <sup>6</sup> rev., sc rev.)
105b	sc w <sup>bl</sup> ec cv . . . . .	
106	sc z ec <sub>62b</sub> . . . . .	
107	sc z sw <sub>Y7G2</sub> (Ives <sub>6</sub> ) . . . . .	
108	sc <sub>2</sub> z w <sub>ec</sub> ct . . . . .	
109	sc <sub>3B</sub> pn/y f:=	
110	sc <sub>3-1</sub> . . . . .	
111	sc <sub>4</sub> w/y f:=	
*	sc <sub>5</sub> . . . . .	794
112	sc <sub>6</sub> . . . . .	
113	sc <sub>7</sub> w <sup>a</sup> . . . . .	
*	sc <sub>8</sub> . . . . .	722, 796, etc.
*	sc <sub>9</sub> . . . . .	784, 801, etc.
*	sc <sub>10</sub> . . . . .	805
*	sc <sub>10-1</sub> . . . . .	(see ac <sup>3</sup> )
114	sc <sub>19</sub> /y Hw . . . . .	
*	sc <sub>D2</sub> . . . . .	838
*	sc <sub>J1</sub> . . . . .	182
*	sc <sub>J4</sub> . . . . .	62
*	sc <sub>S1</sub> . . . . .	769
*	sc <sub>S2</sub> . . . . .	178
*	sc <sub>260-14</sub> . . . . .	837
*	sc <sub>260-15</sub> . . . . .	806
*	sc <sub>260-22</sub> . . . . .	847
*	sc . . . . .	807

* sc <sup>z</sup> . . . . . 846	* vb <sup>2</sup> . . . . . 135
115 scp <sup>t</sup> <sub>8d14</sub>	149 vs
116 sd <sup>5</sup> /y f:=	150 w
117 Sh <sup>2</sup>	151 w m f <sub>3</sub>
118 shf <sup>2</sup>	152 w <sub>11E4</sub> m
* sl <sup>2</sup> . . . . . (in ClB, ClB <sup>36d</sup> )	* w <sub>17G2</sub> . . . . . 199
* sl . . . . . 723	* w <sub>a</sub> . . . . . 108
* sn <sup>2</sup> . . . . . 713	153 w <sub>a2</sub>
* sn <sup>3</sup> . . . . . 161	154 w <sub>a3</sub>
119 sn <sup>3</sup> <sub>3</sub>	155 w <sub>a4</sub>
120 sn <sup>4</sup> lz <sup>y4</sup> v/y f:=	156 w <sub>bf</sub> <sup>5</sup>
121 sn <sup>34e</sup>	157 w <sub>bf2</sub> f
122 sn <sup>36a</sup>	158 w <sub>bf3</sub>
123 sn <sup>36a</sup> /y f:=	* w <sub>Bwx</sub> . . . . . 697
* sp-w . . . . . (see w <sup>sp</sup> )	159 w <sub>ch</sub>
124 spl	160 w <sub>co</sub> wy <sub>2</sub>
* sta . . . . . (see T(1;3)sta)	161 w <sub>col</sub> sn
* su( . . . . . 677	162 w <sub>e</sub>
125 su(dx) dx	163 w <sub>e2</sub>
* su(f) <sup>2</sup> . . . (= su <sup>w</sup> -f) . . 145	164 w <sub>ec3</sub>
126 su(s) <sup>2</sup> v <sub>a</sub> (bw)	165 w <sub>h</sub>
127 su(s) <sup>2</sup> w <sub>a</sub> cv t	166 w <sub>i</sub> f <sub>3</sub> bb <sup>N</sup>
128 su(s) <sup>3</sup> cv v f/y f:=	167 w <sub>sat</sub>
129 su(s) <sup>S</sup> v/FMA3, y <sub>a</sub> (bw)	168 w <sub>sp</sub>
* su(w <sub>S2</sub> ) . . (= su-w <sub>a</sub> ) . . 700, 708, etc.	169 w <sub>t</sub>
* su <sup>-v-pr</sup> . . . . . (see su(s))	170 w <sub>u</sub> fw
130 svr	171 w <sub>u</sub>
131 svr <sub>poi</sub>	172 wy <sub>2</sub>
132 svr <sub>poi-dish</sub> bb <sup>G3</sup>	* wy . . . . . 192
133 svr <sub>poi-dish</sub> bb <sup>G3</sup>	173 y
134 sw	174 y ac v
135 sx vb <sup>2</sup> os <sup>s</sup> /FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B	175 y ct <sup>K</sup> (bw)
* sy . . . . . (see os <sup>s</sup> )	176 y pn
136 t <sub>2</sub>	177 y pn w cm ct <sup>6</sup> sn <sup>3</sup> oc ras <sup>2</sup> v dy g <sup>2</sup> f os <sup>o</sup>
137 t <sub>3</sub> v f	car sw/FM7b, y <sup>3ld</sup> w <sub>a</sub> lz <sup>s</sup> B
138 t <sub>4</sub>	178 y pn w cm ct <sup>6</sup> sn <sup>3</sup> oc ras <sup>2</sup> v dy g <sup>2</sup> f os <sup>o</sup>
* t <sub>5</sub> . . . . . 18	car sw/In(1)sc <sup>S1</sup> , In(1)dl-49, y v B
139 t <sub>5</sub> v r	179 y sc
* tuh-l . . . (= tu-h) . . 673	180 y sc <sub>5</sub> lz <sup>g</sup> v f/y f:=
140 tw/FM1, y <sup>3ld</sup> sc <sup>8</sup> w <sub>a</sub> lz <sup>s</sup> B	181 y sc <sub>D2</sub>
* ty . . . . . 54	182 y sc
* tyl . . . (= ty-l) . . 779, 780	183 y v f mal <sup>bz</sup>
141 un <sub>4</sub> Bx <sup>2</sup> /In(1)AM, ptg	184 y w Co/y f:=
142 un	185 y <sub>2</sub> w spl
143 v	186 y <sub>2</sub>
144 v f Bx <sup>r49k</sup> car/y f:=	187 y <sub>2</sub> cho <sup>2</sup>
145 v <sub>2</sub> f su(f)	188 y <sub>2</sub> cv v f <sub>a</sub>
146 v <sub>36f</sub> fw	189 y <sub>2</sub> sc w <sub>a</sub> ec
147 v <sub>0f</sub>	190 y <sub>2</sub> w <sub>a</sub>
* v . . . . . 781	191 y <sub>2</sub> w <sub>2</sub> w <sub>2</sub> (g <sup>2</sup> partly reverted)
148 vb	192 y wy <sub>g</sub> g <sup>2</sup> (g <sup>2</sup> partly reverted)

193 y<sup>2S</sup>  
 194 y<sup>2S</sup> fw<sup>34e</sup>  
 195 y<sup>3d</sup>/y f:=  
 \* y<sup>3p</sup> . . . . . 812  
 \* y<sup>4</sup> . . . . . 814  
 \* y<sup>3fd</sup> . . . . . 9, 18, etc.  
 \* y<sup>34c</sup> . . . . .  
 196 y<sup>59b</sup>  
 \* y<sup>td</sup> . . . . . 709  
 197 y<sup>v2</sup>  
 198 y<sup>11E4</sup>  
 199 z w

## Chromosome 2

200 a px or  
 201 a px sp  
 202 ab<sup>2</sup>  
 203 ab<sup>2</sup>/T(Y;2)E  
 204 ab<sup>2</sup> ix<sup>2</sup> bw sp<sup>2</sup>/In(2L+2R)Cy, Cy dp<sup>1vI</sup> Bl  
 L sp  
 \* abb . . . . . 403  
 205 abr/In(2L+2R)Cy, Cy hk<sup>2</sup>  
 206 abr/SM5, al<sup>2</sup> Cy lt sp<sup>2</sup>  
 207 ad  
 208 al  
 209 al b c sp<sup>2</sup>  
 210 al dp b bw l(2)ax/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 211 al dp b pr ap<sup>blt</sup> bw/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 212 al dp b pr Bl c px sp/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 213 al dp b pr Bl c px sp/In(2LR)O, dp<sup>1vI</sup> Cy  
 pr cn  
 214 al dp b pr c px sp  
 215 al dp b pr Hx  
 216 al<sup>S</sup> ast ho/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 \* al<sup>2</sup> . . . . . 210, 211, etc.  
 \* alpha-1 . . . . . (see tyr-1)  
 217 Alu  
 218 an/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 219 an/SM1, al<sup>2</sup> Cy sp  
 220 ang  
 221 ant(ro)  
 222 ap<sup>blt</sup>/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 223 ap  
 224 arch chl/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 225 ast<sup>3</sup> ho cl  
 \* ast<sup>4</sup> . . . . . 815  
 226 ast<sup>x</sup> dp cl  
 \* ast . . . . . 300  
 227 Ata . . . . . 868  
 228 b  
 229 b tyr-1

230 b cn beta  
 231 b el rd<sup>s</sup> pr cn  
 232 b Go/In(2LR)Gla  
 233 b Go/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 234 b gp  
 235 b j  
 236 b l(2)Bld pr c px sp/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 237 b lt wx<sup>wxt</sup> bw  
 238 b pr tk/T(Y;2)G  
 239 b sf  
 240 b vg  
 \* ba . . . . . 251  
 241 Bl/In(2L+2R)Cy, Cy bw<sup>45a</sup> sp<sup>2</sup> or<sup>45a</sup>  
 242 Bl/T(2;3)dp  
 243 Bl L<sup>48</sup>/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 244 Bl stw<sup>2</sup> ap<sup>blt</sup> tuf sp/SM5, al<sup>2</sup> Cy lt<sup>v</sup>  
 sp  
 245 Bla/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 246 blo  
 \* blt . . . . . (see ap<sup>blt</sup>)  
 247 bri  
 \* bs<sup>2</sup> . . . . . 380  
 248 bs<sup>3</sup>  
 \* bs . . . . . 328  
 249 bur fs(2)El/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 250 bw  
 251 bw ba  
 252 bw tu  
 253 bw<sup>2b</sup>  
 \* bw<sup>4</sup> . . . . . 686  
 \* bw<sup>45a</sup> . . . . . 241  
 254 bw<sup>D</sup>  
 \* bw<sup>v1</sup> . . . . . 328, 353, etc.  
 \* bw<sup>v32g</sup> . . . . . 352, 739  
 \* bw<sup>v34k</sup> . . . . . 342  
 255 c  
 256 c wt px  
 257 cg c/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 258 cg c/In(2LR)U  
 259 ch  
 260 chl  
 261 chl en/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 262 chl l(2)bw bw<sup>2b</sup> mr/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 263 chy  
 264 ck/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 265 cl<sup>2</sup>  
 266 cl<sup>2</sup>/T(Y;2)E  
 267 cn<sup>2</sup>  
 \* cn (in all stocks containing In(2R)Cy)  
 268 cn bw  
 269 cn en/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>

- 270  $cn_3^1(@)crc/SM5, al^2 Cy lt^v sp^2$   
 271  $cn_3^2 T(Y;2)C$   
 272  $cn_{35K}$   
 \*  $cq$  . . . . . (see  $rk^4$ )  
 273  $cru/In(2L+2R)Cy, Cy (w^e)$   
 274  $Cy Bl bw/SM1, al^2 sp^2$  (no  $Cy$ )  
 275  $d/SM5, al^2 Cy lt^v sp^2$   
 276  $d b/SM5, al^2 Cy lt^v sp^2$   
 277  $da/SM1, al^2 Cy sp^2$   
 277b  $da pr cn/SM5, al^2 Cy lt^v sp^2$   
 278  $dil^2 hv bw sp/SM5, al^2 Cy lt^v sp^2$   
 279  $dke c$   
 280  $dp$   
 281  $dp_2 cn bw$   
 \*  $dp_D$  . . . . . (see  $dp^{lv2}$ )  
 \*  $dp_{lv}$  . . . . . 876  
 282  $dp_{lv2} b/SM5, al^2 Cy lt^v sp^2$   
 \*  $dp_{lvI}$  . . . . . 292, 293, etc.  
 \*  $dp_{Nov}$  . . . . . 204  
 \*  $dp_o$  . . . . . (see  $dp^{ovN}$ )  
 283  $dp_{o2}$   
 284  $dp_{olvR}/SM5, al^2 Cy lt^v sp^2$   
 285  $dp_{ovN}$   
 286  $dp_{Rf}$   
 \*  $dp_{Th}$  . . . . . (see  $dp_{olvR}$ )  
 \*  $dp_{tx}$  . . . . . (see  $dp_{lv}$ )  
 \*  $dp_v$  . . . . . (see  $dp_{lv}$ )  
 \*  $dp_{v2}$  . . . . . 690  
 287  $dp_{vl}$   
 \*  $dp_{vM}$  . . . . . (see  $dp^{vM}$ )  
 288  $dp_{vM}/SM5, al^2 Cy lt^v sp^2$   
 289  $ds_{rv} dp$   
 290  $ds_{rv} ft dp^{v2} 1(2)M b pr/SM5, al^2 Cy lt^v$   
      $sf$   
 291  $ds S_2 G b pr/In(2L+2R)Cy, al^2 Cy lt^3 L^4$   
      $sp$   
 292  $ds_{33K} w/In(2L)Cy t^L R, Su(S) dp^{lv2} pr$   
 \*  $ds_{38K}$  . . . . . 328, 353, etc.  
 293  $ds_{38K}/In(2L)Cy, Cy dp^{lv2} b pr$   
 294  $dsr$   
 295  $dw-24F c1/SM5, al^2 Cy lt^v sp^2$   
 296  $dw-24F 1(2)cg, cg/SM5, al^2 Cy lt^v sp^2$   
 \*  $E(S)$  . . . . . (=  $EN-S$ ) . . . 335, 395, etc.  
 297  $ed Su(dx)^2$   
 298  $el$   
 \*  $en$  . . . . . 261, 269, 748  
 \*  $esc$  . . . . . 816  
 299  $ex$   
 300  $ex ds S^X ast^X/SM1, al^2 Cy sp^2$   
 \*  $fes$  . . . . . (see  $fs(2)B$ )  
 301  $fj 1(2)Su(H)/SM5, al^2 Cy lt^v sp^2$   
 302  $fj wt/SM5, al^2 Cy lt^v sp^2$   
 303  $fr/In(2L+2R)Cy, Cy dp^{lv2}$   
 304  $fr wt/SM5, al^2 Cy lt^v sp^2$   
 305  $Frd/In(2L+2R)Cy, Cy sp^2$   
 \*  $fs 2.1$  . . . . . (see  $fs(2)El$ )  
 306  $fs(2)B Alu lt/SM5, al^2 Cy lt^v sp^2$   
 \*  $fs(2)El$  . . . . . 249  
 307  $ft$   
 \*  $G^{rv}$  . . . . . 291  
 308  $G^{rv}/SM5, al^2 Cy lt^v sp^2$   
 \*  $Go$  . . . . . 232, 233  
 \*  $gp$  . . . . . 234  
 \*  $gt-4$  . . . . . 416  
 \*  $Hia$  . . . . . 439, 440  
 309  $hk$   
 310  $hk_2 pr$   
 \*  $hk_2$  . . . . . 205  
 311  $ho$   
 312  $hv/SM5, al^2 Cy lt^v sp^2$   
 313  $Hx/$  see also 215  
 314  $hy/SM5, al^2 Cy lt^v sp^2$   
 315  $hy a px sp/SM1, al^2 Cy sp^2$   
 \*  $ix_2$  . . . . . 374  
 \*  $ix$  . . . . . 204  
 316  $j$   
 317  $J/In(2L)NS$   
 318  $J_{34e}$   
 319  $kn$   
 320  $L_2$   
 321  $L_4$   
 322  $L_5$   
 323  $L_G$   
 324  $L_r$   
 325  $L_{si}$   
 326  $L$   
 \*  $1(2)301$  . . . . . 367  
 327  $1(2)39 a px slt sp/SM5, al^2 Cy lt^v sp^2$   
 328  $1(2)a bs_3, In(2L)t/bw^{v1}, ds_{33K}$   
 \*  $1(2)ax$  . . . . . 210  
 329  $1(2)ay b c sp/SM5, al^2 Cy lt^v sp^2$   
 \*  $1(2)Bld$  . . . . . 236  
 \*  $1(2)bw$  . . . . . 262  
 \*  $1(2)C$  . . . . . 399  
 \*  $1(2)cg$  . . . . . 296  
 \*  $1(2)crc$  . . . . . 270  
 329b  $1(2)gd a px or/In(2LR)0, Cy dp^{lvI} pr cn^2$   
 330  $1(2)gl a px or/SM5, al^2 Cy lt^v sp^2$   
 331  $1(2)H L/SM5, al^2 Cy lt^v sp^2$   
 \*  $1(2)M$  . . . . . 290  
 332  $1(2)mat/SM5, al^2 Cy lt^v sp^2$   
 333  $1(2)me/SM1, al^2 Cy sp^2$   
 \*  $1(2)mr$  . . . . . 738  
 \*  $1(2)R$  . . . . . 411

* 1(2)Su(H) . . . . .	301, 426	368	pk cn
* 1l . . . . .	363	369	pk tuf (sp <sup>2</sup> /+)
334 1l <sup>2</sup> . . . . .		*	Pm . . . . . (see bw <sup>v1</sup> )
335 lm/In(2L+2R)Cy, Cy S <sup>2</sup> dp <sup>lv2</sup> E(S)		*	Pm . . . . . (see bw <sup>v32g</sup> )
336 lt/T(Y;2)A		370	po <sup>2</sup> vg
337 lt std <sup>3</sup> /SM2, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		371	po <sup>2</sup>
338 lt <sup>3</sup> stw		372	pr
* lt <sup>v</sup> . . . . .	291, 864, 888	373	pr cn/T(Y;2)C
* lt . . . . .	206, 210, etc.	374	pr <sup>cn</sup> ix/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>
339 ltd		375	pr <sup>bw</sup>
340 lw		376	pu <sup>Gr</sup>
* lys . . . . .	691	*	Pu . . . . . 881
341 M(2)173/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		377	puf
* M(2)B <sub>S</sub> . . . . .	(see M(2)z <sup>B</sup> )	378	pw-c/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>
342 M(2)e <sub>S</sub> /In(2L+2R)Cy, Cy, In(2R)bw <sup>v34</sup>		379	px
343 M(2)H <sub>S</sub> /SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		380	px bs (old Berlin stock of Goldschmidt)
344 M(2)l <sub>S</sub> /SM1, al <sup>2</sup> Cy sp <sup>2</sup>		381	px bw sp/T(Y;2)J <sup>v1</sup>
345 M(2)m <sub>S</sub> /SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		382	px bw mr sp/bw <sup>v1</sup> , ds <sup>33k</sup>
346 M(2)S <sub>2</sub> <sup>3</sup> /SM2, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		383	px slt sp
347 M(2)S <sub>2</sub> <sup>9</sup> /SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		384	pym/In(2L+2R)Cy, Cy
* M(2)S <sub>3</sub> . . . . .	(see M(2)S <sub>2</sub> <sup>3</sup> )	385	pys
* M(2)S <sub>5</sub> . . . . .	(see M(2)H <sub>S</sub> <sup>6</sup> )	386	Q
* M(2)S <sub>6</sub> . . . . .	(see M(2)m <sup>2</sup> )	*	rc . . . . . 691
348 M(2)S <sub>7</sub> /SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		387	rd/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>
* M(2)S <sub>9</sub> . . . . .	(see M(2)S <sub>2</sub> <sup>9</sup> )	*	rd <sup>s</sup> . . . . . 231
* M(2)S <sub>11</sub> . . . . .	(see M(2)e <sup>2</sup> )	388	rdo
349 M(2)z/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		389	rdo <sup>2</sup> pr
350 M(2)z <sub>Sk</sub> b/In(2L)Cy, Cy dp <sup>lv2</sup> b pr		*	Rev <sup>B</sup> . . . . . 823
351 M(2)z <sub>B</sub> /SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		*	Rev . . . . . 753
* Mal . . . . .	694	390	rh <sub>4</sub>
352 mi/bw <sup>v32g</sup>		391	rk <sup>4</sup>
353 mr <sub>bs</sub> /bw <sup>v1</sup> , ds <sup>33k</sup>		392	rl
354 mr/In(2R)Cy, cn <sup>2</sup> Bld <sup>2</sup>		*	rn . . . . . 882
355 msf/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		*	Roi . . . . . 441
* N-2G . . . (= N-2) . . .	413	393	rub
356 net		394	Ruf/bw <sup>v1</sup> , ds <sup>33k</sup>
357 net al ex ds S ast shv ho rub/SM1, al <sup>2</sup> Cy sp <sup>2</sup>		*	Rvd . . . . . (see Rev <sup>B</sup> )
358 net ed Su(dx) <sup>2</sup>		395	S/In(2L+2R)Cy, Cy E(S) (K-pn)
359 nub <sub>2</sub> b pr		396	S <sub>2</sub> Sp ab <sup>2</sup> ltd/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>
360 nub <sup>2</sup>		*	S <sub>R</sub> . . . . . 335, 771
361 nw <sup>2</sup> /In(2L)Cy. In(2R)NS		397	S <sub>X</sub> /bw <sup>v1</sup> , ds <sup>33k</sup>
* or <sub>45a</sub> . . . . .	200, 330	*	S . . . . . 300
* or . . . . .	241	398	sca
362 pd		399	sca 1(2)C/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>
363 pd 1l <sup>2</sup>		400	SD-5/SM1, al <sup>2</sup> Cy sp <sup>2</sup>
364 pd 1l <sup>2</sup> sp		401	SD-72/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>
365 Pfd/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		*	sf . . . . . 239
366 pi/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		402	sf <sup>2</sup>
367 pi 1(2)301/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>		403	shr bw <sup>2b</sup> abb sp/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>
* Pin . . . . .	415	404	shv
		405	shv ho



\* Sk . . . . . 350  
 \* slt . . . . . 327, 383  
 406 sm px/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 407 sm px pd/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 408 so<sup>2</sup>  
 409 so<sup>2</sup> b cn  
 \* sp<sup>2</sup> . . . . . 201, 212, etc.  
 410 sp<sup>2</sup> bs  
 411 Sp/In(2L)t, 1(2)R<sub>v</sub><sup>2</sup>  
 412 Sp/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 413 Sp Bl N-2G/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 414 Sp J/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 415 Sp J L Pin/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 416 spd gt-4/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 417 sple  
 418 spt  
 419 std/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 420 stw<sup>2</sup>  
 421 stw<sup>3</sup>  
 422 stw<sup>5</sup>/T(Y;2)B  
 423 stw<sup>48</sup>  
 424 stw<sup>48</sup> ap blt tuf sp<sup>2</sup>  
 \* Su(dx)<sup>2</sup> . . (= Su-dx) . . 665  
 \* Su(dx)<sup>2</sup> . . (= Su<sup>2</sup>-dx) . . 358, 664  
 \* Su(er) . . (= Su-er) . . 693  
 425 Su(h)/In(2L+2R)Cy, Cy pr  
 426 Su(H) whd 1(2)Su(H)/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 \* Su(S) . . . . . 292  
 \* tet . . . . . 668  
 427 Tft/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 \* Tg . . . . . 818  
 \* tk . . . . . 238  
 428 tkd/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 429 tkv  
 430 tri vg<sup>No2</sup>/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 \* tu . . . . . 252  
 \* tu-36a . . (= tu<sup>36a</sup>) . . 604  
 431 tuf ltd  
 432 tyr-1 (p<sup>p</sup>); see also 229  
 433 Uf  
 434 vg  
 435 vg<sub>D</sub> bw  
 436 vg<sub>ni</sub>/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 437 vg<sub>No2</sub>  
 \* vg<sub>np</sub> . . . . . 430  
 438 vg<sub>np</sub>  
 439 vg<sub>nw</sub> Hia/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 440 vg<sub>U</sub> Hia/T(2;3)S<sup>M</sup> In(2L+2R)Cy, Cy  
 441 vg/In(2L)t, Roi, In(2R)Cy, bw sp<sup>2</sup> or  
 442 vst/SM5, al<sup>2</sup> Cy lt<sup>v</sup> sp<sup>2</sup>  
 443 whd

444 wt  
 \* wx<sup>wxt</sup> . . . (= wxt) . . . 237

### Chromosome 3

445 a(3)26  
 \* a-3 . . . . . (see a(3)26)  
 446 aa h  
 447 aa tu-36e  
 448 abd<sup>B</sup>  
 \* Antp . . . . . 826  
 449 app<sup>hg</sup>  
 450 as<sup>hg</sup> e<sup>s</sup>  
 451 as<sup>hg</sup> e<sup>s</sup>  
 \* Ata . . . . . 868  
 452 bar-3  
 \* Bd . . . . . 566  
 453 Bd<sup>G</sup>/In(3R)C, 1(3)a<sup>34e</sup>  
 454 bf/TM6, ss<sup>34e</sup> bx<sup>34e</sup> Ubx<sup>P15</sup> e  
 \* bod . . . . . 563  
 \* bp . . . . . (see bul<sup>bp</sup>)  
 455 bul<sup>bp</sup>  
 456 bul<sup>bp</sup>/TM1, Me ri sbd<sup>1</sup>  
 457 bv  
 \* bx<sup>3</sup> . . . . . 594, 608  
 458 bx<sup>34e</sup> Cbx Ubx bxd pbx/T(2;3)ap<sup>xa</sup>  
 459 bx<sup>34e</sup>  
 \* bxd<sup>107</sup> . . . . . 458, 595, 873  
 \* bxd<sup>107</sup> . . . . . 902  
 \* by . . . . . 576, 577  
 \* c(3)G . . (= c3G) . . . 600  
 460 ca  
 461 ca bv  
 462 ca<sup>2</sup>K-pn  
 463 ca<sup>nd</sup>  
 \* ca<sup>nd</sup> . . . . . 497  
 464 Cbx  
 465 cd  
 466 cmp ca/TM6, ss<sup>-</sup> bx<sup>34e</sup> Ubx<sup>P15</sup> e  
 467 cp  
 468 cp in ri p<sup>p</sup>  
 469 cu  
 470 cu kar<sup>8</sup>  
 471 cu kar ry<sup>8</sup>  
 472 cur  
 473 cv-c  
 474 cv-c sbd<sup>2</sup>  
 475 cv-d  
 \* Cyd . . . . . 489  
 476 D/G1  
 477 D<sup>3</sup> Sb ca<sup>2</sup>/In(3L+3R)P

478	det	516	in
479	Dfd/In(3LR)Cx	517	jv
480	Dfd <sup>r</sup>	518	jv Hn <sup>r</sup> h
481	Dl <sub>3</sub> H e <sup>s</sup> cd/In(3R)P, spr	519	jvl
482	Dl <sub>5</sub> /In(3R)C, e	*	k . . . . . 588
483	Dl <sub>7</sub> /In(3R)C, l(3)a	*	K-pn . . . . . 395, 462
484	Dl <sub>9</sub> /In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup> e <sup>s</sup>	*	kar <sup>2</sup> . . . . . 470, 471, 614
485	Dl <sub>11</sub> /In(3R)C, e	520	kar <sup>2</sup>
486	Dl <sub>12</sub> /In(3L+3R)P, Dfd ca	521	Ki
487	Dl <sub>12</sub> /In(3L+3R)P, Dfd ca	522	l(3)36d10/In(3LR)Cx, D
488	Dl <sub>13</sub> /In(3R)C, Sb e l(3)e	*	l(3)a . . . . . 453, 483
489	Dl <sub>14</sub> /In(3R)Cyd, Cyd	523	l(3)ac e <sup>s</sup> M(3)w/LVM
*	Dl <sub>B</sub> . . . . . 827	*	l(3)e . . . . . 488, 514, etc.
490	Dl <sup>x</sup> /In(3L+3R)P	*	l(3)PL . . . . . (In(3L+3R)P; In(3L+3R)P, Dfd ca)
*	Dr <sup>Mio</sup> . . . . . 556	*	l(3)PR . . . . . same as above <sup>4</sup>
491	Dr <sup>Mio</sup> /TM6, ss bx <sup>34e</sup> Ubx <sup>P15</sup> e	524	l(3)tr Sb/In(3L)P, In(3R)P18, Me Ubx e
492	drb	525	l(3)tr Ubx/TM1, Me ri sbd <sup>1</sup>
*	dsx . . . . . 551	*	l(3)W . . . . . 605, 827
493	dwh/In(3L+3R)P, Dfd ca	*	l(3)XaR . . . . . 867
*	e . . . . . 681, 682, In(3R)C	526	ld
*	e(dp) <sup>v</sup> . . . . . 690	527	Ly/D <sup>3</sup>
494	e <sub>11</sub> wo ro	528	Ly Sb/LVM
495	e <sup>s</sup>	*	M(3)36e . . . . . (see M(3)be <sup>36e</sup> )
496	e <sup>s</sup> ca <sup>nd</sup> /TM6, ss bx <sup>34e</sup> Ubx <sup>P15</sup> e	529	M(3)40130/In(3L+3R)P, Dfd ca
497	e <sup>s</sup> ca <sup>nd</sup> /TM6, ss bx <sup>34e</sup> Ubx <sup>P15</sup> e	*	M(3)124 . . . . . (see M(3)w <sup>124</sup> )
498	eg <sub>2</sub> /In(3LR)Cx	*	M(3)B <sub>2</sub> . . . . . (see M(3)w <sup>B2</sup> )
499	eg <sub>2</sub> /In(3LR)Cx	*	M(3)B <sub>2</sub> <sup>36e</sup> . . . . . (see M(3)w <sup>B2</sup> )
*	er . . . . . 684, 693	530	M(3)be <sup>36e</sup> /In(3R)C, l(3)a
500	eyg	531	M(3)h <sup>S37</sup> /In(3L)P, Me
*	fl . . . . . 541	532	M(3)h <sup>y</sup> /In(3L)P, Me
501	fz	533	M(3)S32/T(2;3)Me
502	gl <sub>2</sub> e <sup>4</sup>	534	M(3)S34/T(2;e)Me
503	gl <sub>3</sub>	535	M(3)S36/T(2;3)Me
504	gl <sub>60j</sub>	*	M(3)S37 . . . . . (see M(3)h <sup>S37</sup> )
505	gl	*	M(3)w . . . . . 523
506	G1 Sb/LVM	536	M(3)w/In(3R)C, e l(3)e
*	gm . . . . . 559, 623	537	M(3)w <sup>124</sup> /In(3R)C, e l(3)e
507	gro	538	M(3)w <sup>B</sup> /In(3R)C, e l(3)e
508	gs	539	M(3)w <sup>B2</sup> /In(3R)C, e l(3)e
509	h <sub>2</sub>	*	M(3)y . . . . . (see M(3)h <sup>y</sup> )
510	h	540	ma
511	H/In(3R)P, hp	541	ma fl
512	H <sub>2</sub> Pr/In(3R)C, e	542	mah
513	H <sub>3</sub> /T(2;3)ap <sup>Xa</sup>	543	Mc/T(2;3)ap <sup>Xa</sup>
514	H <sub>5</sub> <sup>c</sup> /In(3R)C, Sb e l(3)e	544	mwh
*	H . . . . . 620	545	N-X/T(2;3)ap <sup>Xa</sup>
*	Hm . . . . . 878	546	obt
*	Hn . . . . . 879	547	p
*	Hn <sub>r</sub> . . . . . 518	548	p <sup>p</sup>
515	Hn <sub>r3</sub> sr	549	p <sup>p</sup> bx sr e <sup>s</sup>
*	Hu . . . . . 828		

- 550 p<sup>P</sup> cu  
 551 p<sup>P</sup> dsx/TM6, ss<sup>-</sup> bx<sup>34e</sup> Ubx<sup>P15</sup> e  
 552 pb/In(3LR)Cx<sup>Xa</sup>  
 553 pbx/T(2;3)ap<sup>Xa</sup>  
 554 Pc/TM1, Me ri sbd<sup>1</sup>  
 \* Pdr . . . . . 692  
 555 Pr/In(3R)C. e<sup>+</sup>  
 556 Pr Dr/TM3, y<sup>+</sup> ac<sup>+</sup> ri p<sup>P</sup> sep bx<sup>34e</sup> e<sup>s</sup>  
 557 Pt/T(2;3)ap<sup>Xa</sup>, ca  
 558 pyd  
 559 R Ly/In(3L)P, gm  
 560 ra  
 561 red  
 562 ri  
 563 ri bod e<sup>s</sup>/In(3L)P, Me, In(3R)C, Sb e  
 1(3)e  
 564 ri p<sup>P</sup>/T(Y;2;3)F, st  
 565 ro  
 566 ro Bd ca/In(3R)C, 1(3)a  
 567 ro ra ca/T(2;3)Me  
 567b roe p<sup>P</sup>  
 568 rs<sup>2</sup>  
 569 rsd<sup>2</sup>  
 \* rt . . . . . 587  
 570 ru  
 571 ru h th st p<sup>P</sup> H e<sup>s</sup> ro/TM6, ss<sup>-</sup> bx<sup>34e</sup>  
 Ubx<sup>P15</sup> e  
 572 ru h th st cu sr e<sup>s</sup> ca  
 573 ru h th st cu sr e<sup>s</sup> ca/TM3, ru Sb-Ser<sup>34e</sup>  
 574 ru h th st cu sr e<sup>s</sup> Pr ca/TM6, ss<sup>-</sup> bx<sup>34e</sup>  
 Ubx<sup>P15</sup> e  
 575 ru h th st p<sup>P</sup> cu sr e<sup>s</sup>  
 576 ru lxd by  
 577 ru<sup>g</sup> jv se by  
 578 ry<sup>8</sup>  
 \* ry . . . . . 101 . . . 101 . 471  
 579 Sb/In(3LR)Ubx . . . . . Ubx  
 580 Sb H/In(3R)C, cd<sup>Xa</sup>  
 581 Sb Ubx/T(2;3)ap<sup>Xa</sup>  
 582 Sb<sup>63b</sup>/In(3LR)Ubx<sup>130</sup>, Ubx<sup>130</sup> e<sup>s</sup>  
 583 Sb<sup>Spi</sup>/In(3LR)Cx  
 \* Sb<sup>v</sup> . . . . . 885, 886  
 \* sbd<sup>2</sup> . . . . . 603  
 584 sbd<sup>105</sup>  
 \* sbd<sup>1</sup> . . . . . 757  
 \* sbd<sup>1</sup> . . . . . 456, 525  
 585 se  
 586 se h<sup>2</sup>  
 587 se rt<sup>2</sup> th/In(3L)P, Me  
 588 se ss k e<sup>s</sup> ro  
 \* sed . . . . . (see Hn<sup>r3</sup>)  
 \* sep . . . . . 556, 825, 885  
 589 Ser/In(3R)C. e 1(3)e  
 590 snb  
 591 sr  
 592 sr gl  
 593 ss  
 594 ss bx Su(ss)<sup>2</sup>  
 595 ss bxd k e<sup>s</sup>/T(2;3)ap<sup>Xa</sup>  
 596 ss<sup>a</sup>  
 597 ss<sup>aB</sup>  
 598 ss<sup>a40a</sup>  
 599 st  
 600 st c(3)G ca/TM1, me ri sbd<sup>1</sup> (sp<sup>2</sup>)  
 601 st in ri p<sup>P</sup>  
 602 st Ki p<sup>P</sup>  
 603 st sbd e<sup>s</sup> ro ca  
 604 st sr e<sup>s</sup> ro ca (tu-36a)  
 605 st sr H<sup>2</sup> ca/In(3R)P, st 1(3)W ca  
 606 st<sup>sp</sup>  
 \* su(pd)<sub>B</sub> . . (= su-pd)<sub>B</sub> . . 679  
 607 su(pr)<sub>2</sub>/TM6, ss<sup>-</sup> bx<sup>34e</sup> Ubx<sup>P15</sup> e (pr)<sub>2</sub>  
 608 su(Hw)<sub>2</sub> bx bxd/TM1, Me ri sbd<sup>1</sup> (sp<sup>2</sup>)  
 \* Su(ss)<sub>2</sub> . . (= Su<sub>2</sub>-ss) . . 594  
 609 su(t) (t)  
 \* su(tu) . . (= su-tu) . . 693  
 610 su(ve)ru ve h th  
 611 th  
 612 th st cp  
 613 th st pb p<sup>P</sup>/TM6, ss<sup>-</sup> bx<sup>34e</sup> Ubx<sup>P15</sup> e  
 614 th st pb p<sup>P</sup> cu kar su(Hw)<sub>2</sub> jvl ss bx sr  
 gl/TM6, ss<sup>-</sup> bx<sup>34e</sup> Ubx<sup>P15</sup> e  
 \* tra<sub>D</sub> . . . . . 674  
 \* tra . . . . . 844  
 615 Tri/In(3LR)DcxF  
 616 tt wo  
 617 Tu (= Tubby)  
 \* tu-36e . . (= tu<sup>36e</sup>) . . 447  
 \* tuh-3 . . . . . 673  
 618 tx  
 619 Ubx<sup>4</sup>/In(3L+3R)P, Dfd ca  
 620 Ubx<sup>61d</sup>/H<sup>57c</sup>  
 \* Ubx<sup>101</sup> . . . . . 579  
 \* Ubx<sup>130</sup> . . . . . 484, 582, 674, etc.  
 621 ve  
 622 ve h th  
 623 ve R/In(3L)P, gm  
 \* vo-3 . . . . . (see e(dp<sup>v</sup>))  
 624 W  
 625 W Sb/In(3LR)Cx  
 626 We/In(3L)P, Me, In(3R)C, e 1(3)e  
 627 wk/In(3L+3R)P, Dfd ca  
 628 wo

\* Xa . . (= T(2;3)ap<sup>Xa</sup>) . . , 458, 543, etc.

# Multichromosomal Stocks

## Chromosome 4

629 ar/ey<sup>D</sup>  
 630 bt<sup>R</sup>  
 631 bt<sup>D</sup>ey<sup>R</sup>sv<sup>n</sup>  
 632 bt<sup>D</sup>/ci<sup>D</sup>Cat  
 633 Ce<sup>2</sup>/spa<sup>R</sup>  
 634 ci ey<sup>R</sup>sv<sup>n</sup>  
 635 ci ey<sup>R</sup>sv<sup>n</sup>  
 636 ci gvl bt<sup>R</sup>  
 637 ci gvl ey<sup>R</sup>sv<sup>n</sup>  
 638 ci<sup>36</sup>l<sup>57</sup>  
 639 ci<sup>57</sup>g  
 640 ci<sup>D</sup>/ey<sup>D</sup>  
 641 ci<sup>W</sup>/ey<sup>D</sup>  
 642 ci<sup>W</sup>  
 643 ey<sup>2</sup>  
 644 ey<sup>4</sup>  
 645 ey<sup>D</sup>  
 \* ey<sup>R</sup> . . . . . 629, 641, 662  
 \* ey<sup>R</sup> . . . . . 634, 635, etc.  
 646 gvl<sup>R</sup>  
 647 gvl ey<sup>R</sup>sv<sup>n</sup>  
 648 gvl ey<sup>R</sup>sv<sup>n</sup>  
 649 l(4)2<sup>c</sup>/ci<sup>D</sup> (Hochman)  
 650 l(4)4<sup>b</sup>/ci<sup>D</sup> "  
 651 l(4)6<sup>b</sup>/ci<sup>D</sup> "  
 652 l(4)14<sup>b</sup>/ci<sup>D</sup> "  
 653 l(4)15<sup>2</sup>/ci<sup>D</sup> "  
 654 l(4)21/ci<sup>D</sup> "  
 655 l(4)22/ci<sup>D</sup> "  
 656 l(4)25/ci<sup>D</sup> "  
 \* l(4)AM-1 . . . . . (see l(4)22)  
 \* l(4)PT-1 . . . . . (see l(4)6<sup>b</sup>)  
 \* l(4)PT-2 . . . . . (see l(4)2<sup>f</sup>)  
 \* l(4)PT-3 . . . . . (see l(4)4<sup>c</sup>)  
 \* l(4)SLC-1 . . . . . (see l(4)15<sup>2</sup>)  
 \* l(4)ST-1 . . . . . (see l(4)21<sup>b</sup>)  
 \* l(4)ST-2 . . . . . (see l(4)14<sup>b</sup>)  
 \* l(4)ST-3 . . . . . (see l(4)25)  
 \* Mal . . . . . 694  
 657 spa<sup>Cat</sup>/ci<sup>D</sup>  
 658 spa<sup>pol</sup>  
 659 spa<sup>p65</sup>  
 660 spa<sup>35a</sup>  
 661 sv<sup>de</sup>/ey<sup>D</sup>  
 662 sv<sup>n</sup>/ey<sup>D</sup>  
 663 sv<sup>n</sup>

664 br<sup>3</sup>dx<sup>st</sup>;ed Su(dx)<sup>2</sup>(1;2)  
 665 dx<sup>st</sup>;Su(dx)(1;2)  
 666 e(S)<sup>2</sup>/FMA3, y<sup>2</sup>;al S ast ho/SML, al<sup>2</sup> Cy<sup>sp</sup>(1;2)  
 667 lz<sup>D</sup>/In(1)dl-49, m<sup>2</sup>g<sup>4</sup>;bw<sup>vl</sup>/In(2L+2R)Cy,  
 Cy(1;2)  
 668 os<sup>S</sup>;tet(1;2)  
 669 v;bw(1;2)  
 670 v;In(2R)bw<sup>VDel</sup>/SML, al<sup>2</sup> Cy<sup>sp</sup>(1;2)  
 671 w<sup>ch</sup>/FMA3, y<sup>2</sup>;Su(w<sup>ch</sup>)/In(2L+2R)Cy,  
 Cy(1;2)  
 672 sc z w<sup>2</sup>rst;halo(1;3)  
 673 tuh-1;tuh-3(1;3)  
 674 w<sup>a</sup>v/FMA3, y<sup>2</sup>;tra/In(3LR)Ubx<sup>130</sup>, Ubx<sup>130</sup>  
 e<sup>(1;3)</sup>  
 675 w<sup>e</sup>;Dp(2;3)P/TM6, ss<sup>-</sup>bx<sup>34e</sup>  
 Ubx<sup>P15</sup>e(1;3)  
 676 y;mwh(1;3)  
 677 y<sup>2</sup>su(Cbx)<sup>2</sup>v/FMA3, y<sup>2</sup>;Cbx/T(2;3)ap<sup>Xa</sup>(1;3)  
 678 y<sup>2</sup>e(bx)<sup>2</sup>w<sup>1</sup>/FMA3, y<sup>2</sup>;sbd<sup>ss</sup>bx<sup>34e</sup>/TM1,  
 Me ri sbd<sup>1</sup>(1;3)  
 679 ptg;px pd;su(pd)(1;2;3)  
 680 FMA3, y<sup>2</sup>;net;sbd<sup>pol</sup>;spa<sup>pol</sup>(1;2;3;4)  
 681 y f:=;bw;e;ci ey<sup>R</sup>(1;2;3;4)  
 682 y f:=;bw;e;spa<sup>pol</sup>(1;2;3;4)  
 683 al dp b Bl c px sp/In(2L+2R)Cy, Cy;  
 D/In(3L+3R)P(2;3)  
 684 b Su(er)<sup>+</sup>bw;st er(2;3)  
 685 bw<sup>4</sup>;st(2;3)  
 686 bw<sup>vl</sup>;st(2;3)  
 687 bw<sup>vl</sup>, dp b/In(2L+2R)Cy, Cy<sup>sp</sup><sup>2</sup>;Sb/In(3LR)  
 DcxF(ru h ca?)(2;3)  
 688 bw<sup>vl</sup>, ds<sup>33k</sup>/In(2L+2R)Cy, Cy;H/In(3R)Mo,  
 sr(2;3)  
 689 cn;ry<sup>2</sup>(2;3)  
 690 dp<sup>v</sup>;e(dp<sup>v</sup>)(2;3)  
 691 lys rc;ss(2;3)  
 692 px pd;Pdr H, Dp(2;3)P/Pdr(2;3)  
 692b Ubx<sup>P15</sup>e(2;3)  
 693 Su(er)tu bw;st er su(tu)(2;3)  
 694 pr;Mal(2;4)

## Attached-X

695 br ec/y<sup>3d</sup>  
 696 f B/su(s)<sup>S</sup>v  
 \* FMA3, y<sup>2</sup> . . (= FMA3) . . 128, 129, etc.  
 697 w<sup>bf3</sup>/sn<sup>36a</sup>  
 698 y/g<sup>2</sup>ty

699  $y_{pn}/FM6, y^{3ld} sc^8 dm B$   
 \*  $y_{pn} v$  . . . . . 709  
 \*  $y v bb$  . . . . . 786  
 \*  $y v f$  . . . . . 720  
 \*  $y v f car$  . . . . . 780  
 \*  $y w bb$  . . . . . 26  
 \*  $y_{2w} f$  . . . . . 783  
 \*  $y_{2sc} w^a ec^a$  . . . . . 712  
 700  $y_{su(w)} w^a bb/y sc^4L sc^8R$

Attached Autosomal Arms

701 C(2L)P3, +; C(2R)P3, +  
 702 C(2L)P3, j; C(2R)P4, px  
 703 C(2L)P4, dp; C(2R)P4, px  
 704 C(3L)P3, ri; C(3R)P3, sr  
 705 C(3L)P6, +; C(3R)P6, +  
 706 C(4)P1, ci ey<sup>R</sup>/gvl sv<sup>n</sup>  
 707 C(4)P2, ci ey<sup>R</sup>/gvl sv<sup>n</sup>

Attached-XY

708  $v f B, XY/y^2 su(w^a) w^a bb$   
 709  $y^{59b} su(w^a) w^a, XY \cdot Y/y_{pn} v$  (Extra Y  
 present)  
 710  $Y/g^2 B \cdot Y^L$  and  $y f:=(dp^{olv})$  (Stern)  
 711  $Y^S X \cdot Y^L, In(1)EN, In(1)dl-49, Y^S y \cdot Y^L/$   
 $y X \cdot Y; bw; e; ci ey$

Triploid

712a C(1)RM, In(1)dl-49, v<sup>Of</sup> f/FM7  
 712b C(1)RM, y w fa<sup>no</sup>/FM6 ♀ and FM6, y<sup>3ld</sup> sc<sup>8</sup> dm  
 B/B<sup>S</sup> Y y<sup>+</sup> ♂  
 712c C(1)RM, y<sup>2</sup> sc w<sup>a</sup> ec/FM6 ♀ and FM6, y<sup>3ld</sup> sc<sup>8</sup>  
 dm B/B<sup>S</sup> Y y<sup>+</sup> ♂

Extra-Y

713 In(1)w<sup>m4L</sup> N<sup>264-84R</sup>, y sn/FM3, y<sup>3ld</sup> sc<sup>8</sup>  
 dm B 1/Y<sup>+</sup> dm sn ♂ (DIS 28: 137)  
 714 y v f mal/mal<sup>+</sup> Y<sup>+</sup> q; In(1)dl-49, B<sup>M1</sup>, Df  
 (1)mal<sup>+</sup>, y v sn<sup>+</sup>/mal<sup>+</sup> Y ♂  
 715 y v f mal/y<sup>+</sup> mal<sup>+</sup> Y q; l(T2-4a)/y<sup>+</sup> mal<sup>+</sup>  
 Y ♂  
 \* Y<sup>-bb</sup> . . . . . 786  
 716 In(X<sup>c2</sup>)w<sup>vc</sup>/In(1)dl-49, y w lz<sup>s</sup> q; In(1)d  
 dl-49, y w lz<sup>s</sup>/sc<sup>+</sup> Y ♂  
 717 X<sup>c1</sup>, y/y f:=/y<sup>+</sup> Y  
 718 X<sup>c2</sup>, cv v f/ClB, v

Closed-Y

719 R(Y)bw<sup>+</sup>/X; bw ("MYR")  
 \* Y<sup>Lc</sup> bw<sup>+</sup> . . . . . (see R(Y)bw<sup>+</sup>)  
 720 Y<sup>Lc</sup>/y w Y and y v f

DeficienciesDeficiencies-X

721 Df(1)260-1/FM4, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 722 Df(1)B<sup>263-20</sup>/In(1)sc<sup>7</sup>, In(1)AM<sup>sc</sup> car  
 723 Df(1)bb, y sl<sup>2</sup> bb-/FM4, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 724 Df(1)bb, y v car bb-/In(1)AM  
 \* Df(1)bb<sup>1</sup> . . . . . 738  
 725 Df(1)ct<sup>268-42</sup>, y/FM4, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 726 Df(1)g<sup>f B/In(1)AM</sup>  
 727 Df(1)m<sup>259-4c</sup>/FM4, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 728 Df(1)mal/In(1)dl-49, lz<sup>s</sup>  
 729 Df(1)N<sup>8/FM1</sup>, y<sup>3ld</sup> sc<sup>8</sup> w<sup>a</sup> lz<sup>s</sup> B  
 730 Df(1)N<sup>264-39</sup>, w/FM4, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 731 Df(1)N<sup>264-105</sup>/FM1, y<sup>3ld</sup> sc<sup>8</sup> w<sup>a</sup> lz<sup>s</sup> B  
 \* Df(1)rst<sup>2</sup> . . . . . 95  
 \* Df(1)sc<sup>4L</sup> sc<sup>8R</sup> . . . . . 795  
 \* Df(1)sc<sup>8</sup> . . . . . 769  
 732 Df(1)syr<sup>258-11</sup>, Dp(1;f)101, spl/y f:=  
 733 Df(1)w<sup>258-42</sup>, y/In(1)dl-49, y Hw m<sup>2</sup> 4  
 734 Df(1)w<sup>258-45</sup>, y/FM1, y<sup>3ld</sup> sc<sup>8</sup> w<sup>a</sup> lz<sup>s</sup> B  
 735 Df(1)w<sup>258-48</sup>, y/FM4, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 736 Df(1)w<sup>258-48</sup>, y sc spl; Dp(1;3)w<sup>vco</sup>; y f:=

Deficiencies-Y

737 Df(Y)Y<sup>bb-</sup>  
 738 Df(Y)Y<sup>st</sup>      Df(Y)Y<sup>bb-</sup><sub>1</sub><sup>e</sup>, y<sub>1</sub><sup>eq</sup><sub>st</sub> and w<sup>e</sup> bb<sup>1</sup>;Y<sup>+</sup>;In(2L+2R)NS, px sp/l(2)mr<sup>2</sup>

Deficiencies-2

739 Df(2)M33a      Df(2)M33a/bw<sup>V32g</sup>  
 \* Df(2)MB      . . . . . (see Df(2L)M-z<sup>B</sup>)  
 740 Df(2)MS4      Df(2)MS4/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 741 Df(2)MS8      Df(2)MS8/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 742 Df(2)MS10      Df(2)MS10/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 743 Df(2)rl<sup>10a</sup>      Df(2)rl<sup>10a</sup> lt cn/bw<sup>V1</sup>, ds<sup>33k</sup>  
 744 Df(2L)al<sup>B</sup>      Df(2L)al/In(2L+2R)Cy, Cy E(S)  
 745 Df(2L)M-z<sup>B</sup>      Df(2L)M-z/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 746 Df(2L)S2      Df(2L)S2/In(2L+2R)Cy, Cy E(S)  
 747 Df(2L)S3      Df(2L)S3/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 748 Df(2R)42<sup>5</sup>      Df(2R)42<sup>5</sup> en/SM1, Al<sup>2</sup> Cy sp<sup>2</sup>  
 749 Df(2R)bw<sup>VDe2L</sup>      Df(2R)bw<sup>VDe2L</sup>/T(2;3)ap<sup>R</sup>  
 750 Df(2R)bw<sup>2</sup>      Df(2R)bw<sup>2</sup>, In(2R)Cy<sup>R</sup>/Gla<sup>2</sup>  
 751 Df(2R)Px<sup>B</sup>      Df(2R)Px<sup>B</sup>, bw sp/SM1, al<sup>2</sup> Cy<sup>v</sup> sp<sup>2</sup>  
 752 Df(2R)vg<sup>C</sup>      Df(2R)vg<sup>C</sup>/SM5, al<sup>2</sup> Cy<sup>B</sup> lt sp<sup>2</sup>  
 753 Df(2R)vg<sup>C</sup>      Df(2R)vg<sup>C</sup>/In(2LR)Rev<sup>v</sup>  
 754 Df(2R)vg<sup>D</sup>      Df(2R)vg<sup>D</sup>/SM5, al<sup>2</sup> Cy lt<sup>D</sup> sp<sup>2</sup>  
 \* Df(2R)vg      . . . . . (= vg<sup>D</sup>) . . . . . 436

Deficiencies-3

\* Df(3L)Hn      . . . . . 879  
 \* Df(3L)Ly      . . . . . (= Ly) . . . . . 527, 528  
 755 Df(3R)M-S31      Df(3R)M-S31/T(2;3)Me<sup>130</sup>  
 756 Df(3R)ry<sup>105</sup>      Df(3R)ry/In(3LR)Ubx<sup>130</sup> Ubx<sup>130</sup> e<sup>s</sup>  
 757 Df(3R)sbd<sup>105</sup>      Df(3R)sbd<sup>105</sup>, p<sup>p</sup> sbd<sup>105</sup> bx sr e<sup>s</sup>/LVM

Deficiencies-4

758 Df(4)M      Df(4)M/ey<sup>D</sup>

Duplications

\* Dp(1;f)24      . . . . . (= Del(1)24) . . . . . 62, 808  
 759 Dp(1;f)101      Dp(1;f)101;In(1)sc<sup>8</sup>, Df(0+ac).w<sup>a</sup> sc<sup>8</sup>.  
 760 Dp(1;f)107      Dp(1;f)107;In(1)sc<sup>8</sup>, Df(0+ac).w<sup>a</sup> sc<sup>8</sup>.  
 761 Dp(1;f)118      Dp(1;f)118;In(1)sc<sup>8</sup>, Df(0+ac).w<sup>a</sup> sc<sup>8</sup>.  
 762 Dp(1;f)135      Dp(1;f)135, y<sup>2</sup>;In(1)sc<sup>8</sup>, Df(0+ac).w<sup>a</sup> sc<sup>8</sup>.  
 763 Dp(1;f)R<sup>c2</sup>      Dp(1;f)R/y dor /y dor  
 \* Dp(1;f)X<sup>9</sup>      . . . . . (see Dp(1;f)R)  
 764 Dp(1;f)z      Dp(1;f)z<sup>9</sup>, Df(1)sc<sup>4</sup>/y f:=  
 765 Dp(1;1)112      Dp(1;1)112, y<sup>f</sup> (homozygous stock)  
 766 Dp(1;1)1z<sup>S1</sup>      Dp(1;1)1z, lz<sup>50e</sup> lz<sup>y4</sup>/y f:=  
 767 Dp(1;Y)sc<sup>S1</sup>      sc<sup>S1</sup>.Y/y.Y; y f:=;cn bw;(e/+)



768	Dp(1;3)126 <sub>J4</sub>	Dp(1;3)126 <sub>J4</sub> ; v f/In(3L+3R)P, Dfd ca
769	Dp(1;3)sc <sub>vco</sub>	Dp(1;3)sc <sub>J4</sub> /Df(1)sc <sub>a</sub> , w
*	Dp(1;3)w	..... 736
770	Dp(2;2)S	Dp(2;2)S, (S ast) (S ast) net dp cl/In(2L+2R)Cy, Cy E(S)
*	Dp(2;3)P	..... 692
771	Qn(2;2)S	Qn(2;2)S, (ast) <sub>5</sub> , al ho/In(2L+2R)Cy, Cy S <sup>2</sup> E(S)

### Inversions

#### Inversions-X

772	In(1)AB	In(1)AB/y f:=
*	In(1)AM <sub>M1</sub>	..... 14, 88, etc.
773	In(1)B <sub>M2</sub>	In(1)B <sub>M2</sub> , v B <sub>M1</sub> (tan-like); .. see also .. 784, 785, etc.
774	In(1)B <sub>M2</sub>	In(1)B <sub>M2</sub> (rv) f <sub>B27</sub> (reinv.; mosaic)
775	In(1)B <sub>M2</sub>	In(1)B <sub>M2</sub> , f <sub>B27</sub> B <sub>M2</sub> /ClB
776	In(1)B <sub>M2</sub>	In(1)B <sub>M2</sub> , v <sub>rv</sub> B <sub>M2</sub>
*	In(1)bb	..... 723, 724
777	In(1)ClB <sub>36d</sub>	In(1)Cl, sc t <sup>2</sup> v sl B <sub>36d</sub> (= ClB) .. 80, 718
778	In(1)ClB <sub>36d</sub>	In(1)Cl, sc t <sup>2</sup> v sl B <sub>36d</sub> (= ClB <sub>36d</sub> ) .. 851
779	In(1)d1-49	In(1)d1-49, tyl
780	In(1)d1-49	In(1)d1-49, tyl <sup>1</sup> bb <sup>1</sup> /y v f car
781	In(1)d1-49	In(1)d1-49, v <sub>Of</sub> f <sub>n</sub>
782	In(1)d1-49	In(1)d1-49, y fa <sup>2</sup> g <sup>4</sup>
*	In(1)d1-49	In(1)d1-49, y Hw m <sup>2</sup> g <sup>4</sup> .. 733, 798
783	In(1)d1-49	In(1)d1-49, y Su(Hw) Hw m <sup>2</sup> g <sup>4</sup> /y w f; (nub/+)
784	Ins(1)d1-49, B <sub>M1</sub>	In(1)d1-49, In(1)B <sub>M1</sub> , l(1)J1 sc <sup>3</sup> q <sup>2</sup> c <sup>2</sup> p <sup>2</sup> tg B <sub>M1</sub> /In(1)sc <sup>1</sup> sc <sup>8R</sup> , y sc <sup>1</sup> sc <sup>1</sup> pn w ec rb cm ct sn ras g f os os car l/1(1)J1.Y (= "Maxy")
785	Ins(1)d1-49, B <sub>M1</sub>	In(1)d1-49, In(1)B <sub>M1</sub> , sc v <sub>B</sub> B <sub>M1</sub> (homozygous)
786	Ins(1)d1-49, B <sub>M1</sub>	In(1)d1-49, In(1)B <sub>M1</sub> , y/Y <sup>-bb</sup> and y v <sub>bb</sub> /Y <sup>-bb</sup>
787	Ins(1)d1-49, B <sub>M1</sub>	In(1)d1-49, In(1)B <sub>M1</sub> , y sc v cu-X B <sub>M1</sub>
788	In(1)e(bx)	In(1)e(bx), e(bx)/y f:=
*	In(1)EN	..... 711
*	Ins(1)FM1	In(1)FM1, In(1)d1-49, y <sup>3ld</sup> sc <sup>8</sup> w <sup>1z</sup> B (= FM1) .. 18, 27, etc.
*	In(1)FM3	In(1)FM3, y <sup>3ld</sup> sc <sup>8</sup> dm B 1 (= FM3) .. 9, 53, etc.
*	In(1)FM4	In(1)FM4, y <sup>3ld</sup> sc <sup>8</sup> dm B (= FM4) .. 712, 723
789	In(1)FM6	In(1)FM6 y <sup>3ld</sup> sc <sup>8</sup> dm B/y f:=; .. see also 32, 33, etc.
790	In(1)FM7	In(1)FM7a, y <sup>2</sup> w <sup>a</sup> v <sup>Of</sup> B (homozygous) (see DIS 44: 101)
*	In(1)FMA3 <sub>264-84</sub>	In(1)FMA3, y <sup>2</sup> (= FMA3) .. 128, 129
791	In(1)N <sub>3</sub>	In(1)N <sub>3</sub> , y/FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B
792	In(1)rst <sub>3</sub>	In(1)rst <sub>3</sub> , rst <sub>3</sub> (homozygous)
793	In(1)rst <sub>3</sub>	In(1)rst <sub>3</sub> , y rst <sub>3</sub> car bb
*	In(1)S <sub>4</sub>	..... 809, 813
794	In(1)sc <sub>4L</sub> <sup>8R</sup>	In(1)sc <sub>4L</sub> , y <sup>8R</sup>
795	In(1)sc <sub>7</sub> sc <sup>8R</sup>	In(1)sc <sub>7</sub> sc <sub>7</sub> , y; .. see also 700
796	In(1)sc <sub>7</sub>	In(1)sc <sub>7</sub> , sc <sub>7</sub>
797	In(1)sc <sub>7</sub>	In(1)sc <sub>7</sub> , sc <sub>7</sub> w <sup>a</sup>
798	Ins(1)sc <sub>7</sub> , AM	In(1)sc <sub>7</sub> , In(1)AM, sc <sub>7</sub> /In(1)d1-49, y <sup>3ld</sup> Hw m <sup>2</sup> g <sup>4</sup>
799	Ins(1)sc <sub>7</sub> , AM <sub>M1</sub>	In(1)sc <sub>7</sub> , In(1)AM <sub>M1</sub> , sc <sub>7</sub> car/FM4, y <sup>3ld</sup> sc <sup>8</sup> dm (without B)
800	Ins(1)sc <sub>7</sub> , B <sub>M1</sub>	In(1)sc <sub>7</sub> , In(1)B <sub>M1</sub> , sc <sub>7</sub> w <sup>a</sup> B <sub>M1</sub> /y f:=

801	In(1)sc <sup>8</sup>	In(1)sc <sup>8</sup> , sc <sup>8</sup>
802	In(1)sc <sup>8</sup>	In(1)sc <sup>8</sup> , sc <sup>8</sup> cv y f/y f:=
803	In(1)sc <sup>8</sup>	In(1)sc <sup>8</sup> , y <sup>3ld</sup> sc <sup>8</sup> w <sup>a</sup>
804	In(1)sc <sup>8R</sup> dl-49	In(1)sc <sup>8</sup> , In(1)dl-49, y <sup>3ld</sup> sc <sup>8</sup> (homozygous)
*	In(1)sc <sup>9</sup>	. . . . . 700, 784
805	In(1)sc <sup>260-14</sup>	In(1)sc <sup>260-14</sup> Bx f t w <sup>a</sup> (homozygous)
806	In(1)sc <sup>260-22</sup>	In(1)sc <sup>260-22</sup> , sc <sup>260-22</sup>
807	In(1)sc <sup>J1</sup>	In(1)sc <sup>J1</sup> , sc <sup>260-22</sup>
808	In(1)sc <sup>S1</sup>	In(1)sc <sup>S1</sup> ; Dp(1;f)24
*	Ins(1)sc <sup>S1L</sup> dl-49	In(1)sc <sup>S1L</sup> In(1)dl-49, y v B <sup>8</sup> . . . . . 178
*	Ins(1)sc <sup>S1L</sup> , sc <sup>8R</sup>	In(1)sc <sup>S1L</sup> , y sc <sup>8R</sup> sc <sup>8R</sup> pn w ec rb cm ct <sup>6</sup> sn <sup>3</sup> ras <sup>2</sup> g <sup>2</sup> f
809	Ins(1)sc <sup>S1L</sup> , S, sc <sup>8R</sup>	os os <sup>S1L</sup> car l . . . . . 8R . . . . . 784
*	In(1)sc <sup>S1R</sup>	In(1)sc <sup>S1L</sup> , In(1)S, In(1)sc <sup>S1R</sup> , sc <sup>8R</sup> sc <sup>8R</sup> w <sup>a</sup> B (= Miller-5)
810	In(1)w <sup>m4</sup>	. . . . . 813
811	In(1)w <sup>m4</sup>	In(1)w <sup>m4</sup> (bb?)
812	In(1)y <sup>3P</sup>	In(1)w <sup>3P</sup> , y <sup>3P</sup> cv m f/y f:=
813	Ins(1)y <sup>3PL</sup> , S, sc <sup>S1R</sup>	In(1)y <sup>3PL</sup> y <sup>3P</sup> B (B reverted)
814	In(1)y <sup>4</sup>	In(1)y <sup>4</sup> , In(1)S, In(1)sc <sup>S1R</sup> /y f:=; sc <sup>19i</sup> /In(2L+2R)Cy, Cy

2L Inversions

815	In(2L)Cy	In(2L)Cy, al <sup>2</sup> ast <sup>3</sup> b pr (does not carry Cy mutant)
*	In(2L)Cy <sup>L R</sup>	In(2L)Cy, Cy dp <sup>lv2</sup> b pr . . . . . 293, 350
*	In(2L)Cy <sup>L t</sup>	In(2L)Cy <sup>L t</sup> , Su(S) dp <sup>lv2</sup> pr . . . . . 292
*	In(2L)NS	. . . . . 317
*	In(2L)t	. . . . . 328
816	In(2L)t	In(2L)t, esc c sp/SM5, al <sup>2</sup> Cy <sup>lt</sup> sp <sup>2</sup>
817	In(2L)t	In(2L)t, lt l L sp <sup>2</sup> /bw <sup>VI</sup> , ds <sup>33k</sup> sp <sup>2</sup>
*	In(2L)t	In(2L)t, l(2)R . . . . . 411
818	In(2L)Tg	In(2L)Tg, Tg/SM5, al <sup>2</sup> Cy <sup>lt</sup> sp <sup>2</sup>

2L + 2R Inversions

819	In(2L+2R)Cy	In(2L+2R)Cy, al <sup>2</sup> E(S) cn <sup>2</sup> sp <sup>2</sup> (does not carry Cy mutant)
*	In(2L+2R)Cy	In(2L+2R)Cy, al <sup>2</sup> Cy lt <sup>2</sup> L sp <sup>2</sup> . . . . . 291, 864, 888
*		In(2L+2R)Cy, Cy . 45a . 2 . 45a . . . . . 273, 384, etc.
*		In(2L+2R)Cy, Cy bw <sup>lv2</sup> sp <sup>2</sup> or . . . . . 241
*		In(2L+2R)Cy, Cy dp <sup>lvI</sup> . . . . . 303
*		In(2L+2R)Cy, Cy dp <sup>lvI</sup> Bl L sp <sup>2</sup> . . . . . 204
*		In(2L+2R)Cy, Cy dp <sup>lvI</sup> pr . . . . . 838
*		In(2L+2R)Cy, Cy E(S) <sub>2</sub> . . . . . 395, 744, etc.
*		In(2L+2R)Cy, Cy hk <sup>2</sup> . . . . . 205
*		In(2L+2R)Cy, Cy L sp <sup>2</sup> . . . . . 862, 869
*		In(2L+2R)Cy, Cy pr <sup>2</sup> . . . . . 425, 887, etc.
*		In(2L+2R)Cy, Cy sp <sup>2</sup> . . . . . 305, 687
*		In(2L+2R)Cy, Cy S <sub>2</sub> E(S) <sub>2</sub> . . . . . 771
*		In(2L+2R)Cy, Cy S <sub>2</sub> dp <sup>lv2</sup> E(S) <sub>2</sub> . . . . . 335
*	Ins(2L+2R)Cy, bw <sup>V34k</sup>	In(2L+2R)Cy, Cy In(2R)bw <sup>V34k</sup> . . . . . 342
*	Ins(2L)Cy, (2R)NS	In(2L)Cy, Cy dp <sup>lv2</sup> pr, In(2R)NS. 1 px l(2)NS sp . 361
820	In(2L+2R)NS	In(2L+2R)NS, b mr/In(2L+2R)Cy, Cy

* In(2L+2R)NS	In(2L+2R)NS, px sp . . . . .	738
* Ins(2L)t, (2R)Cy	In(2L)t, Roi In(2R)Cy, bw sp <sup>2</sup> or (= Roi) . . . . .	441

2LR Inversions

821 In(2LR)102 <sub>v1</sub>	In(2LR)102 <sub>v1</sub> ds <sup>W</sup> sp <sup>2</sup> /SM1, al <sup>2</sup> Cy sp <sup>2</sup>	
* In(2LR)bw <sub>v32g</sub>	In(2LR)bw <sub>v32g</sub> , ds <sup>33k</sup> . . . . .	328, 353, etc.
* In(2LR)dp	. . . . .	352, 739
* In(2LR)Gla	. . . . . (= Gla) . . . . .	(see T(2;3)dp)
* In(2LR)Pm <sub>2</sub>	. . . . .	232
* In(2LR)Pm <sub>2</sub>	. . . . .	(see In(2LR)bw <sub>v1</sub> )
* In(2LR)Rev <sup>B</sup>	. . . . . (= Rev) . . . . .	(see In(2LR)bw <sub>v32g</sub> )
* In(2LR)Rev <sup>B</sup>	. . . . .	823
* In(2LR)Rvd	. . . . .	753
* In(2LR)SM1	In(2LR)SM1, al <sup>2</sup> Cy sp <sup>2</sup> . . . . . (= SM1) . . . . .	(see In(2LR)Rev <sup>B</sup> )
* In(2LR)SM5	In(2LR)SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup> . . . . . (= SM5) . . . . .	212, 216, etc.
* In(2LR)U	. . . . . (= U) . . . . .	206 210, etc.
		258

2R Inversions

* In(2R)bw <sub>v34k</sub>	. . . . .	342
822 In(2R)bw <sub>vDel</sub>	In(2R)bw <sub>vDel</sub> , b/b lt 1 cn mi sp	
823 In(2R)bw <sub>vDe2</sub>	In(2R)bw <sub>vDe2</sub> /In(2LR)Rev 1	
* In(2R)Cy <sup>K</sup>	In(2R)Cy, cn Bld . . . . .	354
* In(2R)Mo <sup>K</sup>	. . . . .	868
* In(2R)NS	. . . . .	361

3L Inversions

* In(3L)D <sub>3</sub>	. . . . . (= D <sub>3</sub> ) . . . . .	476, 522, etc.
* In(3L)D <sub>3</sub>	. . . . . (= D <sub>3</sub> ) . . . . .	477, 527
* In(3L)P	In(3L)P, gm . . . . .	559, 623
* In(3L)P	In(3L)P, Me . . . . .	531, 532, etc.
* In(3L)P	In(3L)P, Me ca . . . . .	892, 894
824 In(3L)P	In(3L)P, not-36e/R	

3L + 3R Inversions

* In(3L+3R)LVM	. . . . . (= LVM) . . . . .	506 523, 528
* In(3L+3R)P	In(3L+3R)P, 1(3)PL 1(3)PR . . . . . (= Payne) . . . . .	477, 490, etc.
*	In(3L+3R)P, 1(3)PL 1(3)PR, Dfd ca . . . . .	486, 487, etc.
* Ins(3L)P, (3R)C	In(3L)P, Me, In(3R)C, e 1(3)e . . . . .	626
*	In(3L)P, Me, In(3R)C, Sb e 1(3)e . . . . .	563
* Ins(3L)P, (3R)Pl8	In(3L)P, In(3R)Pl8, Me Ubx le . . . . .	524

3LR Inversions

* In(3LR)Cx	. . . . . (= Cx) . . . . .	479, 498, etc.
* In(3LR)Cx	In(3LR)Cx, D . . . . .	522, 893
* In(3LR)DcxF	. . . . . (= In(3LR)CxF) . . . . .	615, 687

* In(3LR)DcxF	In(3LR)DcxF, ru h ca . . . . . 840
825 In(3LR)sep	In(3LR)sep, sep ri pP
* In(3LR)P35	. . . . . (= In(3LR)Pasadena-35) . . . . . 886
* In(3LR)TM1	In(3LR)TM1, Me ri sbd <sup>1</sup> . . . . . (= TM1) . . . . . 456, 525, etc.
* In(3LR)TM3	In(3LR)TM3, y <sup>+</sup> ac <sup>+</sup> ri p <sup>p</sup> sep bx <sup>34e</sup> e <sup>s</sup> (= TM3) . . . . . 556, 885
* In(3LR)Ubx <sup>101</sup>	. . . . . 579
* In(3LR)Ubx <sup>130</sup>	In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup> e <sup>s</sup> . . . . . (= Ubx <sup>130</sup> ) . . . . . 484, 582, 674, etc.

3R Inversions

826 In(3R)Antp <sup>B</sup>	In(3R)Antp <sup>B</sup> , Antp <sup>B</sup> /TM1, Me ri sbd <sup>1</sup>
* In(3R)C	In(3R)C, cd . . . . . 580
*	In(3R)C, e . . . . . 482, 485, etc.
*	In(3R)C, e l(3)e . . . . . 536, 537, etc.
*	In(3R)C, l(3)a . . . . . 453, 483
*	In(3R)C, Sb e l(3)e . . . . . 488, 514
* In(3R)Cyd <sup>B</sup>	In(3R)Cyd <sup>B</sup> , Cyd <sup>B</sup> . . . . . (= Cyd) . . . . . 489
827 In(3R)Dl <sup>B</sup>	In(3R)Dl <sup>B</sup> , st Dl <sup>B</sup> /In(3R)P <sup>W</sup> , st l(3)W ca
828 In(3R)Hu	In(3R)Hu, Hu Sb <sup>Sp1</sup> /In(3L+3R)P <sup>Xa</sup>
829 In(3R)Mo	In(3R)Mo, sr/T(2;3)ap <sup>Xa</sup> , ca; see also 688
* In(3R)P	. . . . . 481, 511
* In(3R)P18 <sup>F1a</sup>	. . . . . (= In(3R)Pasadena 18) . . . . . 524
830 In(3R)P <sup>F1a</sup>	In(3R)P <sup>F1a</sup> (homozygous)
* In(3R)P <sup>W</sup>	In(3R)P <sup>W</sup> , st l(3)W ca . . . . . 605, 827

Translocations-1;Y

831 T(1;Y)1E	T(1;Y)1E, y/y f:=, cn bw
832 T(1;Y)2E	T(1;Y)2E/v car 1(Stern #64)/y f:=; cn bw

Translocations-1;2

833 T(1;2)Bld <sup>257-15</sup>	T(1;2)Bld <sup>257-15</sup> , Bld/C1B (carries In(2R)Cy)
834 T(1;2)f <sup>257-15</sup>	T(1;2)f <sup>257-15</sup> /In(1)AM
835 T(1;2)lt <sup>264-10</sup>	T(1;2)lt <sup>264-10</sup> /In(2L+2R)Cy, Cy (carries eq and possibly su(s) <sup>3</sup> )
836 T(1;2)N <sup>S2</sup>	T(1;2)N <sup>S2</sup> /FM6, y <sup>31d</sup> sc dm B
837 T(1;2)sc <sup>19</sup>	T(1;2)sc <sup>19</sup> /In(2L+2R)Cy, Cy <sup>19i</sup>
838 T(1;2)sc <sup>19</sup>	T(1;2)sc <sup>19</sup> /y f:=;fs(2)B sc <sup>19i</sup> b pr/In(2L+2R)Cy, Cy dp <sup>lvI</sup> pr

Translocations-1;3

839 T(1;3)263-4	T(1;3)263-4, y sc B <sup>1</sup> /In(1)AM
840 T(1;3)143-3	T(1;3)143-3, ru e <sup>s</sup> ca/In(3LR)DcxF, ru h ca
* T(1;3)Del-143 <sup>264-6</sup>	. . . . . (see T(1;3)143-3)
841 T(1;3)N <sup>264-6</sup>	T(1;3)N <sup>264-6</sup> , y/y w dm (= N)
842 T(1;3)04	T(1;3)04/C1B
843 T(1;3)05	T(1;3)05, D/y f:=
844 T(1;3)OR60	T(1;3)OR60/In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup> e <sup>s</sup> q;tra <sup>D</sup> Sb e/In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup> e <sup>s</sup> ♂
845 T(1;3)ras <sup>v</sup> <sub>J4</sub>	T(1;3)ras <sup>v</sup> /y f:=
* T(1;3)sc <sup>J4</sup>	. . . . . 856

846 T(1;3)sc<sup>z</sup>  
 847 T(1;3)sc<sup>z</sup>260-15  
 848 T(1;3)sta  
 849 T(1;3)sta  
 850 T(1;3)v<sup>3ld</sup>  
 851 T(1;3)w<sup>36d</sup>

T(1;3)sc<sup>z</sup>/y f:=  
 T(1;3)sc<sup>z</sup>260-15/FM6, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 T(1;3)sta/FM3, y<sup>3ld</sup> sc<sup>8</sup> dm B I  
 T(1;3)sta/y f:=  
 T(1;3)v, v/FM6, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 T(1;3)w<sup>36d</sup>, v f/C1B

Translocations-1;4

852 T(1;4)B<sup>S</sup>  
 \* T(1;4)N<sup>8a</sup>  
 853 T(1;4)N<sup>8</sup>264-12  
 854 T(1;4)sc<sup>8</sup>  
 855 T(1;4)w<sup>m5</sup>  
 856 T(1;4)w<sup>m5</sup>(1;3)sc<sup>J4</sup>  
 857 T(1;4)w<sup>m258-18</sup>  
 858 T(1;4)w<sup>m258-21</sup>  
 859 T(1;4)w<sup>VD3</sup>  
 \* T(1;4)w<sup>VD3</sup>

T(1;4)B<sup>S</sup>/y f:=  
 . . . . . 264-12 . . . . . 3ld . 8 . . . . . (see T(1;4)N<sup>264-12</sup>)  
 T(1;4)N<sup>8</sup>/FM6, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 T(1;4)sc<sup>8</sup>, B<sup>a</sup>w<sup>D</sup>/y f:=  
 T(1;4)w<sup>m5</sup>/ey  
 T(1;4)w<sup>m5</sup>;T(1;3)sc<sup>J4</sup>(C1B)  
 T(1;4)w<sup>m258-18</sup>, y/ci<sup>D</sup>  
 T(1;4)w<sup>m258-21</sup>/FM1, y<sup>3ld</sup> sc<sup>8</sup> w<sup>a</sup> lz<sup>s</sup> B  
 T(1;4)w<sup>m258-21</sup>, y w<sup>a</sup>/FM4, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 . . . . . (see T(1;4)w<sup>m258-21</sup>)

Translocations-Y;2

\* T(Y;2)A . . . . . 336  
 860 T(Y;2)B T(Y;2)B/b; see also 422  
 \* T(Y;2)C . . . . . 271, 373  
 \* T(Y;2)E . . . . . 203, 266  
 \* T(Y;2)G . . . . . 238  
 \* T(Y;2)J . . . . . 381  
 861 T(Y;2)rl T(Y;2)rl, lt cn/b lt bw

Translocations-Y;2;3

\* T(Y;2;3)F . . . . . 564

Translocations-2;3

862 T(2;3)101 T(2;3)101, al<sup>2</sup> sp<sup>2</sup>/In(2L+2R)Cy, Cy L<sup>4</sup> sp<sup>2</sup>  
 863 T(2;3)101 T(2;3)101;ru h e<sup>2</sup> ro ca/In(3L+3R)P, Dfd ca<sup>3</sup> L<sup>4</sup> sp<sup>2</sup>  
 864 T(2;3)108 T(2;3)108, al c sp/In(2L+2R)Cy, al<sup>2</sup> Cy lt L<sup>4</sup> sp<sup>2</sup>  
 865 T(2;3)109 T(2;3)109, p<sup>P</sup>/In(3L+3R)P, Dfd ca  
 866 T(2;3)A T(2;3)A, Bl;ru h D TA ss e<sup>S</sup>/In(3L+3R)P  
 \* T(2;3)ap<sup>Xa</sup> . . . . . (= Xa) . . . . . 458, 543, etc.  
 \* T(2;3)ap<sup>Xa</sup> T(2;3)ap<sup>Xa</sup>, ca . . . . . 557, 829  
 867 T(2;3)ap<sup>Xa</sup> T(2;3)ap<sup>Xa</sup>/1(3)XaR  
 868 T(2;3)Ata, T(2;3)Ata, Ata/T(2;3)Mot-K  
 869 T(2;3)B T(2;3)B, al sp<sup>2</sup>/In(2L+2R)Cy, Cy L<sup>4</sup> sp<sup>2</sup>  
 870 T(2;3)B<sup>V4</sup> T(2;3)B;ru h D TB ss e<sup>S</sup>/In(3L+3R)P  
 871 T(2;3)bw<sup>V5</sup> T(2;3)bw<sup>V5</sup>/SM1, al<sup>2</sup> Cy sp<sup>2</sup>  
 872 T(2;3)bw<sup>VDe3</sup> T(2;3)bw<sup>VDe3</sup>/SM5, al<sup>2</sup> Cy lt sp<sup>2</sup>  
 873 T(2;3)bw<sup>VDe3</sup> T(2;3)bw<sup>VDe3</sup>;Ubx bxd/In(3LR)Cx<sup>2</sup>  
 874 T(2;3)bw<sup>VDe4</sup> T(2;3)bw<sup>VDe4</sup>/SM5, al<sup>2</sup> Cy lt sp<sup>2</sup>

875	T(2;3)C	T(2;3)C;ru h D TC ss e <sup>s</sup> /In(3L+3R)P
*	T(2;3)dp <sub>D</sub>	..... 242
876	T(2;3)dp	T(2;3)dp, dp/SM1, al <sup>2</sup> Cy <sub>2</sub> sp <sup>2</sup>
877	T(2;3)E	T(2;3)E/SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp
878	T(2;3)Hm	T(2;3)Hm, Hm/In(2L+2R)Cy. Cy
879	T(2;3)Hn	T(2;3)Hn, Df(3L)Hn, Hn/In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup> e <sup>s</sup>
*	T(2;3)Me	..... 533, 534, etc.
*	T(2;3)P	T(2;3)P, P ..... 675, 692
*	T(2;3)p <sub>Gr</sub>	..... (see T(2;3)Pu <sup>Gr</sup> )
880	T(2;3)Pu <sub>Gr</sub>	T(2;3)Pu <sub>Gr</sub> Pu <sub>Gr</sub> <sup>4</sup> C(3)x
881	T(2;3)Pu <sub>Gr</sub>	T(2;3)Pu <sub>Gr</sub> , Pu <sub>Gr</sub> /SM1, al <sup>2</sup> Cy sp <sup>2</sup>
882	T(2;3)rn	T(2;3)rn/In(2R)Cy
883	T(2;3)Dp-S	T(2;3)Dp-S, ho/In(2L+2R)Cy, Cy E(S) (hom. viable)
884	T(2;3)S <sub>M</sub>	T(2;3)S <sub>M</sub> /In(2L+2R)Cy, Cy E(S)
*	T(2;3)S <sub>V</sub>	..... 440
885	T(2;3)Sb <sub>V</sub>	T(2;3)Sb <sub>V</sub> , Sb <sub>V</sub> , In(3R)Mo/TM3, y <sup>+</sup> ac <sup>+</sup> ri p <sup>+</sup> sep <sup>+</sup> bx <sup>34e</sup> e <sup>s</sup>
886	T(2;3)Sb <sub>V</sub>	T(2;3)Sb <sub>V</sub> , Sb <sub>V</sub> , In(3R)Mo, In(3LR)P35/Sm1, al <sup>2</sup> Cy sp <sup>2</sup> ; In(3LR)Ubx <sup>130</sup> Ubx <sup>130</sup> e <sup>s</sup>
*	T(2;3)Xa	..... (see T(2;3)ap <sup>Xa</sup> )

Translocations-2;4

887	T(2;4)a	T(2;4)a/In(2L+2R)Cy, Cy pr; ey <sup>2</sup>
888	T(2;4)ast <sup>v</sup>	T(2;4)ast <sup>v</sup> /In(2L+2R)Cy, al <sup>2</sup> Cy lt <sup>3</sup> L <sup>4</sup> sp <sup>2</sup>
889	T(2;4)b	T(2;4)b/In(2L+2R)Cy, Cy pr; ey <sup>2</sup>
890	T(2;4)d	T(2;4)d, al dp px sp/In(2L+2R)Cy, Cy pr; ey <sup>2</sup>
891	T(2;4)d	T(2;4)d/In(2L+2R)Cy, Cy pr

Translocations-3;4

892	T(3;4)A2	T(3;4)A2/In(3L)P, Me ca
893	T(3;4)A12	T(3;4)A12/In(3LR)Cx, D
894	T(3;4)A13	T(3;4)A13, ve ca/In(3L)P, Me ca
895	T(3;4)A28	T(3;4)A28, ve ca (homozygous)
896	T(3;4)c	T(3;4)c/In(3L+3R)P, Dfd ca
897	T(3;4)e	T(3;4)e/In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup> e <sup>s</sup>
898	T(3;4)e	T(3;4)e, h th st cu sr e <sup>s</sup> ca/In(3L+3R)P, Dfd ca
899	T(3;4)f	T(3;4)f/In(3L)P, Me
900	T(3;4)f	T(3;4)f, h th st cu sr e <sup>s</sup> ca/In(3L+3R)P, Dfd ca

Transpositions

901	Tp(3)bxd <sup>100</sup>	Tp(3)bxd <sup>100</sup> , ri/T(2;3)Me
902	Tp(3)bxd <sup>107</sup>	Tp(3)bxd <sup>107</sup> , bx bxd <sup>107</sup> sr e <sup>s</sup> /bx <sup>34e</sup> Mc
903	Tp(3)Vno	Tp(3)Vno/H <sup>2</sup>



ROMA, ITALY: CITTA' UNIVERSITARIA  
Istituto di Genetica

Wild Stocks

A 1 Canton-S  
A 2 Marzi  
A 3 Oregon

Chromosome 1

B 1 car bb  
B 2 e(w<sup>e</sup>)w<sup>e</sup>/C(1)DX, y f  
B 3 gt v  
B 4 mal  
B 5 pn  
B 6 r<sup>9</sup>/C(1)DX, y f  
B 7 sc cv v f B/C(1)DX, y f  
B 8 sc z ec  
B 9 sw  
B10 w  
B11 w<sup>a</sup>  
B12 w<sup>Bwx</sup>  
B13 w<sup>cf</sup>  
B14 w<sup>cf</sup>/C(1)DX, y f  
B15 w<sup>ch</sup>  
B16 w<sup>bl</sup>  
B17 y  
B18 y ac sc pn/C(1)DX, y f  
B19 y cv v g/C(1)DX, y f  
B20 y fa<sup>n</sup> sn<sup>3</sup>  
B21 y w  
B22 y w<sup>a</sup> spl rb  
B23 y<sup>2</sup> v mal  
B24 y<sup>2</sup> w<sup>cf</sup>  
B25 y<sup>3ld</sup> v<sup>Of</sup> w<sup>a</sup> f sn/C(1)  
DX, y f  
B26 y l(1)J l<sup>259</sup> w m f/S-5/  
sc<sup>8</sup>.Y

Chromosome 2

C 1 b cn c bw  
C 2 b cn vg bw  
C 3 b pr cn  
C 4 Bl L/Cy Su(Cy)  
C 5 Bl L<sup>2</sup>/SM5, al<sup>2</sup> Cy lt<sup>v</sup>  
cn<sup>2</sup> sp<sup>2</sup>  
C 6 Bl stw<sup>48</sup> ap<sup>blt</sup> tuf sp/  
SM5, al<sup>2</sup> Cy lt<sup>v</sup> cn<sup>2</sup> sp<sup>2</sup>  
C 7 bw  
C 8 bw<sup>D</sup>  
C 9 cn bw  
C10 cn en/SM5, al<sup>2</sup> Cy lt<sup>v</sup>  
cn<sup>2</sup> sp<sup>2</sup>  
C11 cv-2  
C12 ds S G b br/Cy, al<sup>2</sup> lt<sup>3</sup>  
L<sup>4</sup> sp<sup>2</sup>  
C13 Frd/SM5, al<sup>2</sup> Cy lt<sup>v</sup>  
cn<sup>2</sup> sp<sup>2</sup>  
C14 L<sup>2</sup>/SM5, al<sup>2</sup> Cy lt<sup>v</sup> cn<sup>2</sup> sp<sup>2</sup>  
C15 Pin/Cy  
C16 Pin L/Cy  
C17 pr cn  
C18 px bw sp  
C19 S Sp ab<sup>2</sup> ltd/SM5, al<sup>2</sup> Cy  
lt<sup>v</sup> cn<sup>2</sup> sp<sup>2</sup>  
C20 Sp/Cy  
C21 Sp Bl L/Cy  
C22 Sp Bl L Pin/Cy  
C23 Sp J Pin/Cy  
C24 Sp b pr cn L<sup>2</sup> Pin/SM5,  
al<sup>2</sup> Cy lt<sup>v</sup> cn<sup>2</sup> sp<sup>2</sup>  
C25 Tft/Cy

Chromosome 3

D 1 bul  
D 2 ca  
D 3 ca K-pn  
D 4 D<sup>3</sup> H/In(3L)P, Me  
D 5 eg/In(3LR)Cx  
D 6 eg<sup>2</sup>/In(3LR)Cx  
D 7 Gl/In(3LR)bx<sup>130</sup>  
D 8 Gl Sb/LVM  
D 9 Gl Sb/In(3LR)Ubx<sup>130</sup>  
D10 Hn<sup>r-3</sup> sr  
D11 H/Sb sr Ir(3R)Me  
D12 H<sup>2</sup>/Tp(3)Vno  
D13 Ly Sb/LVM  
D14 Ly Sb/In(3LR)Ubx<sup>130</sup>  
D15 Ly st/In(3LR)Ubx<sup>130</sup>  
D16 R Ly/In(3L)P, gm  
D17 ry<sup>2</sup>  
D18 ru h th st cu sr e<sup>s</sup> ca  
("rucuca")  
D19 ru h th st p<sup>p</sup> cu sr e<sup>s</sup>  
("rupes")  
D20 Sb/In(3LR)Ubx<sup>130</sup>  
D21 st  
D22 st c-3-G ca/In(3LR)Ubx<sup>130</sup>  
D23 st Sb/In(3LR)Ubx<sup>130</sup>  
D24 st ve  
D25 th  
D26 th st cp  
D27 th st cp in ri p<sup>p</sup>

Chromosome 4

E 1 ci ey<sup>R</sup>  
E 2 ey  
E 3 spa<sup>pol</sup>

Multichromosomal

F 1 y<sup>2</sup>;bw(1;2)  
F 2 y<sup>2</sup> cho;lys rc(1;2)  
F 3 y;Gl Sb/TM2(1;3)  
F 4 y;spa<sup>pol</sup>(1;4)  
F 5 y pn;C(4)RM, ci ey<sup>R</sup>/O(1;4)  
F 6 al L<sup>4</sup> Cy Sp/bw<sup>V1</sup>;H/Sb sr In(3R)Me(2;3)  
F 7 bw;e(2;3)  
F 8 bw;st(2;3)  
F 9 lys rc;ss(2;3)  
F10 ("sifter O")S Sp P<sup>-</sup> T(2;3), Ins CXF/  
dp<sup>txI</sup> Cy, Ins 05 pr cn<sup>2</sup>;Dl H e p<sup>i</sup>(2;3)  
F11 Sp/Cy;Ly Sb/TM2(2;3)  
F12 Sp/Cy;Gl Sb/TM2(2;3)  
F13 Sp/Cy;spa<sup>pol</sup>(2;4)  
F14 "Basc" In(1)sc<sup>8</sup> S<sup>1</sup>L sc<sup>8</sup>R+S, sc<sup>8</sup>S<sup>1</sup> sc<sup>8</sup> w<sup>a</sup>  
B;Sp/Cy;e(1;2;3)  
F15 y;bw;st(1;2;3)

F16 y B;Sp/Cy;spa<sup>pol</sup>(1;2;4)  
F17 "Basc" In(1)sc<sup>8</sup> S<sup>1</sup>L sc<sup>8</sup>R+S, sc<sup>8</sup>S<sup>1</sup> sc<sup>8</sup> w<sup>a</sup> B;  
SM1/T(2;3)S9 bw e/TM2;spa<sup>pol</sup>(1;2;3;4)  
F18 y;bw;st;spa<sup>pol</sup>(1;2;3;4)  
F19 y;SM1/T(2;3)S9 bw e/TM2;spa<sup>pol</sup>(1;2;3;4)  
F20 y/Y.y<sup>+</sup>;SM1/T(2;3)S9 bw e/TM2;spa<sup>pol</sup>  
(1;2;3;4)  
F21 y B;Sp/Cy;st ve;spa<sup>pol</sup>(1;2;3;4)  
F22 y B;Sp/Cy;st ca;spa<sup>pol</sup>(1;2;3;4)  
F23 y B/Y.y<sup>+</sup>;SM1/T(2;3)S9 bw e/TM2;spa<sup>pol</sup>  
(1;2;3;4)  
F24 y w;SM1/T(2;3)S9 bw e/TM2;spa<sup>pol</sup>(1;2;3;4)

Translocations

G 1 T(1;4)B<sup>S</sup>(16 Al), y<sup>2</sup> cv v B<sup>S</sup> car/C(1)DX,  
y f  
G 2 T(1;4)w<sup>m5</sup>  
G 3 T(Y;2)A/lt

G 4 T(Y;2)B/b  
 G 5 T(Y;2)E/ab<sup>2</sup>  
 G 6 T(Y;2)G/b pr tk  
 G 7 T(Y;2)J/px bw sp  
 G 8 T(Y;2)rl, lt cn/b lt bw  
 G 9 T(2;3)63-3  
 G10 T(2;3)63-18  
 G11 T(2;3)63-23  
 G12 T(2;3)64-33  
 G13 T(2;3)bw<sup>VDe4</sup>/SM5, al<sup>2</sup> Cy lt<sup>v</sup> cn<sup>2</sup> sp<sup>2</sup>  
 G14 T(2;3)A;B1;ru h D TA ss e<sup>S</sup>/In(3L+3R)P  
 G15 T(2;3)B;ru h D TB ss e<sup>S</sup>/In(3L+3R)P  
 G16 T(2;3)dp  
 G17 T(2;4)a/In(2L+2R)Cy, Cy pr;ey<sup>2</sup>  
 G18 T(2;4)ast<sup>v</sup>/In(2L+2R)Cy, al<sup>2</sup> Cy lt<sup>3</sup> L<sup>4</sup> sp<sup>2</sup>  
 G19 T(2;4)b/In(2L+2R)Cy, Cy pr;ey<sup>2</sup>

Inversions

H 1 In(1)65, y f/B<sup>S</sup> Y  
 H 2 In(1)d1-49 Hw m<sup>2</sup>/y w<sup>a</sup> N<sup>614-10</sup>  
 H 3 In(1)d1-49, In(1)B<sup>M1</sup>, y/Y-bb & y v bb/Y-bb  
 H 4 In(1)d1-49, y fa<sup>n</sup>  
 H 5 In(1)d1-49, y Hw m<sup>2</sup> g<sup>4</sup>/Df(1)N<sup>8</sup>  
 H 6 In(1)d1-49, v<sup>Of</sup> f  
 H 7 In(1)EN, y bb/sc<sup>8</sup>.Y  
 H 8 In(1)l-v 231, y l(1)v 231/C(1)RM, y w/sc<sup>8</sup>.Y  
 H 9 In(1)N<sup>264-84</sup>, y/FM6, y<sup>3ld</sup> sc<sup>8</sup> dm B  
 H10 In(1)rst<sup>3</sup>, y rst<sup>3</sup> car bb  
 H11 In(1)sc<sup>4</sup>, sc<sup>8R</sup>, y<sup>sc4+8</sup> cv v f/C(1)DX, y f  
 H12 In(1)sc<sup>4</sup>, y sc<sup>4</sup>  
 H13 In(1)sc<sup>7</sup>, In(1)AM, sc<sup>7</sup>/In(1)d1-49, y Hw m<sup>2</sup> g<sup>4</sup>  
 H14 In(1)sc<sup>8</sup>, sc<sup>8</sup>  
 H15 In(1)sc<sup>8</sup>, y<sup>3ld</sup> sc<sup>8</sup> w<sup>a</sup>  
 H16 ("Basc")In(1)sc<sup>S1L</sup> sc<sup>8R+S</sup>, sc<sup>S1</sup> sc<sup>8</sup> w<sup>a</sup> B  
 H17 ("Binscy")In(1)sc<sup>S1L</sup> sc<sup>8R+d1-49</sup>, y sc<sup>S1</sup> sc<sup>8</sup> v B  
 H18 ("new Binsc")y sc<sup>S1</sup> B In 49 sc<sup>8</sup>/C(1)DX, y f  
 H19 In(1)w<sup>m4</sup>(bb?)  
 H20 In(2R)bw<sup>VDe1</sup>, b/b lt 1 cn mi sp  
 H21 In(2R)bw<sup>VDe2</sup>/In(2LR)Rev 1  
 H22 In(2L)Cy, Cy dp<sup>lv2</sup> b pr/ds<sup>38k</sup>  
 H23 In(2L)Cy<sup>L</sup> t<sup>R</sup>, Su(S)dp<sup>lv2</sup> pr/ds<sup>W</sup>  
 H24 In(2L)NS/J  
 H25 In(2L)t, lt 1 L<sup>4</sup> sp<sup>2</sup>/bw<sup>V1</sup> ds<sup>33k</sup>  
 H26 l(2)a bs<sup>3</sup>, In(2L)t/bw<sup>V1</sup> ds<sup>33k</sup>  
 H27 In(2L+2R)Cy, al<sup>2</sup> E(S)cn<sup>2</sup> sp<sup>2</sup>  
 (does not carry Cy mutant)  
 H28 In(2L+2R)NS, px sp/s<sup>2</sup> Cy pr B1 cn<sup>2</sup> L<sup>4</sup> bw sp  
 H29 In(2LR)102, ds<sup>W</sup> sp<sup>2</sup>/SM1, al<sup>2</sup> Cy cn<sup>2</sup> sp<sup>2</sup>  
 H30 In(2LR)dp/B1 (see T(2;3)dp....G 16)  
 H31 In(2LR)Gla/b Go

H32 Bld, In(2R)Cy/mr<sup>2</sup>  
 H33 In(2R)Mo<sup>K</sup>/T(2;3)Ata, Ata  
 H34 In(3R)C, l(3)a/Bd<sup>G</sup>

Deficiencies and Duplications

I 1 Df(2R)bw<sup>5</sup>, sp<sup>2</sup>/T(2;3)ap<sup>Xa</sup>  
 I 2 Df(2R)bw<sup>VDe2L</sup> Cy<sup>R</sup>/In(2LR)Gla  
 I 3 Df(2R)MS2<sup>8</sup>/SM1, al<sup>2</sup> Cy cn<sup>2</sup> sp<sup>2</sup>  
 I 4 Df(2R)MS2<sup>10</sup>/SM1, al<sup>2</sup> Cy cn<sup>2</sup> sp<sup>2</sup>  
 I 5 Df(2R)MS4/SM1, al<sup>2</sup> Cy cn<sup>2</sup> sp<sup>2</sup>  
 I 6 Df(2)rl<sup>10a</sup> lt cn/bw<sup>V1</sup>, ds<sup>33K</sup>  
 I 7 Df(2R)vg<sup>C</sup>/Rev<sup>B</sup>  
 I 8 M(2)c<sup>33a</sup>/al<sup>2</sup> InMis Cy cn<sup>2</sup> sp<sup>2</sup>  
 I 9 M(2)H<sup>S5</sup>/SM5, al<sup>2</sup> Cy lt<sup>v</sup> cn<sup>2</sup> sp<sup>2</sup>  
 I10 M(2)e<sup>S</sup>/Cy, bw<sup>V34</sup>  
 I11 Dp(1;f)z<sup>9</sup>, Df(1)sc<sup>J4</sup>/C(1)DX, y f  
 I12 Dp(1;1)B<sup>S</sup> RMG, y w<sup>a</sup> B<sup>S</sup>/In(1)sc<sup>S1</sup>, In(1)d1-49, v  
 I13 px pd;Pdr H, Dp(2;3)P/Pdr

X Chromosomes with a Y arm attached

L 1 Y<sup>S</sup> X.Y<sup>L</sup>, In(1)EN, v f B/O;C(4)RM, ci ey<sup>R/O</sup>  
 L 2 Y<sup>S</sup> X.Y<sup>L</sup> Inv(1)EN, Y<sup>S</sup> B f v y.Y<sup>L</sup> y<sup>+</sup>/y v bb/O  
 L 3 Y<sup>S</sup> X.Y<sup>L</sup> Inv(1)EN, Y<sup>S</sup> v cv y.Y<sup>L</sup> y<sup>+</sup>/y<sup>2</sup> su(w<sup>a</sup>)w<sup>a</sup> bb/O  
 L 4 Y<sup>S</sup> X.(P-7), In(1)EN, Y<sup>S</sup> y f/C(1)RM, y v f/Y  
 L 5 Y<sup>S</sup> X.Y<sup>L</sup> Inv(1)EN, y<sup>+</sup> Y<sup>S</sup> y.Y<sup>L</sup> y<sup>+</sup>/y<sup>2</sup> su(w<sup>a</sup>)w<sup>a</sup> bb/O  
 L 6 X.Y<sup>L</sup>(c-2), y cv v f car bb<sup>-</sup>.Y<sup>L</sup>/C(1)RA (ND-27)sc v f--In(1)sc<sup>8</sup>, f v sc<sup>8</sup>.Y"  
 L 7 Y<sup>S</sup> X.(FR1), Y<sup>S</sup> y cv v f./C(1)DX, y f/Y

Attached XY Chromosomes

M 1 X.Y<sup>S</sup>.Y<sup>L</sup>(110-8 Parker), y<sup>2</sup> su(w<sup>a</sup>)w<sup>a</sup> Y<sup>S</sup>.Y<sup>L</sup> y<sup>+</sup>/C(1)RM, y v bb/O  
 M 2 XY<sup>S</sup>.Y<sup>L</sup>(129-16 Parker), y<sup>2</sup> su(w<sup>a</sup>)w<sup>a</sup> Y<sup>S</sup>.Y<sup>L</sup> y<sup>+</sup>/C(1)RM, y v bb/O  
 M 3 Y<sup>S</sup> X.Y<sup>L</sup>, In(1)EN+d1 49, Y<sup>S</sup> car f v y.Y<sup>L</sup>/C(1)RM, y<sup>2</sup> su(w<sup>a</sup>)w<sup>a</sup> bb/O

Compound Chromosomes

N 1 C(1)DX, y f (see B2, B6, B', B14, B18, B19, B25, G1, H11, H18, I11 and L7)  
 N 2 C(1)RA(ND-27)sc v f--In(1)sc<sup>8</sup>, f v sc<sup>8</sup>. (see L6)  
 N 3 C(1)RM, y v f (see L4)  
 N 4 C(1)RM, y w (see H8)  
 N 5 C(2L)Pr, dp;C(2R)P4, dp  
 N 6 C(3L)RM #4, ri;C(3R)RM #4, sr  
 N 7 C(4)RM, ci ey<sup>R</sup> (see F5 and L1)

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Wild Stocks

1 Canton-S	51 y <sup>2</sup> v f
2 Canton-S-C (highly inbred)	52 y <sup>2</sup> w <sup>a</sup> cv sn <sup>55a</sup> v/M-5
3 Cockaponsett Forest, Conn.	53 y <sup>2</sup> w <sup>a</sup> w/y f:=
4 IF-38, Idaho Falls, Idaho	54 w <sup>a</sup> fw <sup>49c</sup> /y f:=

Chromosome 2

5 NB-1, New Britain, Conn.	55 al
6 OZL, New Haven, Conn.	56 al b c sp <sup>2</sup>
7 Oregon-R	57 al dp b pr c px sp
8 Oregon-R (highly inbred)	58 b
9 Oregon-K	59 b cn vg
10 Sevelen	60 b vg
11 Sevelen (highly inbred)	61 bs <sup>2</sup>
12 Swedish-b	62 bw
13 Swedish-b (highly inbred)	63 bw bs <sup>cy</sup>

Chromosome 1

14 B	64 cn
15 bi	65 cn bw
16 bi ct <sup>6</sup> g <sup>2</sup>	66 cn bw Kr/Pm
17 car	67 dp
18 ct <sup>6</sup>	68 dp bw <sup>a</sup>
19 dor/C1B	69 L <sup>2</sup> /Cy sp <sup>2</sup>
20 dor/FM <sub>4</sub> , y <sup>3ld</sup> sc <sup>8</sup> dm B	70 ltd <sup>37b</sup>
21 fa	71 ltd <sup>37b</sup> vg
22 fu/C1B	72 M(2)l <sup>2</sup> /SML, al <sup>2</sup> Cy sp <sup>2</sup>
23 fs(1)N/M-5	73 net al ex ds S ast/SML, al <sup>2</sup> Cy sp <sup>2</sup>
24 g/C1B	74 pr
25 g <sup>50e</sup>	75 rc
26 g <sup>50e</sup> /y f:=	76 sca
27 Hw <sup>49c</sup> /M-5	77 vg
28 l(1)mys/M-5	78 vg c
29 lz <sup>50e</sup>	79 vg tu-bw

Chromosome 3

30 na/FM <sub>3</sub> , y <sup>3ld</sup> sc <sup>8</sup> dm B 1	80 Dfd <sup>r-1</sup>
31 pn <sup>2</sup>	81 e
32 sc ec v g f/C1B	82 e <sup>4</sup> wo ro
33 sc ec cv ct <sup>6</sup> v g f/FM <sub>3</sub> , y <sup>3ld</sup> sc <sup>8</sup> dm B 1	83 ell
34 sc ec v g f	84 e <sup>s</sup>
35 sc <sup>S1</sup> B, In-S, w <sup>a</sup> sc <sup>8</sup> (M-5)	85 Gl Sb/LVM
36 sn <sup>3</sup> /y f:=	86 Ly/D <sup>3</sup>
37 sn <sup>3</sup> v	87 Ly Sb/LVM
38 sn <sup>3</sup> v B	88 ru h st cu sr e <sup>s</sup> ca
39 sn <sup>4</sup> oc ptg <sup>3</sup> /+:=	89 ry
40 v	90 se
41 w	91 se e
42 w m f	92 ss
43 w spl	93 ss <sup>a</sup>
44 w <sup>a</sup>	94 st
45 w <sup>a</sup> v B	95 Ubx <sup>130</sup> e <sup>s</sup> /Xa
46 w <sup>bf</sup>	
47 w <sup>bf</sup> /FM <sub>4</sub> , y <sup>3ld</sup> sc <sup>8</sup> dm B	
48 w <sup>bl</sup>	
49 w <sup>e</sup>	
50 y <sup>2</sup> sc w <sup>a</sup> ec/y f:=	

Chromosome 496 ci ey<sup>R</sup>

97 ey <sup>2</sup>
98 Scn/ey <sup>D</sup>
99 sv <sup>de</sup> /ey <sup>D</sup>
100 sv <sup>n</sup>
101 spaCat/ci <sup>D</sup>

Multichromosomal

102 ct <sup>45e</sup> v;bw;e(1;2;3)
103 g;cn(1;2)
104 v;bw(1;2)
105 v;bw;e(1;2;3)
106 v;bw;e;ey <sup>2</sup> (1;2;3;4)
107 sc <sup>S1</sup> B, In-S, w <sup>a</sup> sc <sup>8</sup> ;In SML, al <sup>2</sup> Cy sp <sup>2</sup> / dp b Pm ds <sup>33k</sup> ;C Sb/Ubx <sup>130</sup> e <sup>s</sup> (1;2;3)
108 v;e(1;3)
109 w;e(1;3)
110 w <sup>a</sup> v;e(1;3)
111 w <sup>e</sup> ;cn(1;2)
112 y w;ant(1;2)
113 y <sup>2</sup> v f;bw(1;2)
114 bw;e(2;3)
115 bw;st(2;3)
116 cn bw;e(2;3)
117 cn;se(2;3)
118 dp;e(2;3)
119 Pm, dp b/Cy sp <sup>2</sup> ;Sb/CxF (ru h ca?)(2;3)
120 pr;ey <sup>2</sup> (2;4)
121 e;ey <sup>2</sup> (3;4)

Closed-X

122 X <sup>C</sup> , y/y f:=
123 In(X <sup>C2</sup> )w <sup>vc</sup> f/dl-49, y w lz <sup>s</sup>

Deficiencies

124 Df(1)g <sup>1</sup> , f B/In(1)AM
125 Df(1)N <sup>8</sup> /dl-49, y Hw m <sup>2</sup> g <sup>4</sup>
126 Df(1)N <sup>45e</sup> /dl-49, y Hw m <sup>2</sup> g <sup>4</sup>
127 Df(1)w <sup>258-11</sup> /dl-49, y Hw m <sup>2</sup> g <sup>4</sup>
128 Df(1)w <sup>258-42</sup> /dl-49, y Hw m <sup>2</sup> g <sup>4</sup>
129 Df(1)w <sup>258-45</sup> /dl-49, y Hw m <sup>2</sup> g <sup>4</sup>
130 Df(1)y sc/M-5(Vogt)

Translocation

131 T(Y;2)C/pr cn
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BERLIN-DAHLEM, GERMANY: INSTITUT FÜR GENETIK DER FREIEN UNIVERSITÄT BERLIN

Wild Stocks			Chromosome 2	
	107	su(s) <sup>2</sup> w <sup>a</sup> cv t		307 st
	108	v		308 Tu
1 Berlin wild B	109	w	201 al dp b pr c px sp	
2 Berlin wild K	110	w <sup>a</sup>	202 b cn vg	Chromosome 4
3 Canton-S	111	w <sup>bf</sup>	203 bw	
4 Oregon-R	112	w <sup>e</sup>	204 vg	401 ar/ey <sup>D</sup>
	113	w sn <sup>3</sup>		402 bt ey <sup>R</sup> sv <sup>n</sup>
Chromosome 1	114	w <sup>co</sup> sn <sup>2</sup>	Chromosome 3	403 ci ey <sup>R</sup>
	115	w <sup>ch</sup> wy		404 ci gvl bt
101 B	116	wy	301 bx <sup>34e</sup>	405 ey <sup>2</sup>
102 car	117	y ac sc pn	302 Dfd <sup>r-L</sup>	
103 cv	118	y cv v f	303 e <sup>ll</sup>	Multichromosomal
104 f	119	y w <sup>llE4</sup>	304 jv se	
105 m	120	z w <sup>llE4</sup>	305 ri	501 su(s) <sup>2</sup> v;bw
106 sc ec ct v g f			306 ru h th st cu sr	502 bw;st
			e <sup>s</sup> ca	503 ci;ss
				504 v <sub>3</sub> ;e

Inversions

601 In(1)Cl, sc l(1)t <sup>2</sup> v sl B/+ (=ClB)	610 In(1)scS1L sc <sup>8R</sup> + S, scS1 sc <sup>8</sup> w <sup>a</sup> B;
602 In(1)dl-49, tyl bb <sup>1</sup> /C(1)RM, y v f car	In(2LR)SM1, al <sup>2</sup> Cy cn <sup>2</sup> sp <sup>2</sup> /In(2LR)bw <sup>V1</sup> ,
603 In(1)sc <sup>8</sup> + dl-49, yS1 v f B	dp b bw <sup>V1</sup> ds <sup>33k</sup> ;In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup>
604 In(1)sc <sup>8</sup> + dl-49, yS1 v f B;bw;	e <sup>s</sup> /C Sb;spa <sup>pol</sup>
e;spa <sup>pol</sup>	
605 In(1)scS1L sc <sup>8R</sup> + S, scS1 sc <sup>8</sup>	Attached-X and -XY
w <sup>a</sup> B (=M-5)	701 C(1)RM, y/+
606 In(2L)Cy/L <sup>2</sup>	702 C(1)RM, y f/+;bw;e;spa <sup>pol</sup>
607 In(2LR)SM5, al <sup>2</sup> Cy lt <sup>v</sup> cn <sup>2</sup> sp <sup>2</sup> /Bl L <sup>2</sup>	703 Y <sup>SX</sup> .Y <sup>L</sup> , In(1)EN + dl-49, Y <sup>S</sup> car f v y.Y <sup>L</sup> /
608 In(3L)D <sup>3</sup> /Ly	C(1)RM, y <sup>2</sup> su(w <sup>a</sup> )w <sup>a</sup> bb/O
609 In(3LR)TM3, y <sup>+</sup> ri p <sup>D</sup> sep Sb bx <sup>34e</sup> e <sup>s</sup>	704 BSY y <sup>+</sup> /y v;bw
Ser/Me <sup>65d</sup> h th	

Selected Stocks

801 Berlin wild, DDT-resistant 1

IOWA CITY, IOWA: UNIVERSITY OF IOWA  
Department of Zoology (Milkman)

cv-2

A variety of polygenic crossveinless stocks.

OXFORD, ENGLAND: UNIVERSITY OF OXFORD  
Department of Biochemistry, Genetics Laboratory

Wild Stocks	Chromosome 1	Chromosome 2	Chromosome 3
Oregon-R	gr	ab	e
Various Swedish and	w	bw	Ns/Antp <sup>B</sup>
Jugoslav strains	w <sup>a</sup>	cn	
	w m f	cn bw	Chromosome 4
	w m f/ClB	dp cn bw	
	M-5	Cy L <sup>4</sup> /Pm	ey <sup>2</sup>
		vg	ey <sup>opt</sup>
		l(2)Me	

SEOUL, KOREA: CHUNGANG UNIVERSITY  
Department of Biology

Wild Stocks

- 1 Canton-S
- 2 Daekwanryung (Korea)
- 3 Damyang (Korea)
- 4 Keuksando-1 (Korea)
- 5 Keuksando-2 (Korea)
- 6 Kwangju-1 (Korea)
- 7 Oregon-B
- 8 Oregon-R-C
- 9 Oregon-S
- 10 Samarkand
- 11 Seoul-1 (Korea)
- 12 Seoul-2 (Korea)
- 13 Seoul-3 (Korea)
- 14 Seoul-4 (Korea)
- 15 Suwon (Korea)
- 16 Swedish-c
- 17 Yangdong (Korea)
- 18 Dangjin (Korea)
- 19 Wonju (Korea)
- 20 Ansung (Korea)

Chromosome 1

- 101 B
- 102 bo
- 103 br
- 104 Bx<sup>3</sup>
- 105 cm
- 106 cc
- 107 fa
- 108 rg
- 109 sc cv v eq
- 110 sc cv v f

- 111 t
- 112 t<sup>2</sup> v f (
- 113 v
- 114 w
- 5 w<sup>a</sup>
- 116 w<sup>bf2</sup>
- 117 w<sup>ch</sup>
- 118 w<sup>col</sup>
- 119 w<sup>e</sup> bb<sup>1</sup>/ClB
- 120 y
- 121 y ac v
- 122 y sc mf<sup>2</sup>
- 123 y<sup>2</sup> cv v f
- 124 Basc/y sc<sup>8</sup> y

Chromosome 2

- 201 a px sp
- 202 ab
- 203 al
- 204 al bc sp<sup>2</sup>
- 207 bw
- 208 bw ba
- 209 Bl/Cy, bw<sup>45a</sup> sp<sup>2</sup> or<sup>45a</sup>
- 210 c
- 211 c l
- 212 cn bw
- 213 Cy/Pm
- 214 ex
- 215 ho
- 216 L
- 217 L<sup>4</sup>
- 219 pr
- 220 vg
- 221 wt

Chromosome 3

- 301 aah
- 302 bul
- 304 cu
- 305 gl
- 306 h
- 308 ra
- 309 ro
- 310 ru
- 311 se
- 312 ss
- 313 st

Chromosome 4

- 401 bt
- 403 ci gvl bt
- 404 ey

Multichromosomal

- 504 Cy/Pm;Sb/Ubx(2;3)
- 505 Cy/Pm;D/Bd(2;3)

Inversions

- 801 Vg<sup>nw</sup> Hia/SM5, al<sup>2</sup> Cy  
lt<sup>L</sup> Sp
- 802 Vg<sup>u</sup>/Roi, bw sp or

Translocations

- 901 T(2;3)Xa/Sb bx<sup>D</sup>

PITTSBURGH, PENNSYLVANIA: UNIVERSITY OF PITTSBURGH  
Department of Biology

Wild Stocks

- Canton-S
- Umgazi River
- Roma
- Pavia
- Bisignano
- Sciolze

Chromosome 1

- B
- Hw<sup>49c</sup>/FM1
- Hw<sup>49c</sup> f<sup>5</sup>/ClB
- Hw<sup>49c</sup> sn<sup>3</sup>/ClB

- w
- y f:= & Hw<sup>49c</sup> sn<sup>3</sup>
- v<sup>36f</sup>
- v
- su<sup>s2</sup>-v-pr v/FM3(bw)
- vf su<sup>w</sup>-f
- "Basc"(sc<sup>s1</sup> B Ins w<sup>a</sup> sc<sup>8</sup>)
- y Hw & RM, sc<sup>8</sup> B w<sup>a</sup> sc<sup>s1</sup>:=
- y Hw In l B<sup>M1</sup> & y f:=
- y

Chromosome 2

- bw

Chromosome 3

- h
- ry
- su<sup>b</sup> pr/In(3R)C, e;(pr)
- su<sup>2</sup>-Hw bx bxd/TM1, Mc, ri(TM1)
- h<sup>1</sup> gs th

Multichromosomal

- bw;st
- bw<sup>5</sup>;st
- vg;e<sup>11</sup>
- Cy 05/PM;Ubx/Sb

MILANO, ITALY: UNIVERSITA' DI MILANO  
Istituto di Genetica

<u>Wild Stocks</u>				<u>Translocations</u>
1 Canton-S	33 b cn vg	65 w;vg		89 T(1;4)B <sup>S</sup> (16A <sub>1</sub> ),
2 Chieti-v	34 blt	66 y;al bw sp		y <sup>2</sup> cv v B <sup>S</sup> car/
3 Crkwenica	35 blt <sup>S</sup>			y f:=
4 Gaino	36 bsp	<u>Inversions on X Chromosome</u>		
5 Jaslo o.c.	37 bw ba	67 ClB/+		<u>Special Stocks</u>
6 Moltrasio	38 c wt px	68 ClB y/yg <sup>4</sup>		90 "sz e" Y <sup>Lc</sup> /X.Y <sup>S</sup> &
7 Oregon-R	39 cn	69 1(1)7/dl-49 y hw m <sup>2</sup> g <sup>4</sup>		y v f.;e
8 Pavia	40 cn c wt px	70 Muller-5		91 "szw" Y <sup>Lc</sup> /X <sup>w</sup> .Y <sup>S</sup>
9 S. Maria	41 dp cl b	71 Muller-5/lozenge		92 y <sup>2</sup> su <sup>wa</sup> w <sup>a</sup> b b/v
10 Sevelen	42 ft			f B.X.Y
11 Suna	43 ll <sup>2</sup>	<u>Inversions on Chromosome 2</u>		
12 Urbana	44 net	72 Cy sp/Pm		<u>Stocks selected for</u>
13 Valdagno	45 S.o.	73 Cy E-S/S		<u>Tumor manifestation</u>
14 Varese	46 so <sup>2</sup> b cn	74 Cy cn <sup>2</sup> bw sp/Gla In LR		
15 Aspra	47 So <sup>C</sup>	75 Gla/spd gt-4		
16 Ponza	48 sp <sup>2</sup> bs <sup>2</sup>			
17 Giannutri	49 al bc sp			
18 S. Antioco				
	<u>Chromosome 3</u>	<u>Inversions on Chromosome 3</u>		
<u>Chromosome 1</u>	50 cp	76 H/Sb sr In(3R)Me		93 tu A1
19 B	51 gl <sup>3</sup>	77 ltr/Sb sr In(3R)Me		94 tu B1
20 N <sup>B</sup> -S	52 mwh	78 Me ca/ru cu ca		95 tu B3
21 ptg <sup>2</sup>	53 mwh se	79 ve h th C3 G Sb Ubx/		96 tu C1
22 sc ec cut y g f	54 mwh ri ss k	st C3 G ca		97 tu C2
23 v	55 ru	80 Florida In(3R)Payne		98 tu C3
24 sd	56 ve	81 Cy L <sup>4</sup> sp/Pm;H/Sb sr In(3R)Me		99 tu C4
25 w <sup>a</sup>	57 obt	82 y w;CyL <sup>4</sup> sp/Pm;H/Sb sr		100 tu C5
26 w <sup>e</sup>	58 th	In(3R)Me		101 tu D
27 y w	59 tx	83 y sc <sup>S1</sup> In-49 sc <sup>8</sup> ;bw st p <sup>P</sup>		102 tu So <sup>C</sup>
28 abw	60 th tx	84 bsp/bsp;Sb Me/H		103 tu Aspra
29 wmf	61 h	<u>Deficiencies</u>		104 Freckled/Curly
	62 c	85 Df(2)Px <sup>2</sup> Df(2)Px, bw sp/SM1,		105 Frd/Cy L.
	63 se cp e	al Cy sp <sup>2</sup>		106 yw;Cy L/Frd;Sb
<u>Chromosome 2</u>	<u>Multichromosomal</u>	86 Df(2)bw <sup>5</sup> Df(2)bw <sup>5</sup> sp <sup>2</sup> /Xa		Me/H
30 b cn	64 px <sup>43j</sup> co;ru	87 Minute(2)Bridges		107 yw;Cy L/Pm(Frd);
31 sp a px	jv se st ca	88 M(2)33a/al <sup>2</sup> In Mis Cy		Sb Me/H
32 ab		cn <sup>2</sup> sp <sup>2</sup>		108 q 156 melanotic
				109 e 144 melanotic
				110 lm
				111 lnd

CLAYTON, QLD., AUSTRALIA: MONASH UNIVERSITY  
Department of Genetics and Psychology

<u>Wild Stocks</u>	<u>Chromosome 1</u>	<u>Chromosome 2</u>	<u>Chromosome 3</u>
Oregon K	B	bw	e <sup>11</sup>
Oregon R	w	cn	e
Riverside	y	vg	se
East African	w, y	cn bw	ve
Bermuda	t	pd	ry
Hikone	mal	b pr vg	
Canton S	v	Cy Bl L	
		pr	
		pym/cy	
		dy stw bw	



LIMBE, MALAWI: UNIVERSITY OF MALAWI  
Department of Biology, Genetics Section

<u>Wild Stocks</u>	sn	dp	<u>Chromosome 4</u>
	v	sca	
São Paulo	w	vg	ci <sup>57g</sup>
Limbe	w cv sn		ci <sup>w</sup>
	w:= and +	<u>Chromosome 3</u>	spa <sup>pol</sup>
<u>Chromosome 1</u>	<u>Chromosome 2</u>	e	<u>Multichromosomal</u>
B		Ly/D <sup>3</sup>	
cv	bw <sup>60g</sup>	ss <sup>a</sup>	Cy/Pm;D/Sb
1z <sup>cl</sup> /Muller-5	cn bw	ve	y;bw;st
m			Sm5/Bl <sup>a</sup> ;Tm3/Sb

SOUTH ORANGE, NEW JERSEY: SETON HALL UNIVERSITY  
Department of Biology

<u>Wild Stocks</u>	<u>Chromosome 1</u>	w <sup>a</sup>	<u>Chromosome 2</u>
		w <sup>e</sup>	
Oregon-R	B	w B	b
Canton-S	Basc	w f	b vg
Urbana-S	f	w m f	bw
Swedish-c	g <sup>2</sup>	y	
3 South Orange Strains	m	y cv v f	<u>Chromosome 3</u>
	m <sup>DB</sup> (homozygous)	y B & y f:=	
	ras <sup>2</sup>		e
	w		ru h th st cu sr e <sup>s</sup> ca
			st

CLEVELAND, OHIO: CLEVELAND STATE UNIVERSITY  
Department of Biology

<u>Wild Stocks</u>	s	J <sup>34e</sup>	U/cg C	rsd
	f	rk <sup>4</sup>	Px <sup>2</sup> bw sp/SM1 al <sup>2</sup>	ca
Oregon-R	Bx <sup>3</sup>	b	Cy sp <sup>2</sup>	bv
Lausanne-S	car	Coi	Bl L <sup>2</sup> /Cy dp <sup>2</sup>	ca K-pn
	y <sup>2</sup> w <sup>a</sup> ct <sup>6</sup>	hk	b Go/Gla	ve h th
<u>Chromosome 1</u>	y cv v f	Bl/esc	Pu <sup>2</sup> /SM1, al <sup>2</sup> Cy sp <sup>2</sup>	se ss k e <sup>s</sup> ro
	w m f	Alu	Ruf/ds <sup>38k</sup> Pm	D/G1
y	fa fa <sup>no</sup> sn	stw		G1 Sb H/Payne
br	ec dx	ad	<u>Chromosome 3</u>	Bd <sup>8</sup> /In3R C, 1(3)a
pn <sup>2</sup>	cv f	L <sup>2</sup>		Mc/ap <sup>Xa</sup>
w	f B	L <sup>2</sup>	ve	
w <sup>sat</sup>	dor/C1B	c	se	<u>Chromosome 4</u>
w <sup>co</sup>	N <sup>8</sup> /y Hw In49 m <sup>2</sup> g <sup>4</sup>	bw <sup>2b</sup>	eyg	
w <sup>Bwx</sup>	y w sn <sup>3</sup> B & y f:=	Pin	app	spa <sup>pol</sup>
Ax	sc ct <sup>6</sup> car/y f:=	Hx	th	bt
fa	sc <sup>51</sup> B InS w <sup>a</sup> sc <sup>8</sup>	al b cv sp	st	ey <sup>2</sup>
fa <sup>no</sup>	("Basc")	al dp b pr	cp	ar/ey <sup>D</sup>
ec		b cn bw	w	
bo	<u>Chromosome 2</u>	b vg	drb	<u>Multichromosomal</u>
cx		Cy/Pm	bx <sup>34e</sup>	
cm	net	Cy/Pm; st	Cbx	cn;st
sn <sup>3</sup>	al	dp <sup>02</sup> dp <sup>1v1</sup> b/Cy	ell	b;ell
v	ho	Bl L	cd	bw;e
dy	dp	ds <sup>38k</sup> /dy(2L)dp <sup>2</sup>	bar-3	
		b pr		

HONOLULU, HAWAII: UNIVERSITY OF HAWAII  
Department of Genetics

Wild Stocks

1 Oregon-R  
2 Canton-S

Chromosome 1

3 B  
4 ec  
5 m  
6 sc cv v f

7 v  
8 w  
9 w<sup>a</sup>  
10 y  
11 y w spl

Chromosome 2

12 dp cn bw  
13 b  
14 b vg

15 Bl/In(2L+2R)Cy,  
Cy bw<sup>45a</sup> sp<sup>2</sup> or<sup>45a</sup>  
16 L<sup>2</sup>  
17 vg

Chromosome 3

18 cu  
19 e<sup>s</sup>  
20 gl<sup>1</sup>  
21 p<sup>p</sup> cu

Chromosome 4

22 ey

Others

23 bw;st  
24 f BB/y f:-  
25 70 2nd & 3rd chromosomal lethals from Korea

OSAKA, JAPAN: OSAKA UNIVERSITY  
Medical School, Department of Genetics

Wild Stocks

1 Canton-S  
2 Oregon-R (iso)  
3 Hikone

Chromosome 1

101 B  
102 car  
103 cm  
104 Eag  
105 f  
106 Hk<sup>1</sup>  
107 Iz<sup>50e</sup>  
108 Muller-5  
109 phenylfulthyroless-1  
110 rb  
111 sn<sup>3</sup>  
112 Sk<sup>5</sup>  
113 v  
114 w  
115 w<sup>a</sup>  
116 y  
117 y<sup>2</sup>  
118 y<sup>34c</sup>  
119 y<sup>2s</sup>  
120 y<sup>v2</sup>  
121 y w f  
122 y w m f

Chromosome 2

201 b  
202 b gp  
203 b vg  
204 bw  
205 cl

206 cn  
207 cn bw  
208 cn Cy/Pm  
209 cn fes(K)bw/Cy  
210 cn vg bw  
211 l(2)gl cn bw/Cy  
212 px  
213 ry<sup>1</sup>  
214 ry<sup>2</sup>  
215 vg

Chromosome 3

301 bar<sup>-3</sup>  
302 ca<sup>2</sup>  
303 e<sup>11</sup>  
304 jv  
305 ld  
306 se  
307 ss<sup>a</sup>  
318 st  
309 tx<sup>52j</sup>

Chromosome 4

401 pol<sup>1</sup>  
402 sv<sup>2</sup>

Multichromosomal

1:2  
501 v;bw  
502 v;cn  
1:3  
503 y;e<sup>11</sup>

2:3  
504 albas p rucuca  
505 b;se  
506 bw;st ss  
507 cl<sup>57j</sup>;ss<sup>a</sup>  
508 cn;st  
509 vg;e<sup>11</sup>

2:3;4  
510 bw;st;sv<sup>n</sup>  
511 cn;ss;gvl

Attached X

601 yy:+ (Oregon-R)

Special Stocks(A) Insecticide-resistant

701 Hikone-R (multiple)(Japan)  
702 HL2-Q (multiple)(USA)  
703 KSL (multiple)(Sweden)  
704 TG-57 (multiple)(Korea)  
705 WMB (multiple)(Japan)

(B) Amylase

801 Amy<sup>1</sup> (Hikone)  
802 Amy<sup>1</sup> (Oregon-R)  
803 cn Amy<sup>1</sup> bw  
804 Amy<sup>1.3</sup> (L<sup>2</sup>)  
805 cn L<sup>2</sup> Amy<sup>1.3</sup> bw  
806 L<sup>2</sup> Amy<sup>1.3</sup>  
807 Amy<sup>1.6</sup> (Suyama)  
808 Amy<sup>2.6</sup> (Hikone)  
809 cn Amy<sup>2.6</sup> bw  
810 Amy<sup>3.6</sup> (Kyoto)  
811 Amy<sup>4.6</sup> (ad<sup>60</sup>)

UPTON, NEW YORK: BROOKHAVEN NATIONAL LABORATORYWild Stocks

W-1 Canton-S  
W-2 Oregon-R

X Chromosome

X-1  $pn^2$   
X-2 w  
X-3 y cv v f car  
X-4 y v

Chromosome 2

2-1 b vg  
2-2 bw  
2-3 dp

Chromosome 3

3-1 e

Chromosome 4

4-1  $spa^{Cat}/ci^D$

Multichromosomal Stocks

X,3-1 C(1)RM, y f/Y;ca K-pn  
2,3-1 bw;e  
2,3-2 In(2LR)SM1,  $al^2$  Cy  $cn^2$   $sp^2$ /In(2LR)bw<sup>V1</sup>, dp b bw<sup>V1</sup> ds<sup>33k</sup>;In(3R)C,  
Sb/In(3LR)Ubx<sup>130</sup>, Ubx<sup>130</sup> e

Inverted Chromosomes

INX-1 In(1)FM6, y<sup>3ld</sup>  $sc^8$  w<sup>FM6</sup>  $dm^+$  B  
INX-2 In(1)sc<sup>4L</sup>  $sc^8R$  + S, y  $sc^4$   $sc^8$  w<sup>a</sup> B/C(1)RM, y<sup>2</sup> su<sup>(w<sup>a</sup>)</sup> w<sup>a</sup> bb/y<sup>+</sup>Y  
INX-3 In(1)sc<sup>8L</sup>  $sc^8R$  + dl-49,  $sc^1y^8$   $sc^a$  v f B/y l(1)Jl<sup>259</sup> w m f/y<sup>+</sup>Y  
INX-4 In(1)sc<sup>S1L</sup>  $sc^8R$  + S,  $sc^1$   $sc^a$  w<sup>a</sup> B

Attached XY

XY-1 Y<sup>S</sup>X.Y<sup>L</sup>, In(1)EN, ptg oc sn<sup>5</sup>/C(1)RM,  $sc$  ct<sup>n</sup> oc ptg.In(1)dl-49, car  
sn<sup>X2</sup> y

Y Derivatives

Y-1 y<sup>+</sup>Y/In(1)dl-49, y  $sc^{S1}$  B v f/C(1)RM, y f

NORWICH, ENGLAND: JOHN INNES INSTITUTEWild Stocks

1 Bayfordbury  
2 Hampton Hill  
3 Oregon-K  
4 Samarkand  
5 Teddington

Inbred Lines

6 Bayfordbury (A)  
7 Bayfordbury (B)  
8 Oregon (v marker)  
9 b pr

Chromosome 1

10 v  
11 w  
12 y w

Chromosome 2

13 b pr

Chromosome 3

14 b pr vg  
15 bw  
16 cn  
17 dp b cn bw  
18 vg

Chromosome 3

19 st

Multichromosomal

20 Cy L<sup>4</sup>/Pm;H/Sb  
21 bw;e

Inversions

22 Muller-5

PORTO ALEGRE, BRAZIL: UNIVERSIDADE FEDERAL DO RIO GRANDE DO SUL  
Instituto de Biociências, Departamento de Genética

Wild Stocks

Buenos Aires  
Oregon  
Leningrado  
Brisbane  
Eldorado (Rio Grande  
do Sul, Brazil)  
Some inbred strains

Chromosome 1

$pn^2$   
w  
w<sup>e</sup>  
w<sup>bl</sup>  
w<sup>h</sup>  
ras<sup>2</sup>

f  
sc cv v f  
y v  
y

Chromosome 2

pr

ltd  
cn  
bw  
pd  
vg  
L  
st bw

Chromosome 3

p  
e  
se  
e se  
p v

SYDNEY, AUSTRALIA: UNIVERSITY OF SYDNEY  
Department of Animal Husbandry

<u>Wild Stocks</u>	y w sc ec cv ct <sup>6</sup> vg <sup>2</sup> f/ FM3, y <sup>3ld</sup> sc <sup>8</sup> dm B 1 e <sup>11</sup>	<u>Chromosome 3</u>
4 strains from N.S.W. and Victoria		
<u>Chromosome 1</u>	<u>Chromosome 2</u>	<u>Multichromosomal</u>
In rst <sup>3</sup>	b j	In(2L+2R)Cy, Cy bw <sup>45a</sup> sp <sup>2</sup> or <sup>45a</sup> ; In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup> /T(2;3)ap <sup>Xa</sup>
w	net	In(1)sc <sup>S1L</sup> sc <sup>8R+S</sup> , sc <sup>S1</sup> sc <sup>8</sup> wa <sup>a</sup> B; In(2L+2R)Cy, Cy bw <sup>45a</sup> sp <sup>2</sup> or <sup>45a</sup> ; In(3LR)Ubx <sup>130</sup> , Ubx <sup>130</sup> / T(2;3)ap <sup>Xa</sup>
w <sup>bl</sup>	vg	
y	Cy/Pm	

BRNO, CZECHOSLOVAKIA: J.E. PURKYNĚ UNIVERSITY  
Department of Genetics

<u>Wild Stocks</u>	12 y v 13 Muller-5	<u>Chromosome 3</u>	<u>Multichromosomal</u>
1 Oregon K (inbred)		25 se	35 w;e
2 Hikone R (inbred)	<u>Chromosome 2</u>	26 e	36 w <sup>a</sup> ;e
3 Suchumi (inbred)		27 se e	37 b;se
4 Moravec (inbred)	14 dp	28 ru cu ca	38 Cy L/Pm;H/Sb
5 Krnov 65 (inbred)	15 cn	29 Gl Sb/LVM	39 Cy/Pm;Ly/D <sup>3</sup>
6 Krnov 66 (inbred)	16 bw	30 Me Sb e/He	40 Cy/Pm;D/Sb
7 Moskva (inbred)	17 cn vg/Oregon K	31 Gl/Ubx <sup>130</sup>	41 Pm dp b/Cy sp <sup>2</sup> ;Sb/D
	18 cn vg/Suchumi	32 Sb/Ubx <sup>130</sup>	Cx F
<u>Chromosome 1</u>	19 b cn vg	<u>Chromosome 4</u>	42 Sb Ubx/T(2;3)Xa
8 y	20 dp b cn bw		43 Cy/Pm;H/C Sb
9 w	21 al b c sp	33 pol	44 y w;Cy L <sup>4</sup> sp/Pm;H/Sb
10 v	22 al dp b pr c px sp	34 ey <sup>2</sup>	sr, In(3R)Me
11 B	23 Cy/B1 L		
	24 Cy/al b pr lt ltd cn a px pd bw		

BERKELEY, CALIFORNIA: UNIVERSITY OF CALIFORNIA  
Department of Zoology

<u>Wild Stocks</u>	156 y sc m f <sup>5</sup> 159 y sn <sup>3</sup> 160 y w 170 FM7, In(1)sc <sup>8</sup> , dl-49, y <sup>3ld</sup> sc <sup>8</sup> wa v <sup>0f</sup> B 172 FM7-lz/sc <sup>10-1</sup>	324 ss <sup>a</sup> 353 h H/h hp 354 st
3 Samarkand		
5 + <sup>3</sup>		<u>Chromosome 4</u>
<u>Chromosome 1</u>		408 ci ey <sup>R</sup> 431 ci ey x ci ey/+
101 "Basc" (In(1)sc <sup>S1L</sup> sc <sup>8R</sup> + S, sc <sup>S1</sup> sc <sup>8</sup> wa <sup>a</sup> B)	<u>Chromosome 2</u>	<u>Multichromosomal</u>
105 cm ct <sup>6</sup> sn <sup>3</sup>	212 bw	504 y ac;esc <sup>D</sup>
108 f <sup>36a</sup> car/y f	215 cg c/U	508 y;mwh
109 Hw <sup>49c</sup> /FM1, y <sup>3ld</sup> sc <sup>8</sup> wa lz <sup>S</sup> B	232 vg	523 b;Msc/+
116 sc ec cv ct <sup>6</sup> v g <sup>2</sup> f/ FM3, y <sup>3ld</sup> sc <sup>8</sup> dm B 1	<u>Chromosome 3</u>	<u>Triploid</u>
142 y ac sn <sup>3</sup> v	310 H/In(3)hp	
144 y ac sn <sup>3</sup> sx yb <sup>2</sup> sv/ v sc <sup>S1</sup> B In-49 v wa sc <sup>8</sup>	314 mwh e 320 se h	559 ec rb cv/FM6, y <sup>3ld</sup> sc <sup>8</sup> dm B

BOGOTÁ, COLOMBIA: UNIVERSIDAD DE LOS ANDES  
Instituto de Genética

Fusa XXI	Vermellón	Vestigial
Caracolico X	Ebony	Dumpy
Orito (Putumayo)	Sepia	Cy L/Pm
Fusa XXV B	White	

BHAGALPUR, INDIA: BHAGALPUR UNIVERSITY  
Department of Zoology, Drosophila Laboratory

<u>Chromosome 1</u>	<u>Chromosome 3</u>
White eye	Se Cu
	Se Vg

NAPLES, ITALY: UNIVERSITY OF NAPLES  
Institute of General Biology and Genetics & Institute of Zoology

<u>Wild Stocks</u>	Muller-5	b	cn vg bw	se
	v	b cn	Cy/B1 L <sup>2</sup>	se h
Oregon-R	w	b cn bw	Cy/Pin	se Sb
	w <sup>a</sup>	b cn vg		st
<u>Chromosome 1</u>	w cv mal	b cn vg bw	<u>Chromosome 3</u>	
	y	bw		<u>Multichromosomal</u>
cv		bw <sup>D</sup>	cd	
cv mal	<u>Chromosome 2</u>	cl	Sb	cv mal;se
ct		cn	Sb/In(3R)Na	w;vg
mal	awu <sup>2</sup>		<del>xx</del>	

SAPPORO, JAPAN: HOKKAIDO UNIVERSITY  
Faculty of Science, Zoological Institute

<u>Wild Stocks</u>	<u>Chromosome 1</u>	<u>Chromosome 2</u>
Canton S	y w m f	Cy
Sapporo, Hokkaido, Japan	w	v vg
Otoinepp, Hokkaido, Japan		vg;se(2;3)

STOCKHOLM, SWEDEN: UNIVERSITY OF STOCKHOLM  
Institute of Genetics

See DIS 44 (1969): 19-20 and revision in DIS 46 (1971): 33.

KENSINGTON, N.S.W., AUSTRALIA: THE UNIVERSITY OF NEW SOUTH WALES  
School of Wool and Pastoral Sciences

<u>Wild Stocks</u>	<u>Chromosome 1</u>	<u>Chromosome 2</u>	<u>Multichromosomal</u>
Oregon R C	B	dp b j fr	w;vg;e
7 Eastern Australian strains	y cv v f		

SAN BERNARDINO, CALIFORNIA: CALIFORNIA STATE COLLEGE  
Natural Sciences Division

<u>Wild Stocks</u>	<u>Chromosome 2</u>	12 se h
1 Oregon-R-C	6 al b c sp <sup>2</sup>	13 st
<u>Chromosome 1</u>	7 cn bw	<u>Chromosome 4</u>
	8 vg <sup>D</sup> /SM5, al <sup>2</sup> Cy lt <sup>v</sup> sp <sup>2</sup>	14 ci <sup>D</sup> /ey <sup>D</sup>
2 br w <sup>e</sup> ec rb t <sup>4</sup> /Fm1, y <sup>3ld</sup> sc <sup>8</sup> w <sup>a</sup> lz <sup>s</sup> B	<u>Chromosome 3</u>	15 bw <sup>V1</sup> , ds <sup>33k</sup> /In(2L+2R)Cy, Cy;H/In(3R)Mo, sr (2;3)
3 f BB/y f:=	9 Ly/D <sup>3</sup>	16 T(2;3)Hn T(2;3)Hn, Df(3L)
4 g <sup>2</sup>	10 Ly Sb/LVM	Hn, Hn/In(3LR)Ubx <sup>130</sup> ,
5 w	11 red	Ubx <sup>130</sup> es

ASSUIT, U.A.R.: UNIVERSITY OF ASSUIT  
Department of Genetics

<u>Wild Stocks</u>	<u>Chromosome 1</u>	<u>Chromosome 2</u>	<u>Chromosome 3</u>	<u>Multichromosomal</u>
ORK	w	L	st	dp;e
Nobareya	w <sup>a</sup>	bw	e	bw;st
Ghorayeb	B	vg	PP	M-5;Cy/L
Mattmar	y	Cy		
	Sc <sup>81</sup> B Fn S w <sup>a</sup> sc <sup>8</sup>	dp		
		Cy/L		

LONDON, ENGLAND: St. BARTHOLOMEW'S HOSPITAL MEDICAL COLLEGE  
Zoology Department, Genetics Laboratory

<u>Wild Stocks</u>	<u>Chromosome 1</u>
(a) mass mated	(balanced with Inscy)
Kaduna	dm mgt sld
Oregon-K	fin pun slm
	gt w <sup>a</sup> rst <sup>2</sup> ty g <sup>2</sup>
(b) inbred by brother-sister mating	lf sla
Kaduna	
Oregon-K	
	M-5

UTICA, NEW YORK: MASONIC MEDICAL RESEARCH LABORATORY  
Aging Program

<u>Wild Stocks</u>	<u>Chromosome 1</u>
Oregon-R	w
Swedish-c	

MEXICO CITY, MEXICO: NATIONAL COMMISSION OF NUCLEAR ENERGY  
Genetics and Radiobiology Program

Same as DIS #46: 36 except: Delete a3, f3, f4, j2, j8, j17 and m3  
 Correct h8 to read: st c3G en In(3LR)Ubx<sup>130</sup> Ubx<sup>130</sup>,es

PADOVA, ITALY: UNIVERSITA' DEGLI STUDI DI PADOVA  
Istituto di Biologia Animale

<u>Wild Stock</u>	<u>Chromosome 1</u>	<u>Chromosome 2</u>	<u>Chromosome 3</u>
1 Varese	2 sc ec ct v gt f	8 b cn vg	12 ru b ss p <sup>P</sup> st e <sup>S</sup>
	3 v	9 cn	13 se
	4 w <sup>a</sup>	10 dp cl b	
	5 w <sup>bl</sup>	11 net	<u>Inversion on 2</u>
	6 w <sup>e</sup>		
	7 y w		14 Cy sp/Pm

TÜBINGEN, GERMANY: UNIVERSITY OF TÜBINGEN  
Department of Genetics

<u>Wild Stocks</u>	<u>Chromosome 1</u>	<u>Chromosome 2</u>	<u>Chromosome 3</u>
Oregon	Muller 5	vg bw	L Cy/Pm
Wien 1965	ClB/v ptg oc sn		L/Cy
Wien 1966	y v f		
Ponza III			

MELANOGASTER - NEW MUTANTS

Report of D.E. Jeffery

The following chromosomal rearrangements were all X-ray induced. All involve rearrangements of the hairy<sup>+</sup> allele, and all exhibit position effect when in heterozygous condition with hairy.

In(3L)h<sup>+</sup>38: Inversion (3L) hairy<sup>+</sup> Jeffery 67a9. Breakpoints 66D,80F.

T(2;3)h<sup>+</sup>40: Translocation (2;3) hairy<sup>+</sup> Jeffery 67a12. In addition to the reciprocal exchange of the distal ends of 2L and 3L, 3L has section 66D-73C inverted. Breakpoints: 30B, 66D, 73C. New Order: 40-30C/66D-61; 80-73C/66E-73B/30B-21.

T(3;4)h<sup>+</sup>44: Insertional translocation of 63A-66F of 3 into 101F of 4 Jeffery 67b8. Breakpoints 63A, 66F, 101F.

T(2;3)h<sup>+</sup>47: double translocation between 2L and 3L, and 2L and 3R Jeffery 67b10. Breakpoints 23A,34C,66D,98F. New Order: 40-34C/98A-100; 80-66D/23A-21; 81-98F/34C-23A/66D-61.

Report of I.R. Franklin and W. Rumball

Mdh-NADP<sup>0.90</sup>: NADP-dependent malic dehydrogenase, slow form

Mdh-NADP<sup>1.0</sup>: NADP-dependent malic dehydrogenase, intermediate form

Mdh-NADP<sup>1.1</sup>: NADP-dependent malic dehydrogenase, fast form 3-53.1 Three variants of a NADP-dependent malic dehydrogenase have been found in natural populations from New South Wales. Mdh-NADP<sup>1.0</sup> is the most common allele, with a frequency usually exceeding 95%. Variants were identified by polyacrylamide gel electrophoresis in a continuous 0.1M tris-borate buffer, pH 8.9. Each gel was stained in 50 ml of 0.2M tris-HCl buffer, pH 8.5 with 3.5 mg NADP, 12.5 mg nitro blue tetrazolium, 1.0 mg phenazine methosulphate and 20 mg sodium hydrogen malate. The locus was located, in test crosses to 'rucuca', between curled (3-50.0) and stripe (e.62.0). Eighty-six recombinants between these two loci were tested, and in sixty-four cases the Mdh-NADP alleles were associated with curled. The Mdh-NADP locus is therefore located at 3-53.1 ± 0.6.

Report of I.R. Franklin and G.K. ChewTo-1<sup>F</sup>: Tetrazolium oxidase, fast form

To-1<sup>S</sup>: Tetrazolium oxidase, slow form 3-33.3 Several unstained regions are apparent on gels stained for dehydrogenases with tetrazolium salts. One of these oxidases has been found to be polymorphic in several natural populations of *D. melanogaster*. The frequency of the rarer slow allele has been observed at frequencies between 5-10%. The multiply marked third chromosome stock 'rucuca' is homozygous for the slow allele. Heterozygotes are characterized by three bands. To-1 maps between hairy and Esterase-6, 2.6 centiMorgans from Est-6.

Report of M. Carfagna and I. Melon

awu<sup>2</sup>: augenwulst Spontaneous in a wild stock. The eye's phenotype corresponds exactly to that of lost awu described by Volkart, 1959, DIS 33: 100. Expression variable, often asymmetrical. Complete penetrance in homozygote; incomplete, about 22%, in heterozygote, which always shows minor expression. Viability is good to excellent. The location is not exactly corresponding to  $2-56.8 \pm 1.4$  as indicated by Volkart for awu. Extensive recombination studies gave the location  $2-53.7 \pm 0.7$ . It has not been studied cytologically. RK3.

Report of D.J. Fox and K. Madhavan

Hexokinase-3 map position reported as 2-79± in DIS 46: 42 should read 2-73±.

## DROSOPHILA SPECIES - STOCKS

POUGHKEEPSIE, NEW YORK: MARIST COLLEGE  
Department of Biology

D. pseudoobscura

Payson, Arizona (3 strains)  
 Pine Creek, Ariz. (3 strains)  
 Baker Butte, Ariz. (3 strains)  
 Flagstaff, Ariz. (1 strain)  
 Lake Mary, Ariz. (3 strains)  
 Grand Canyon, N.Rim, Ariz. (3 strains)  
 Prescott, Ariz. (4 strains)  
 Sierra Ancha Mtns., Ariz. (1 strain)  
 Portal, Ariz. (1 strain)  
 Crystal Lake, Calif. (3 strains)  
 Sequoia Nat. Pk., Calif. (3 strains)  
 Yosemite Nat. Pk., Calif. (3 strains)  
 Nederland, Colo. (1 strain)  
 Black Canyon, Colo. (1 strain)  
 Custer, S. Dakota (3 strains)  
 Logan, Utah (1 strain)

D. persimilis

Crystal Lake, Calif. (1 strain)  
 Sequoia Nat. Pk., Calif. (2 strains)  
 Yosemite Nat. Pk., Calif. (3 strains)

D. busckii

Princeton, N.J. (1 strain)

D. hydei

Poughkeepsie, N.Y. (1 strain)

D. robusta

Princeton, N.J. (1 strain)  
 Poughkeepsie, N.Y. (1 strain)

D. immigrans

Poughkeepsie, N.Y. (2 strains)

D. affinis

Poughkeepsie, N.Y. (2 strains)

D. melanogaster

Princeton, N.J. (1 strain)  
 Poughkeepsie, N.Y. (4 strains)



GENEVA, SWITZERLAND: UNIVERSITY OF GENEVA  
Department of Genetics

D. hydei Chromosomes are numbered according to Berendes' map. The correlation with Spencer's linkage groups, and with chromosome elements, is as follows:

Element	A	B	C	D	E	F
D. melanogaster chromosome	1=X	2 L	2 R	3 L	3 R	4
D. hydei chromosome						
Spencer 1949	X	IV	III	V	II	VI
Berendes 1963	1=X	3	5	4	2	6

Mutant names and symbols are the same as in D. melanogaster only where actual homology is well established. Different names were chosen deliberately in most other instances. In the case of eye color in particular, ommochrome-deficient mutants are listed as "red A, B, etc." according to their localization on chromosome elements A-F, different loci on the same element as "red A-1, -2, etc.", and similarly drospterin-deficient mutants as "bn A, etc." Homology of the cinnabar mutant has been established, but it is not certain which of the present loci is cn.

Abbreviations: loc = localization; rec = received; ref = reference; syn = synonymous. (For descriptions of these mutants see Drosophila Species - New Mutants, this DIS).

Wild Stocks

W-1 São Paulo 56  
W-2 Leiden 65  
W-3 Alicante 67  
W-4 Madeira 68  
W-5 Zürich 70

C-7 bed pb redC-1Sp sca  
C-8 vg pb redC-1Sp sca  
C-9 C(1)RM, wiv<sup>y</sup>Lt<sup>691</sup>;redC-1Sp bnC-1;Ex

Element D

✓D-1 c-V se jv ic blr  
D-2 c-V se jv ht redD-1G1 ruw

Element A

A-1 C(1)RM, wiv<sup>y</sup>Lt/Y x w<sup>sw</sup>/Y  
A-2 wak  
A-3 wrz  
A-4 whg  
A-5 redA-3Gr  
A-6 bnA-1Ko bnA-2Gr  
A-7 v<sup>55</sup>bnA-2Gr  
A-8 v Ax sc y m redA-2Sp bb  
A-9 C(1)RM wiv<sup>y</sup>Lt.yLtN<sup>691</sup>wiv/Y x v sc sn/Y  
A-10 v sc y m f<sup>64</sup>redA-2Sp bb  
A-11 sdx  
A-12 y<sup>tlmtl</sup>redA-2H  
A-13 y m bnA-1Sp

Element E

E-1 Bls/+  
E-2 Dl<sup>64</sup>/+; ruw  
E-3 H/+  
E-4 H<sup>67</sup>/+  
E-5 edbnE-1G1  
E-6 aa5  
E-7 aa3 redE-1Hy;nt;h;sca  
E-8 Sp<sup>2</sup>Dl<sup>59</sup>Kf/Tp<sub>2</sub>Anp

Element F

✓F-1 Ex/Ci<sup>67</sup>  
F-2 Ex/Ex  
✓F-3 Ci/Ci;T(Y;2;3;5)Do

Element B

B-1 blu ng nt  
B-2 Smr/+  
B-3 pci  
B-4 kb/blu ng nt  
B-5 lpl/blu ng nt

Chromosome aberrations

T-1 In(1)f<sup>3</sup>  
T-2 Tp(1)w<sup>m1</sup>  
T-3 In(1)w<sup>m2</sup>/wiv x wiv  
T-4 In(1)w<sup>m2</sup>Hess  
T-5 In(1LR)w<sup>m3</sup>  
T-6 In(2)Lew  
T-7 T(1;2)v<sup>t3</sup>/+ x T(1;2)v<sup>t3</sup>  
T-8 T(1;2;4)H<sup>Jag</sup>/+  
E-8 Tp(2)Anp/Sp<sup>2</sup>Dl<sup>59</sup>Kf  
F-3 T(Y;2;3;5)Do;Ci/Ci  
A-1 C(1)RM, wiv<sup>y</sup>Lt/Y x w<sup>sw</sup>/Y  
A-9 C(1)RM wiv<sup>y</sup>Lt.yLtN<sup>691</sup>wiv/Y x v sc sn/Y

Element C

C-1 Ap<sup>68</sup>/+  
C-2 Gk/+  
C-3 Stw/Stw  
C-4 Sf pb redC-1Sp sca/pb redC-1Sp sca  
C-5 mt  
C-6 tk redC-1Sp sca

OSAKA, JAPAN: OSAKA UNIVERSITY  
Medical School, Department of Genetics

<u>D. virilis</u>	<u>Chromosome 2</u>	<u>Multichromosomal</u>	<u>Chromosome 2</u>
<u>Wild Stocks</u>	10 eb	15 ru;mt w <sup>e</sup> sb 16 v;es(1;5)	18 net
1 Hikone (Japan)	<u>Chromosome 3</u>	<u>D. simulans</u>	<u>Chromosome 3</u>
2 Kaidema (Japan)	11 cn	<u>Wild Stocks</u>	19 jv se 20 st se
3 Kochi (Japan)	<u>Chromosome 4</u>	15 strains	<u>Other species</u>
4 New York (USA)	12 cd	<u>Chromosome 1</u>	D. ananassae 1 strain (USA)
5 Pasadena (USA)	<u>Chromosome 5</u>	16 v 17 y w	D. funebris 1 strain (Japan)
<u>Chromosome 1</u>	13 st B <sup>3</sup> pe 14 st es		
7 v <sup>4</sup>			
8 w <sup>a</sup>			
9 y			

TURKU, FINLAND: UNIVERSITY OF TURKU  
Department of Genetics

<u>simulans</u>	<u>Chromosome 2</u>	<u>Chromosome 3</u>
wild	b bw dh b pm(py sd?) net net b py sd pm py <sup>2</sup> py <sup>2</sup> up stw (from su-bb) up	st D1 <sup>2</sup> pe/st pe H <sup>h</sup> pe jv st pe rd rd jv se st pe st Ubx pe/st pe
<u>Chromosome 1</u>		
f <sup>2</sup>		
v		
y w		

HONOLULU, HAWAII: UNIVERSITY OF HAWAII  
Department of Genetics

<u>D. immigrans</u>	Oahu, Hawaii Rochester, New York	RSB-7-Im w 40A'-Im RSS-18-Im O-3-Im 29cId-Im RS-3-Im SFRSB-7-Im S-1-Im S-4-Im S-6-Im S-7-Im S-11-Im
Hawaii, Hawaii	sl w e	
Oahu, Hawaii	m	
Kwangju, Korea	v pm vl v d	
<u>D. mercatorum mercatorum</u>	<u>Parthenogenetic</u>	
<u>Bisexual</u>	SO-1-Im RSB-6-Im OB-2-Im	
El Salvador		
Manizales, Colombia		

D. paranaensis Please write for current list of Hawaiian species.

SAPPORA, JAPAN: HOKKAIDO UNIVERSITY  
Faculty of Science, Zoological Institute

D. virilis (2 strains)	D. sordidula (1)	D. albomicans (1)	D. ezoana (1)
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CHANDIGARH, INDIA: PANJAB UNIVERSITY  
Department of Zoology

<i>D. melanogaster</i>	<i>D. nepalensis</i>	<i>D. malerkotliana</i>	<i>D. panjabiensis</i>
<i>D. takahashii</i>	<i>D. suzukii</i>	<i>D. jambulina</i>	<i>D. immigrans</i>

CHICAGO, ILLINOIS: UNIVERSITY OF CHICAGO  
Department of Biology

<u><i>D. americana</i></u>	<u>Chromosome 1</u>	<u>Chromosome 5</u>	
1 Independence	6 w <sup>50</sup> 112	13 B <sup>3</sup> pe	21 b;sv t tb gp <sup>2</sup> ;cd;pe
2 Anderson		14 pe	22 b;sv t tb gp <sup>2</sup> ;pe
	<u>Chromosome 2</u>	15 ru	23 cd;pe
<u><i>D. texana</i></u>		16 ru st mh	24 cn;pe
	7 b bk dt	17 ru st mh pe	25 gp <sup>2</sup> ;pe
3 New Orleans	8 va	18 st es pe <sup>Jap</sup>	26 gp <sup>2</sup> S/gp <sup>2</sup> +;ru st mh
			27 pe;gl
<u><i>D. virilis</i></u>	<u>Chromosome 3</u>	<u>Multichromosomal</u>	28 "scute"(II);pe <sup>m3</sup>
<u>Wild Stocks</u>			29 t;cd
	9 gp <sup>2</sup>	19 b;cn;B <sup>3</sup> pe	30 v <sup>48a</sup> ;pe
4 Pasadena lethal-free	10 S/+	20 b;tb gp <sup>2</sup> ;cd;pe	31 v <sup>48a</sup> w;pe
5 Texmelucan	11 sv t tb gp <sup>2</sup>		32 v <sup>48b</sup> ;pe
			33 y <sup>40a</sup> ;pe

MISIMA, JAPAN: NATIONAL INSTITUTE OF GENETICS

*D. ananassae*

<u>Wild Stocks</u>			
Barro Collorado, Panama 69	w <sup>65</sup> ty y <sup>51</sup>	Dl ext	M-d ru <sup>2</sup>
(low elevation)	ty	ext	ri M-d
Turrialba, Costa Rica 101	kk <sup>c</sup>	ext se	ri
(high elevation)	w <sup>65</sup> kk <sup>c</sup>	cd Dl bw <sup>R</sup>	mot
Hawaii	od	<u>Chromosome 3</u>	<u>Chromosome 4</u>
D-pp (Pago Pago, dark)	<u>Chromosome 2</u>	px	bb <sup>67</sup>
L-pp (Pago Pago, light)	b <sup>65</sup>	px <sup>2</sup>	bb <sup>2</sup>
IM-4 (Madras, India)	ba <sup>65</sup>	ru	
L-Upolu (light)	bw <sup>R</sup>	ru <sup>2</sup>	<u>Multichromosomal</u>
F2 (Peng-Hu Is.)	ma (Hinton)	px <sup>2</sup> ru	f;cd (Hinton)
F3 ( " )	se <sup>T</sup>	sm <sup>66</sup>	f;cd se <sup>T</sup>
F5 ( " )	Arc bw <sup>R</sup>	sm <sup>66</sup> px <sup>2</sup>	f y <sup>66</sup> ;cd ba <sup>65</sup>
F8 ( " )	Arc se <sup>T</sup>	bri Rf	w <sup>65</sup> f <sup>49</sup> ;cd
Ph-5 (Malaybalay, Philippines)	b ma	bri Rf px	kk <sup>c</sup> ;cd bw <sup>R</sup>
Ph-15 ( " )	b se <sup>T</sup>	bri ru	w <sup>65</sup> ;px
D-Tonga	bw <sup>R</sup> ba <sup>65</sup>	M <sup>65</sup> px	w <sup>65</sup> y <sup>51</sup> ;M-d
V-Truk	cd ba <sup>65</sup> - 12	M <sup>65</sup> ru	w <sup>65</sup> ;bb <sup>67</sup>
	cd bw <sup>R</sup>	bri M <sup>65</sup> ru	f;cd;px
<u>Chromosome 1</u>	cd se <sup>T</sup> ba <sup>65</sup>	M-b	b se <sup>T</sup> ;px <sup>2</sup>
y (Hinton)	j b	M-b pc	bw <sup>R</sup> ;bri
w <sup>65</sup>	ma ba <sup>65</sup>	pc	bw <sup>R</sup> ;ru
w <sup>65</sup> y <sup>51</sup>	se <sup>T</sup> ba <sup>65</sup>	M-c	j b se <sup>T</sup> ;ru
w <sup>65</sup> f <sup>49</sup> y <sup>51</sup>	b bn <sup>67</sup>	M-c px	ma;bri
w <sup>65</sup> sn <sup>65</sup> y <sup>51</sup>	cd bw <sup>R</sup> bn <sup>67</sup>	Sn <sup>p</sup> M-c	se <sup>T</sup> ;ru
kk	pe	Sn <sup>p</sup> bri ru	Dl;px
w <sup>65</sup> kk y <sup>51</sup>	Dl	Sn <sup>p</sup>	Dl bw <sup>R</sup> ;ru
	Dl pe	M-d	cd bw <sup>R</sup> ;M-d

LEXINGTON, KENTUCKY: UNIVERSITY OF KENTUCKY  
Department of Zoology

D. affinis: Lexington, Kentucky	D. robusta: Lexington, Kentucky
D. busckii: Lexington, Kentucky	D. tripunctata: Lexington Kentucky
D. hydei: Lexington, Kentucky	D. immigrans: Lexington, Kentucky
D. putrida: Lexington, Kentucky	

Some of these stocks are not continuously available since they are difficult to maintain under laboratory conditions; however, most can be field collected from March through October.

MILANO, ITALY: UNIVERSITA' DI MILANO  
Istituto di Genetica

D. simulans

<u>Wild Stocks</u>	<u>Chromosome 2</u>	<u>Chromosome 3</u>	<u>Stocks selected for tumor manifestation</u>
1 Aspra	4 net	5 st	6 tu Bl
2 Serpentara			7 tu Aspra
3 Giannutri			

MEXICO CITY, MEXICO: NATIONAL COMMISSION OF NUCLEAR ENERGY  
Genetics and Radiobiology Program

3 D. virilis	9 D. pseudoobscura (Mexico City)	
4 D. hydei	10 D. immigrans	" "
5 D. ananassae	13 D. virilis	" "
7 D. neohydei	12 D. busckii	" "
8 D. simulans	14 D. funebris	" "

NORWICH, ENGLAND: JOHN INNES INSTITUTE

1 D. simulans

UTICA, NEW YORK: MASONIC MEDICAL RESEARCH LABORATORY  
Aging Program

D. funebris	D. pseudoobscura	D. saltans	D. virilis
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STOCKHOLM, SWEDEN: UNIVERSITY OF STOCKHOLM  
Institute of Genetics

All of our D. pseudoobscura lines, listed in DIS 43: 75 (1968) have been discarded.

PADOVA, ITALY: UNIVERSITA' DEGLI STUDI DI PADOVA  
Istituto di Biologia Animale

1) D. hydei	2) D. pseudoobscura	3) D. simulans
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CLEVELAND, OHIO: CLEVELAND STATE UNIVERSITY  
Department of Biology

D. virilis

<u>Wild Stocks</u>	Brazil	<u>Multichromosomal</u>
	Chile	
Argentina	Texmelucan	b;tb gp;cd;pe

SEOUL, KOREA: CHUNGANG UNIVERSITY  
Department of Biology

<u>D. auraria</u>	D. immigrans (2 wild strains)
	D. lutea (2 wild strains)
Race A (10 wild strains)	D. pseudobscura
Race B (4)	D. suzukii (2 wild strains)
Race C (10)	D. virilis (6 wild strains)

LIMBE, MALAWI: UNIVERSITY OF MALAWI  
Department of Biology, Genetics Section

Twenty species of Drosophila are kept in stock.

TÜBINGEN, GERMANY: UNIVERSITY OF TÜBINGEN  
Department of Genetics

<u>D. subobscura</u>	Norwegen	<u>D. pseudoobscura</u>
	Griechenland	
Küsnacht	Ba/1 <sup>8</sup>	wild
Belgrad	Va/+	
Lipari 1	Va/Ba <sup>210</sup>	<u>D. ambigua</u>
Ponza 2/3	ch cu	wild

SANTIAGO, CHILE: UNIVERSIDAD DE CHILE  
Facultad de Medicina, Departamento de Genética

- D. brncici: Colombia  
D. buskii: Chile (La Serena)  
D. camaronensis: Chile (Azapa)  
D. funebris: Chile (La Serena, Valdivia, Tierra del Fuego y Punta Arena)  
D. gasici: Chile (Arica), Colombia, Bolivia (Cochabamba)  
D. gaucha: Brasil (M. Capoes, C. de Jordan and Taimbas), Argentina (Cordoba, San Luis), Bolivia  
D. hydei: Chile (Camarones, Azapa, Copiapó, Antofagasta), Bolivia (Cochabamba).  
D. immigrans: Chile (Valdivia, La Granja, Las Viscachas)  
D. mercatorum: Chile (Arica, Azapa, Antofagasta), Bolivia (Cochabamba)  
D. mesophragmatica: Bolivia (La Paz), Perú (Cuzco, Machu-Picchu)  
D. pavani: Chile (Copiapó, Vallenar, La Serena, Curicó, Viña del Mar, Olmúe, Bellavista, Colbún, Los Queñes, Chillán, La Granja, Las Viscachas, Algarrobo)  
D. repleta: Chile (Arica, La Cisterna, Azapa)  
D. simulans: Chile (Azapa)  
D. viracochi: Perú (Machu-Picchu)  
D. virilis: Chile (Santiago)

PÓRTO ALEGRE, BRAZIL: UNIVERSIDADE FEDERAL DO RIO GRANDE DO SUL  
Instituto de Biociências, Departamento de Genética

D. willistoni

Wild strains from: Florida, Perú, Cuba, Guatamala, Equador, Brasil: Tracua, Serra do navio (Amapá), Manaus and Tabatinga (Amazonas), Pôrto Velho (Guaporé) Bélem (Pará), Maranguape (Ceará), Salvador (Bahia) Xingú (Mato Grosso) Brasília, Chapadinha (Distrito Federal), Tijuca (Guanabara), Itatiaia and Angra dos Reis (Rio de Janeiro), Ilha das Cobras (Paraná), Iperoba, Tubarão and Florianópolis (Santa Catarina), São Pedro and Eldorado (Rio Grande do Sul)

Chromosome 1w<sup>e</sup> y sn ru(Inv)/lethalChromosome 2207/Broad  
Em phChromosome 3

pink(Inv)/lethal

D. paulistorum

Wild strains from: Brazil: Xingú (Mato Grosso), Maranguape (Ceará), Florianópolis (S.C.). F68 - H82

D. pavloskiana (12) ApoteriD. equinoxialis Cuernavaca - TefeD. tropicalis Palmas (Goiás) 1975.1 Belém - BrasilD. cubana Trinidad - S. Domingos  
fr. J.I. TownsendD. nebulosa

Radioresistant and radioresistant strains  
Wild strains from Lima (Perú), Tingo Maria (Perú), Posto Duque (Amazonas)

D. polymorpha several wild strains and select morphsD. pseudoobscura PP10, AR/STD. stoni T-10D. victoria GH-3D. lebanonensis Casteeli, LebanonD. Pattersoni Texas

BOGOTA, COLOMBIA: UNIVERSIDAD DE LOS ANDES  
Instituto de Genética

D. paulistorum

Yaguaracaca A2

Yaguaracaca A1

Yaguaracaca B1

Yaguaracaca B2

Mitú 1A

Mitú 1B

Mitú 2B

Valparaiso 1

Valparaiso 2

Caripe 8 (Ven)

British Guiana

Gigante (Huila)

Raposo 95 - a,b,c,d

Fusa

Angra 24 (Brasil)

Chocó

Belem 11 (Brasil)

D. Willistoni - Banano

Marco 2

Yaguaracaca A (Bajo)

Yaguaracaca B (Alto)

San Pablo (Brasil)

Bosque

Recuerdo

Condoto

Valparaiso (Caquetá)

Mitú 2

Macarena

Sasaima

Manizalez

Mitú 1

Piojó

Equinoxialis

Mitú 2A

San Pablo (Bosque)

Mitú 2A

Mitú 1

Marco 1 (Brasil)

Mesas

Valparaiso

Turbo B2

Yaguaracaca A (Bajo)

Yaguaracaca B1

Macarena

Umaripunta bosque  
(Vaupes)Tropicalis

Mitú 2

Valparaiso (Caquetá)

San Pablo (Bosque)

Macarena

Umaripunta Est. 2  
(Vaupes)

LEEDS, ENGLAND: UNIVERSITY OF LEEDS  
Department of Zoology

D. busckiiD. funebrisD. hydeiD. immigransD. littoralisD. obscuraD. phalerata

(several strains)

D. subobscuraD. transversa

KALYANI, INDIA: KALYANI UNIVERSITY  
Department of Zoology, Genetics Laboratory

D. ananassae

1. wild

2. se px po fa

BARCELONA, SPAIN: UNIVERSITY OF BARCELONA  
Department of Genetics, Faculty of Sciences

D. affinis - Nebraska (USA)	D. immigrans - Spanish stocks	D. simulans - Spanish stocks
D. ambigua - Spanish stocks	D. littoralis - Spanish stocks	D. subobscura - Spanish stocks
D. bifasciata - Pavia (Italy)	D. mercatorum - Spanish stocks	mutant stocks
D. busckii - Spanish stocks	D. obscura - Spanish stocks	D. testacea - Spanish stocks
D. buzzati - Spanish stocks	D. phalerata - Spanish stocks	D. tolteca - Medellin (Colombia)
D. cameraria - Spanish stocks	D. persimilis	D. transversa - Spanish stocks
D. emarginata	D. pseudoobscura - Texas (USA)	D. tristis - Snery (Switzerland)
D. funebris - Spanish stocks	D. repleta - Spanish stocks	D. victoria - Prescott (USA)
D. hydei - Spanish stocks		Megaselia scalaris - Spanish

VARANASI, INDIA: BANARAS HINDU UNIVERSITY  
Department of Zoology

Wild Stocks

(a) D. ananassae (7 strains)	(c) D. malerkotliana	(f) D. raychaudhurii
(b) D. bipectinata (Calcutta)	(d) D. nasuta	(g) D. latifshahi
	(e) D. kikkawai	(h) D. seguyi

Mutants of D. ananassae

<u>Chromosome 1</u>		<u>Chromosome 3</u>	<u>Unlocated mutants</u>
y	cu se		dct
wa	b se		sp
vs	cu b	px pc	ci
	b	stw pc	arch
	cu	stw px	
	se	stw	
<u>Chromosome 2</u>	ic	px	
	cu bw	pc	
cu b se	ssa		

HELSINKI, FINLAND: UNIVERSITY OF HELSINKI  
Department of Genetics

D. simulans (2 strains)	D. testacea (6 strains)	D. funebris (7 strains)
D. tristis Zürich	D. kunzei Interlaken,	D. littoralis (11 strains)
D. obscura (23 strains)	Switzerland	D. subarctica (8 strains)
D. bifasciata (9 strains)	D. transversa (15 strains)	D. willistoni (1 strain from
D. subobscura (8 strains)	D. limbata (2 strains)	Galápagos)
D. ambigua (3 strains)	D. busckii (7 strains)	Chymomyza costata (14
D. hydei (5 strains)	D. immigrans (1 strain from	strains)
D. phalerata (11 strains)	Finland & 1 from Zürich)	Scaptomyza pallida (1 strain)

Note: If not mentioned especially, stocks are collected from natural populations in Finland, Sweden and Norway.

BHAGALPUR, INDIA: BHAGALPUR UNIVERSITY  
Department of Zoology, Drosophila Laboratory

D. ananassaeDarjeeling populationBhagalpur population

ST<sup>2</sup>/ST<sup>2</sup>, ST<sup>3</sup>/ST<sup>3</sup>  
 AL/AL, ST<sup>3</sup>/ST<sup>3</sup>  
 ST<sup>2</sup>/ST<sup>2</sup>, DE/DE  
 AL/AL, DE/DE  
 New gene arrangement in IIL

ST<sup>2</sup>/ST<sup>2</sup>, ST<sup>3</sup>/ST<sup>3</sup>  
 AL/AL, ST<sup>3</sup>/ST<sup>3</sup>  
 ST<sup>2</sup>/ST<sup>2</sup>, DE/DE  
 AL/AL, DE/DE  
 New gene arrangement in IIL  
 Pericentric inversion between IIL & IIL

## DROSOPHILA SPECIES - NEW MUTANTS

Report of F.M.A. van Breugel

w<sup>m</sup>CoY: white-mottled Confluens Y Cytology: Dp(1;Y)16B<sub>2</sub>-17B<sub>1</sub> NOR?;Y<sup>L</sup>. Insertional duplication (probably including part of the X chromosomal nucleolus organizer region) by transfer of a small fragment of X into the Y chromosome. Derived from w<sup>ml</sup> and found as a single exceptional male from a cross: Dp(1)Co<sup>Nt</sup>, w<sup>hg</sup>/w<sup>iv</sup> ♀ x w<sup>ml</sup>, y, m♂. The male was y<sup>+</sup>m<sup>+</sup> and showed large-spotted type of mottling. Males carrying the duplication Y chromosome in addition to being mottled in the presence of a white-marked X chromosome, show a Confluens phenotype, a wing vein character caused by duplication of the Notch<sup>+</sup> locus. The w<sup>m</sup>Co duplication not infrequently may become transferred by some exchange mechanism, in the male, to the X chromosome. w<sup>m</sup>Co serves as a suitable marker of the Y chromosome. RK1.

Xw<sup>m</sup>Co: white-mottled Confluens X Cytology: Dp(1;1)16B<sub>2</sub>-17B<sub>1</sub> NOR?;X<sup>R</sup>. Product of X-Y exchange in X<sup>w</sup>/w<sup>m</sup>CoY males. Insertional duplication of euchromatin of X, including w<sup>+</sup> and N<sup>+</sup>, into the heterochromatic (=Right) arm of X. Insertion quite terminally in heterochromatin most likely neighboring the X chromosomal nucleolus organizer region. RK1-2.

C(1)RM w<sup>hg</sup>.w<sup>hg</sup> (white-honing) Spontaneous as a single homozygous w<sup>hg</sup> female in a w<sup>iv</sup> marked attached-X stock of Gregg (1957). Rather dark white allele, amber to prunelike in color. w<sup>hg</sup> is also available in detached condition (X-rayed). RK1.

C(1)RM In(1)N<sup>68</sup>, w<sup>hg</sup>.w<sup>hg</sup> Inversion Notch<sup>68</sup> in one arm of the C(1)RM chromosome. Cytology: C(1)RM In(1) 17A<sub>1,2</sub>-18A<sub>1,2</sub>. X-ray induced. Females have w<sup>hg</sup> eye color and notched wings and often malformed eyes. The w locus is included in the inversion. In(1)N<sup>68</sup>, w<sup>hg</sup> is also available in detached condition (X-rayed). RK1.

C(1)RM Dp(1)Co<sup>Nt</sup>, w<sup>hg</sup>.w<sup>hg</sup> Insertional duplication Confluens-Net in one arm of C(1)RM w<sup>hg</sup>.w<sup>hg</sup>, including the N<sup>+</sup> and w<sup>hg</sup> loci. Cytology: C(1)RM Dp(1)16A-18D; 20D. X-ray-induced. Females have a darker eye color than ordinary attached w<sup>hg</sup>, indicating triplication of the w<sup>hg</sup> locus. The abnormal wing venation is caused by a triplication of the N<sup>+</sup> locus. Dp(1)Co<sup>Nt</sup>, w<sup>hg</sup> is also available in detached condition (X-rayed). RK2(attached). RK3(detached).

w<sup>r2</sup> (white-roze) Arose spontaneously; found as two pink-eyed males in a w<sup>m2</sup>/w<sup>iv</sup> stock containing supernumerary Y chromosomes. RK1.

(Supplementary information will be given in Genetica 41, 1970 and 42, 1971, in press).



hydeiReport of H. Gloor

(Abbreviations: loc = localization; rec = received; ref = reference; syn = synonymous. Some of these have been reported previously. See also Geneva's hydei stock list, this issue.)

## white alleles:

w<sup>ak</sup>: white-abrikoos Hess 65e. syn: w<sup>a</sup>. loc: A. X-ray induced as w<sup>ak</sup>/w<sup>iv</sup>. Eye color light orange, the same in both sexes. rec: 68b from Hess, Tübingen, Germany as stock w<sup>a</sup>. ref: van Breugel 1970.

w<sup>hg</sup>: white-honing van Breugel 68k. loc: A. Spontaneous in stock C(1)RM w<sup>iv</sup> y<sup>Lt</sup> as ♀ C(1)RM w<sup>hg</sup> y<sup>Lt</sup>. Eye color dark brown-red. ref: van Breugel 1970.

w<sup>iv</sup>: white-ivoor Clausen 23. syn: w. loc: A. Spontaneous. Eye color off-white, ivory. rec: 62b from Gregg, Oxford, Ohio as stock 56. ref: Clausen 1923, Spencer 1949, van Breugel 1970.

w<sup>rz</sup>: white-roze van Breugel 68b. loc: A. Spontaneous as 2♂♂ in cross In(1LR)w<sup>m2</sup>/w<sup>iv</sup>/Y x w<sup>iv</sup>/Y(Y). Eye color pink. ref: van Breugel 1970.

w<sup>sw</sup>: white-sneeuw Holleman 67f. loc: A. Spontaneous as 1 ♂ in wild stock São Paulo. Eye color pure white. ref: van Breugel 1970.

## red eye mutants

v: vermillion Clausen (?). loc: A. Eye color a brilliant scarlet. Homology with v of D. melanogaster tested through interspecific eye disk transplantation by G.W. Beadle. rec: 61k from Gregg, Oxford, Ohio as stock 59. ref: Clausen 1923, Spencer 1949, Gregg & Smucker 1965, van Breugel et al. 1968.

v<sup>55</sup>: vermillion-55 Gloor 55c. loc: A. X-ray induced allele of v.

red A-2<sub>Sp</sub>: red eye-2 of Spencer on element A Spencer. syn: ch<sup>I</sup>: cherry-I. loc: A. Spontaneous. Eye color orange brown, darkens with age. rec: 61g from Gregg, Oxford, Ohio as stock 59. ref: Spencer 1949, Gregg & Smucker 1965.

red A-2<sub>H</sub>: red eye-2 of Hess on element A Hess 63b. syn: ch<sup>t</sup>: cherry-tomato. loc: A. X-ray induced as 1♂. Allelic and similar in phenotype with red A-2<sub>Sp</sub>. rec: 67m from Green, Davis, California as stock y<sup>tl</sup> m<sup>tl</sup> ch<sup>to-1</sup>. ref: Hess & Green 1965.

red A-3<sub>Gr</sub>: red eye-3 of Green on element A Green 63k. syn: to<sup>3</sup>: tomato-3. loc: A. X-ray induced as 1 ♂. Eye color red orange, darkens with age to dull brown. rec: 64c from Hess, Tübingen, Germany as stock to<sup>3</sup>. ref: Hess & Green 1965.

red C-1<sub>Sp</sub>: red eye-1 of Spencer on element C Spencer 37. syn: cn<sup>A62</sup> 37: cinnabar (?). Eye color a brilliant scarlet. rec: 62k from Gregg, Oxford, Ohio as stock 97a. ref: Spencer 1949, Gregg & Smucker 1965.

red D-1<sub>G1</sub>: red eye-1 of Gloor on element D Gloor. loc: D. Spontaneous. Eye color scarlet.

red E-1<sub>Hy</sub>: red eye-1 of Hyde on element E Hyde 15. syn: st: scarlet. loc: E. Eye color bright scarlet, darkens with age. rec: 61i from Wheeler, Austin, Texas as stock st jv pb sca. ref: Hyde 1915, Spencer 1949, Gregg & Smucker 1965.

red E-G1: red eye-1 of Gloor on element E Gloor 68. loc: E. Allele of red E-1Hy, but much darker. From wild population on Madeira.

#### brown eye mutants

bn A-1Sp: brown eye-1 of Spencer on element A Spencer 37. syn: gn<sup>w93</sup> 37: garnet. loc: A. Heterozygous wild ♀ collected in Wooster, Ohio. Eye color translucent purple brown, darkens with age. rec: 62g from Gregg, Oxford, Ohio as stock 62d. ref: Spencer 1949, Gregg & Smucker 1965.

bn A-1Ko: brown eye-1 of Kobel on element A Kobel 67c. loc: A. Spontaneous as several ♂♂. Allele of bn A-1Sp.

bn A-2Gr: brown eye-2 of Green on element A Green 63j. syn: pn: prune. loc: A. X-ray induced as 1 ♂. Eye color purplish dark brown. rec: 64a from Green, Davis, California as stock XX w Lt x pn. ref: Hess & Green 1965.

bn C-1: brown eye-1 on element C syn: br<sup>A</sup> 135 38: brown (?) Spencer 38. loc: C. Eye color slightly darker than wild type, in combination with red C-1Sp bright orange. rec: 65g from Gregg, Oxford, Ohio as stock XX w Lt;or;Ex.

bn E-1G1: brown eye-1 of Gloor on element E Gloor 55. syn: peach. loc: E. X-ray induced. Eye color waxy orange brown. Allelic with eye color 52 of Gregg & Smucker 1965.

se: sepia Spencer 38. loc: D. From wild population at Gatlinburg, Tennessee. Eye color dark brown, changing to black with age; accumulation of sepiapterin. rec: from Gregg, Oxford, Ohio as stock 121a. ref: Spencer 1949, Gregg & Smucker 1965.

#### morphological mutants

aa: abnormal abdomen Gloor. loc: E. A frequent phenodeviant in wild populations and especially after X-irradiation, the abnormal abdomen character often has a hereditary component. Sternites and tergites incomplete, hairs correspondingly eliminated. Penetrance and expressivity highly variable, depending on environment and genetic background. Larval and pupal segmentation normal. aa3 and aa5 are stocks which differ in several properties (dominance, expressivity, penetrance), both containing an allelic major factor on element E. Ref: Kobel & van den Bosch 1970.

Apl<sup>68</sup>: Aeroplan-68 van Breugel 68. loc: C. Spontaneous as 1 ♀. Wings spread. Homozygous lethal.

Ax: Abruptex Kobel 66h. loc: A. 1 ♀ in F<sub>2</sub> of X-irradiated first instar larvae. Veins of Ax/+ shortened (at 18°C nearly wild type), acrostichals fewer with corresponding pigment pattern changed. Enhanced by H. Lethal in ♂; Durchbrenner are small, weak, sterile, and show extreme vein erosion.

bb: bobbed Spencer. loc: A. Frequently found in wild strains. Females show reduction in bristle size, abnormal abdominal sclerites, lowered fertility, late emergence, late sexual maturation after emergence. rec: 61g from Gregg, Oxford, Ohio as stock 59. ref: Spencer 1949.

bed: breed Gloor 60. loc: C. Spontaneous. Wings broad, long, curved downwards.

Bls: Blaas Gloor 63m. loc: E. X-ray induced. Homozygous lethal. Wings spread at right angle to body, often blistered; alula without marginal hairs; viability and fertility low.

blr: blister Spencer 38. loc: D. syn: bl. From wild population at Azusa. Blister in 2nd posterior cell, penetrance lower at 18°C. rec: from Gregg, Oxford, Ohio as stock 121a. ref: Spencer 1949.

ble: blue Gloor. loc: B. Spontaneous. Color blueish gray, especially of thorax.

br: broad Kobel 66g. loc: A. Spontaneous. Wings shortened, broad. Distance between crossveins equals length of posterior crossvein. Legs (femur, tibia) shortened.

c-V: crossveinless-V Clausen (?). loc: D. Posterior crossvein missing. rec: 611 from Gregg, Oxford, Ohio as stock 121a. ref: Spencer 1949.

✓ Ci: Cubitus-interruptus Spencer 37. syn: Gp<sup>W</sup> 90 37: Gap. loc: F. Found as 1 ♂ in Wooster, Ohio. Heterozygotes with gap in L-5 proximal to posterior crossvein, homozygotes without L-5 and with gap in L-4. rec: 611 from Gregg, Oxford, Ohio as stock 122b. ref: Spencer 1949.

✓ Ci<sup>67</sup>: Cubitus-interruptus-67 Kobel 67m. loc: F. Spontaneous as 1 ♂ Do; Ci<sup>67</sup>. L-5 with gap proximal to posterior crossvein; enlarged lateral pigment spots on thorax. Homozygous lethal.

Dl<sup>59</sup>: Delta-59 Gloor 59. syn: Dl<sup>1</sup>. loc: E. Veins broadened into deltas at junction with margin. Weak allele. Suppressed by H. Homozygous lethal. ref: Gloor & Kobel 1966.

Dl<sup>63</sup>: Delta-63 Gloor 63m. loc: E. Spontaneous. Veins thickened, especially at junction with margin and with crossveins, L-1,2 extremely broad. Not entirely suppressed by H. Allelic to Dl<sup>59</sup>. Homozygous lethal.

Dl<sup>64</sup>: Delta-64 Gloor 64d. loc: E. Spontaneous. Phenotype intermediate between Dl<sup>59</sup> and Dl<sup>63</sup>. Enhancer of ruw. Homozygous lethal.

e<sup>d</sup>: ebony-dunkel Gloor 55. loc: E. X-ray induced. Larval spiracle sheaths, puparium, body, veins and wing blade darker than wildtype; tarsal bristles black instead of golden brown. Heterozygote indistinguishable from wild type.

Ex: Extension Spencer 38. loc: F. From wild population at Azusa, California. Heterozygotes with enlarged lateral pigment spots on thorax. In homozygotes spotted thorax pattern obliterated by evenly distributed brown pigment; thorax humpy. rec: 65g from Gregg, Oxford, Ohio as stock XX w Lt; or; Ex. ref: Spencer 1949.

f<sup>64</sup>: forked-64 Gloor 64c. loc: A. X-ray induced as single ♂. Bristles gnarled with ends forked; thorax color darker than wild. ref: Hess & Green 1965.

Gk: Gekneused Gloor. loc: C. X-ray induced. Anterior crossvein missing, L-1 in marginal cell squashed, marginal bristle pattern disarranged. Homozygous lethal.

H: Hairless Gloor. loc: E. Anterior dorsocentrals missing, sockets present. Frequently some of the acrostichal and abdominal hairs lost. Veins not shortened. Interaction with Ax, sca, Dl and other mutants. Homozygous lethal.

H<sup>67</sup>: Hairless-67 Kobel 67. loc: E. Spontaneous as 1 ♀. Similar to H.

ht: hart Davids 61d. loc: D. Spontaneous as 1 ♂. Thorax forshortened, creased anteriorly. Campaniform sensillum of anterior crossvein transformed into bristle.

✓ ic: incomplete Spencer 37. loc: D. From wild population at Azusa, California. L-2 and L-5 incomplete distally or gap in L-2; wings slightly warped. rec: 611 from Gregg, Oxford, Ohio as stock 121a. ref: Spencer 1949.

jv: javelin Spencer 37. loc: D. From wild population at Azusa, California. Bristles long, erect, not tapered, often hooked or broken. Thorax dark. rec: 611 from Wheeler, Austin Texas as stock 139 Spencer. ref: Spencer 1949.

kb: kabeuter Gloor. loc: B. Spontaneous. Wings spread, short, rounded, curved downwards. Posterior crossvein absent or interrupted; thorax short, groove between thorax and

scutellum incomplete; abdominal hairs bristly. All leg segments shortened. Development slow; sterile.

Kf: Kerbflugel Gloor. loc: B. X-ray induced. Wing margin notched, especially costal, 2nd and 3rd posterior cell. Expression variable. Homozygous lethal. ref: Gloor & Kobel 1966.

lpl: lethal-polyploid Gloor. loc: B. Spontaneous in wild stock Rabat, Morocco. High degree of polyploidy in the imaginal anlagen. Lethal at late larval stage. ref: Gloor 1951, Staiger & Gloor 1952, Gloor & Staiger 1954.

m: miniature Spencer. loc: A. Spontaneous as 1 ♂. Small, shortened dark wings, flies small. rec: 61l from Gregg, Oxford, Ohio as stock 59. ref: Spencer 1949.

mtl: miniature-Tübingen-1 Green 63i. loc: A. X-ray induced as 1 ♂. Allele of m. rec: 67m from Green, Davis, California as stock y<sup>tl</sup> m<sup>tl</sup> ch<sup>to-1</sup>. ref: Hess & Green 1965.

mt: mat van Breugel 66. loc: C. Bromouracil-induced. Eyes dim, bristles broken, wings somewhat smaller and slightly warped.

N<sup>691</sup>: Notch-691 Frei 691. loc: A. X-ray induced. Wing notched, veins thickened, eyes often reduced. Homo- and hemizygous lethal.

ng: no-groove Kobel 67. loc: B. Spontaneous. Groove between thorax and scutellum imperfect, scutellum with nick, bristle pattern on thorax disarranged.

nt: netvleugel Baumgartner 49. syn: confluent. loc: B. Extra veins off posterior crossvein and in marginal cell, wing frequently blistered. Suppressed by Smr. rec: 55 from Mainx, Vienna, Austria as stock confluent. ref: Mainx 1949.

pb: pearly-body Spencer, syn: pearly (?), brown-thorax. loc: C. Thorax color brown-gray, spotted pigment pattern obscured. rec: 62k from Gregg, Oxford, Ohio as stock 97a.

pci: posterior-crossvein-interrupted Gloor 67. loc: B. Spontaneous. Posterior crossvein missing or interrupted, legs (femur, tibia) often short, thorax large with pigment pattern slightly abnormal.

ruw: ruw-oog Spencer (?). syn: r: rough. loc: D. Eyes rough, facets irregular. Enhanced by Dl. rec: 61i from Wheeler, Austin, Texas as stock 1878.3.

sc<sup>1</sup>: scute-1 Spencer. loc: A. Spontaneous as 1 ♂ in a cherry stock. Removes anterior orbitals and posterior sternopleurals; effect on other bristle groups variable. rec: 61k from Gregg, Oxford, Ohio as stock 59. ref: Spencer 1949.

sca: scabrous Spencer. syn: roughest. loc: C. From Gatlinburg wild population. Eyes large, bulging, some facets very large, rounded and irregularly distributed. Extra acrostichal rows; most bristles subject to twinning. rec: 62k from Gregg, Oxford, Ohio as stock 97a.

sdx: spreadex Gloor 67d. syn: spread. loc: A. Spontaneous as 1 ♂. Wings spread at an angle of 90°; flies small. Allelic with spread Green 63 (DIS 40).

Sf: Spitzflügel Gloor 55h. loc: C. X-ray induced. Wings long, pointed. Homozygous lethal.

Smr: Small-rough-eye Green 63. loc: B. X-ray induced. Eyes slightly smaller and texture rough by irregularity of facets and corresponding hairs. Suppressor of nt. Homozygous lethal. rec: 64a from Green, Davis, California as stock R/+. ref: Spencer 1949.

sn: singed Spencer. loc: A. Spontaneous as single ♀ sn/+. Short gnarled bristles and hairs. Reported as female sterile, but now fertile. rec: 61k from Gregg, Oxford, Ohio as stock 59. ref: Spencer 1949.

Sp<sup>2</sup>: Spaltthorax-2 Gloor 55h. loc: E. X-ray induced. Thorax and scutellum cleft by longitudinal groove; in weaker expression only nicked scutellum and disturbed hair pattern on thorax. Homozygous lethal. ref: Gloor & Kobel 1966.

Stw: Straw Kobel 66. loc: C. Spontaneous as 1 ♀. Body color and wings pale yellow; spotted pigment pattern on thorax obscured; bristles black; larval spiracle sheaths pale. Homozygous viable.

tk: thick Kobel 66. loc: C. Spontaneous as several flies. Tarsal segments swollen, wings blistered.

vg: vestigial Spencer 38. loc: C. From Gatlinburg wild population. Wings variably scalloped and nicked. rec: 55 from Spurway, London, as stock vg. ref: Spencer 1949.

y: yellow Clausen. loc: A. Body, wings and bristles yellow; wildtype thoracic pigment pattern indistinct; larval mouth parts lighter. rec: 61k from Gregg, Oxford, Ohio as stock 59. ref: Spencer 1949.

y<sup>Lt</sup>: yellow-Light Spencer 37. syn: Light. loc: A. From wild population at Azusa, California as 1 ♂. Dark thoracic spots reduced, in homozygotes reduction more pronounced; body color lighter, bristles black. Phenotype of males intermediate between homo- and heterozygotes. rec: 62d from Gregg, Oxford, Ohio as stock 56. ref: Spencer 1949, Hess & Green 1965.

y<sup>tl</sup>: yellow-Tübingen-1 Hess 621. loc: A. X-ray induced as 1 ♂. Phenotypically like y. rec: 76m from Green, Davis, California as stock y<sup>tl</sup> m<sup>tl</sup> ch<sup>to</sup>-1. ref: Hess & Green 1965.

#### chromosome aberrations

In(1)f<sup>3</sup>: Inversion-forked-3 Green 631. In(1)8C;18B van Breugel. X-ray induced as 1 ♂. Allelic with f<sup>64</sup>. Homozygous viable. rec: 64m from Hess, Tübingen, Germany as stock In(X)f<sup>2</sup>. ref: Hess & Green 1965, van Breugel et al. 1968.

In(1LR)w<sup>m3</sup>: Inversion-white-mottled-3 Green 64b. In(1LR)16D3-4-17A1-2;NOR<sup>D</sup>. New order: h<sup>L</sup>NOR? / 16D3-4 - 1Ah<sup>S</sup>.h<sup>L</sup>NOR / 17A1-20 van Breugel. X-ray induced as 1 ♀. Heterozygotes over white have small and large patches on reddish background. Homozygotes have facet-sized spots on yellow background. Males nearly wildtype, but mottling distinct. rec: 67m from Green, Davis, California as stock w<sup>m3</sup>/w Lt. ref: Hess & Green 1965, van Breugel 1970.

Tp(1)w<sup>m1</sup>: Transposition-white-mottled-1 Green 631. Tp(1h;le)1<sup>M</sup>;17B1-2. New order: 20-17B2 / h<sup>L</sup>NOR / 17B1 - 1h<sup>S</sup>.h<sup>L</sup> / Nor / h<sup>L</sup> van Breugel. X-ray induced as 1 ♀. w<sup>m1</sup>/w<sup>iv</sup> has large wild sectors on orange brown background; homozygotes and males nearly wild type. rec: 67m from Green, Davis, California as stock w<sup>m1</sup>/w Lt. ref: Hess & Green 1965, Mukherjee 1965, van Breugel 1970.

In(1)w<sup>m2</sup>: Inversion-white-mottled-2 Green 631. In(1L)h<sup>S</sup> - 1A;16C-D1 + In(1LR)9C3;NOR<sup>P</sup>. New order: h<sup>L</sup>NORh<sup>L</sup>(?) / 9C4 - 16CD1 / h<sup>S</sup>.h<sup>L</sup> / 9C3 - 1Ah<sup>S</sup> / 16D1 - 20 van Breugel. X-ray induced as 1 ♀. w<sup>m2</sup>/w<sup>iv</sup> has single dark facets on a yellowish background. Modified by extra Y chromosomes. Hemizygous lethal but viable with Y from stock w<sup>m2H</sup> (w<sup>m2</sup>-Hess, derived from w<sup>m2</sup>) in this condition darker mottling of the eyes. Homozygotes light yellowish. rec: 67m from Green, Davis, California as stock w<sup>m2</sup>/w and 66 from Hess, Tübingen, Germany as w<sup>m2</sup>/w<sup>m2</sup>. ref: Hess & Green 1965, Mukherjee 1965, Gloor et al. 1967, van Breugel 1970.

In(2)Lew: Inversion-Lewontin Lewontin 55. In(2)23D;35B van Breugel. From wild population at Raleigh, North Carolina. Present in several wild populations. Phenotype wild. rec: 60a from Lewontin, Rochester, New York as stock In(2)Lew/+. ref: Warters 1944.

Tp(2)Anp: Transposition-Antennapedia Gloor 55h. Tp(2)22D1;26D4;34A1. New order: 21-D1 / 26D4 - 34A1 / 26D4 - 22D1 / 34A1 - 48 Berendes. X-ray induced as 1 ♂. Antenna frequently transformed into mesothoracic leg. Penetrance and expressivity variable, environment-dependent. Homozygous lethal. ref: Berendes 1962, Gloor & Kobel 1966, van Breugel et al. 1968.

T(1;2)v<sup>t3</sup>: Translocation-vermilion-Tübingen-3 Green 63k. T(1;2)18C1;44C van Breugel. X-ray induced as 1 ♂. Allelic to v. Homozygous lethal. rec: 68a from Hess, Tübingen, Germany as stock T(X,2)v<sup>t3</sup>. ref: Hess & Green 1965, van Breugel et al. 1968.

T(1;2;4)H<sup>Jag</sup>: Translocation-Hairless-Jaguar Gloor 63m. T(1;2;4)1h;39D;76C. New order: 20-1. / 76C - 94; 21 - 39D / 1h; 77-76C / 39D - 48 van Breugel. X-ray induced as 1 ♀. Allelic to H. Homozygous lethal. ref: van Breugel et al. 1968.

T(Y;2;3;5)Do: Translocation-Doorn Gloor 61c. T(Y;2;3;5)Y;24A;26D5;70C2;97B1. New order: 21 - 24A / 97B1 - 122; 49 - 70C2 / 26D5 - 48; 95 - 97B1 / 24A - 26D5 / Y; Y- / 70C van Breugel & Kobel. X-ray induced. Tarsus of first leg somewhat shortened; sensilla campaniformia on L-3 transformed into bristles. Penetrance variable, enhanced by ht. Heterozygous females obtained by crossing to C(1) are sterile. ref: Berendes 1962.

References: Berendes, H.D. 1962, Chromosoma (Berl.) 14: 195-206; Breugel, F.M.A.v. 1970, Genetica 41 (Nr 4); Breugel, F.M.A.v. et al. 1968, Genetica 39: 165-192; Clausen, R.E. 1923, Amer. Nat. 57: 52-58; Gloor, H. 1951, Rev. Suisse Zool. 54: 520-521; Gloor, H. & H. R. Kobel 1966, Rev. Suisse Zool. 73: 229-252; Gloor, H. & H. Staiger 1954, J. Hered. 45: 289-293; Gregg, T.G. & L.A. Smucker 1965, Genetics 52: 1023-1034; Hess, O. & M.M. Green, 1965, DIS 40: 37-39; Hyde, R.R. 1915, Amer. Nat. 49: 183-185; Kobel, H.R. & Bosch, J.J.v.d. 1970, Genetica 41: 119-140; Mainx, F. 1949, DIS 23; Spencer, W.P. 1949, Genetics, Paleontology, and Evolution, 23-44. Princeton Univ. Press; Staiger, H. & H. Gloor 1952, Chromosoma 5: 221-245; Warters, M. 1944, Univ. Texas Publ. 4445: 129-174.

#### pseudoobscura

#### Report of S.E. Moyer and V.J. Merluzzi

w<sup>ec</sup>: white-ecru. Spontaneous during synthesis of Ba/Δ Bl/L from Bl Sc pr(ST) and upt bx Ba gl/Δ; or L/or kindly provided by Mr. Boris Spassky from Rockefeller University. Allelic with w (Spassky) on X. Lighter than w<sup>ec</sup> and w<sup>1</sup> of melanogaster kindly provided by Dr. Irwin I. Oster, Bowling Green State University. Eyes light buff, light yellow or "green gold", depending on age. Testes colorless. Eye color comparisons for sex, w<sup>ec</sup> and the F<sub>1</sub> ♀ w<sup>ec</sup> x w♂ are as follows:

$$F_1\delta = w^{ec}\delta > w^{ec}\phi > F_1\phi$$

Lancefield (1922 Genetics 7: 335) has described another multiple allele of this locus, white-eosin. Tests for pseudoallelism in more than 80,000 progeny from w<sup>ec</sup> w females has not resulted in crossing over; including 40,000 from w<sup>ec</sup> w; Ba/+;Bl/+ females designed to increase crossing-over on X (see Welshons and Nicoletti, 1963, DIS 38: 80).

#### QUOTABILITY OF NOTES

For previous listings see DIS 38, 42, 43, 44, and 45

Angus, D. 35:71; 42:96-97; 42:112; 45:109; 45:153-155  
 Bryant, P.J. 46:94  
 David, J. 46:84  
 David, J. and J. Bouletreau-Merle 46:83  
 Doane, W.W. 45:189  
 Langjahr, S. 46:126  
 Mather, W.B. 27:101; 33:146-147; 33:147; 34:91-92; 37:104;  
 38:55; 40:66; 41:125-126; 41:126-128; 42:85; 43:96; 43:100-  
 101; 43:101; 44:72; 44:89; 44:98; 45:74; 45:111; 45:156-157  
 46:80  
 Roberts, R. 46:158

Nickla, H.\* Arizona State University, Tempe, Arizona. Riboflavin content in Malpighian tubes of *D. melanogaster*.

Larvae of many eye color mutants of *D. melanogaster* have less yellow pigment (YP) in their Malpighian tubes (MPT) than wild type larvae<sup>1</sup>. No mutants other than those influencing eye color have been found to

alter the color of MPT<sup>2</sup>. Kikkawa<sup>3</sup> has suggested that the YP in MPT of *D. melanogaster* is 3-hydroxykynurenine, an intermediate in the synthesis of brown eye pigments<sup>4</sup>, while Forrest and Mitchell<sup>5</sup> have implicated sepiapterin, an intermediate in the synthesis of red eye pigments<sup>6</sup>. However, YP in MPT of the American roach (*Periplaneta americana*), *Tribolium confusum*, and *Ephestia kuhniella* is mainly riboflavin<sup>7,8,9</sup>. In this communication, results are presented which suggest that the major component of YP in MPT of *D. melanogaster* is riboflavin.

The technique used for chromatographic separations followed that of Hadorn and Mitchell<sup>10</sup>. Ascending paper chromatography was carried out in the dark with n-propanol and 5% ammonia (2:1) as the solvent. Flies were reared on standard agar-cornmeal-brewer's yeast-molasses-sucrose-propionic acid medium.

For comparison with YP, sepiapterin was obtained from heads of sepi mutant flies by paper chromatography and was identified by its fluorescent color (yellow) and Rf value<sup>10,11</sup>. Following elution in 50% aqueous acetone<sup>5</sup>, the visible portion of the absorption spectrum was determined (in 50% acetone) at pH values 1.8, 7.4, and 13.4. In alkaline solution, there is a maximum at 441nm; in neutral and acid solution, there is a maximum at 418nm. Riboflavin was also chromatographed and eluted in 50% aqueous acetone. At pH 6.1 the absorption maxima are 445nm and 375nm; 447nm and 375 nm at pH 8.9; and 450nm and 353nm at pH 12.0. Yellow pigment from MPT was separated chromatographically from abdomens of approximately 1,300 wild type (Urbana) female flies. These chromatograms, when viewed under ultraviolet light reveal a bright yellow spot with approximately the same Rf value as riboflavin and sepiapterin. Several mutants (light and clot), which have pale MPT upon visual examination, were chromatographed. In both cases there was a reduction in concentration of this yellow fluorescent spot indicating that it is responsible for the yellow color of MPT in wild type flies. Following elution, the visible portion of the absorption spectrum of YP was determined at pH values 7.6 and 13.3. The absorption spectra obtained at these pH values are similar, which suggests the absence of large amounts of sepiapterin. In addition, the portion of the spectrum from 430nm to 540 nm resembles that of riboflavin.

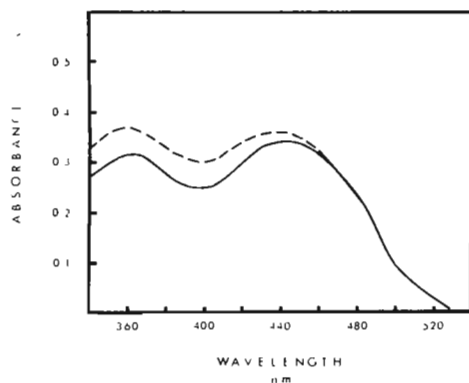


Figure 1. Absorption spectrum of riboflavin before (solid line) and after (broken line) treatment with bromine.

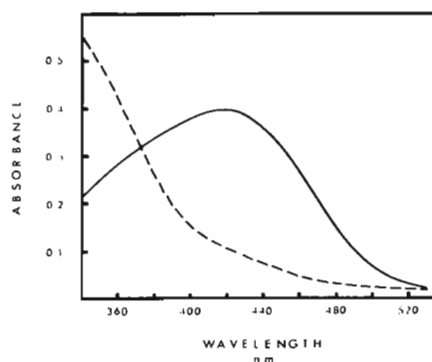


Figure 2. Absorption spectrum of sepiapterin before (solid line) and after (broken line) treatment with bromine.

Bromine causes immediate decolorization of sepiapterin producing two blue fluorescent compounds<sup>5</sup>, without altering the characteristic yellow color of riboflavin<sup>12</sup>. Figures 1 and 2 present the absorption spectra of riboflavin and sepiapterin respectively before and after treatment with bromine (0.1ml of bromine/3ml of solution). Inspection of the sepiapterin

solution revealed a complete absence of yellow color after bromine treatment. Similar treatment of YP from MPT did not alter its absorption spectrum (Figure 3) and did not cause loss of color. The absence of the typical riboflavin spectrum (Figure 3) probably results from incomplete purification of YP as only single dimension chromatography was used.

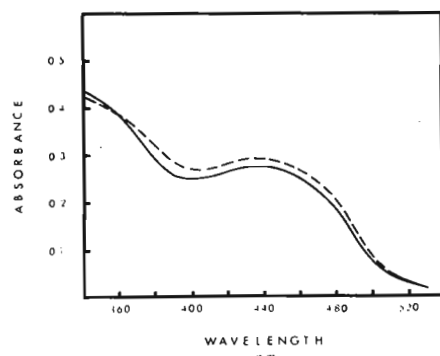


Figure 3. Absorption spectrum of the yellow pigment from the Malpighian tubes of wild type flies before (solid line) and after (broken line) treatment with bromine.

flavin in MPT is under control of a single gene whereas the amount stored is controlled by a small number of different genes. Sang<sup>15</sup> demonstrated that riboflavin is an essential dietary factor for normal development in *D. melanogaster*. The results presented in this

Table 1. Mean Rf values of riboflavin, sepiapterin, and the yellow pigment from Malpighian tubes following treatment with bromine. Kynurenine not previously treated with bromine is also given.

Compound	Visible Color	Fluorescent Color	Rf*	
			A	B
Riboflavin	Yellow	Yellow	.34	.22
Sepiapterin	Colorless	Blue	.16	.17
Yellow Pigment	Yellow	Yellow	.35	.23
Kynurenine	Yellow	Blue	.55	.36

\*Solvents: (A) n-propanol and 1% ammonia (2:1), (B) n-butanol, acetic acid, and water (4:1:1).

After bromine treatment, riboflavin, sepiapterin, and YP were chromatographed using two solvent systems and their Rf values were determined (Table 1). The fluorescent color was determined with an ultraviolet light. The absence of a blue fluorescent spot in bromine-treated YP strongly suggests that little, if any, sepiapterin is present in MPT. Since 3-hydroxykynurenine is an alpha amino acid, its presence can be determined by the ninhydrin test. Chromatograms (developed in n-propanol and 5% ammonia) containing kynurenine, sepiapterin, and YP from MPT were sprayed with a 0.2% (in acetone) solution of ninhydrin. Only kynurenine produced a positive ninhydrin reaction. Hadorn and Mitchell<sup>10</sup> also found that ninhydrin-positive materials are found only in trace amounts in chromatographed MPT.

Weber and Roberts<sup>14</sup> demonstrated that the primary site of riboflavin storage in *Tribolium confusum* is in the MPT. They concluded that the ability to store riboflavin in MPT is under control of a single gene whereas the amount stored is controlled by a small number of different genes. Sang<sup>15</sup> demonstrated that riboflavin is an essential dietary factor for normal development in *D. melanogaster*. The results presented in this communication support the hypothesis that riboflavin accumulates in MPT<sup>13</sup>, and that the ability to absorb riboflavin from the diet and store it in MPT is intricately related to eye pigment metabolism in *D. melanogaster*. The nature of this relationship is under investigation.

References: <sup>1</sup>Brehme, K.S., and Demerec, M., *Growth* 6:351 (1947); <sup>2</sup>Brehme, K.S., *Proc Natl. Acad. Sci. U.S.* 27:254 (1941); <sup>3</sup>Kikkawa, H., *Adv. in Genetics* 5:107 (1953); <sup>4</sup>Butenandt, A., Weidel, W., and Schlossberg, H., *Z. Naturforsch.*, 4b:242 (1949); <sup>5</sup>Forrest, H.S., and Mitchell, H.K., *J. Am. Chem. Soc.* 76:5658 (1954); <sup>6</sup>Kaufman, S., *Ann. Rev. Biochem.* 1:171

(1967); <sup>7</sup>Metcalf, R.L., and Patton, R.L., *J. Cell. Compar. Physiol.* 19:373 (1947); <sup>8</sup>Weber, J. and Roberts, C.W., *Canad. J. Genet. Cytol.* 8:796 (1966); <sup>9</sup>Caspari, E., and Blomstrand, I., *Genetics* 43:679 (1958); <sup>10</sup>Hadorn, E., and Mitchell, H.K., *Proc. Natl. Acad. Sci. U.S.* 37:650 (1951); <sup>11</sup>Gregg, T.G., and Smucker, L.A., *Genetics* 52:1023 (1965); <sup>12</sup>Wagner-Jauregg, T., in *The Vitamins* (edit. by Sebrell, W.H., and Harris, R.S.), 299 (Academic Press, New York, 1954); <sup>13</sup>Ziegler, I., *Adv. in Genetics*, 10:349 (1961); <sup>14</sup>Weber, J., and Roberts, C.W., *Canad. J. Genet. Cytol.* 9:302 (1967); <sup>15</sup>Sang, J.H., *Exp. Biol.* 33:45 (1955).

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Kuroda, Y. National Institute of Genetics, Misima, Japan. Fibroblastic cells derived from pupal ovary of *D. melanogaster* in culture.

slightly modified and supplemented with 0.1 mg/ml fetuin, 5 mg/ml peptone and 15% fetal bovine serum.

After several hours of cultivation, many fibroblastic cells came out from the cut end of ovarian fragments explanted, and stretched their cytoplasm on the surface of culture flasks. They increased gradually in number in further cultivation and formed a network around the original explants after 24 hours of cultivation (Fig. 1). Mitotic figures were frequently observed. Under a phase microscope some small granules were observed in the cytoplasm and one or two nucleoli were found in the nucleus. From their morphology and behavior, these fibroblastic cells seem to be derived from the lumen cells in the ovarian cavity.

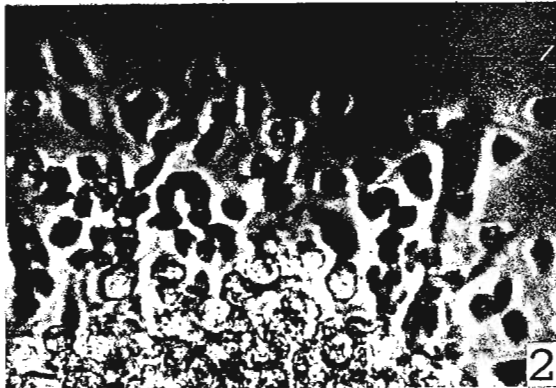
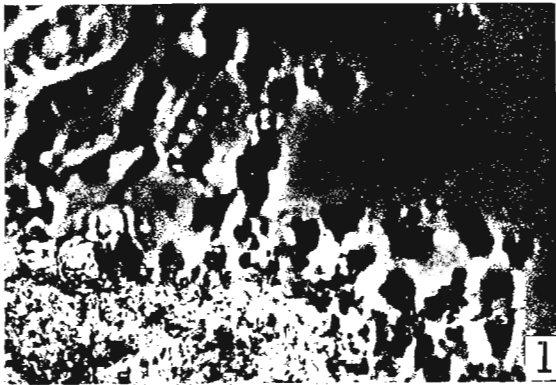


Fig. 1. Fibroblastic cells derived from a pupal ovarian fragment. After 24 hours of cultivation. Phase. x 540.

Fig. 2. Fibroblastic cells derived from an embryonic fragment. After 24 hours of cultivation. Phase. x 540

Fig. 3. Slender spindle-shaped cells derived from another embryonic fragment. After 24 hours of cultivation. Phase. x 540.

Of interest is the finding that the fibroblastic cells derived from ovarian fragments were very similar in morphology to one type of cells derived from embryonic fragments (Fig. 2). Among several types of cells derived from fragments of 12-hour embryos two types of cells were predominantly observed under the culture conditions employed: the fibroblastic cells and more slender spindle-shaped cells (Fig. 3).

Some insect cell lines established by Grace (3,4) were derived from ovarian tissues of *Antheraea eucalypti* and *Bombyx mori*. Although it is uncertain that Grace's cell lines originated from the same type of cells in the ovary as the fibroblastic cells observed in the present study, these fibro-

blastic cells may have some advantage in growing under in vitro culture conditions.

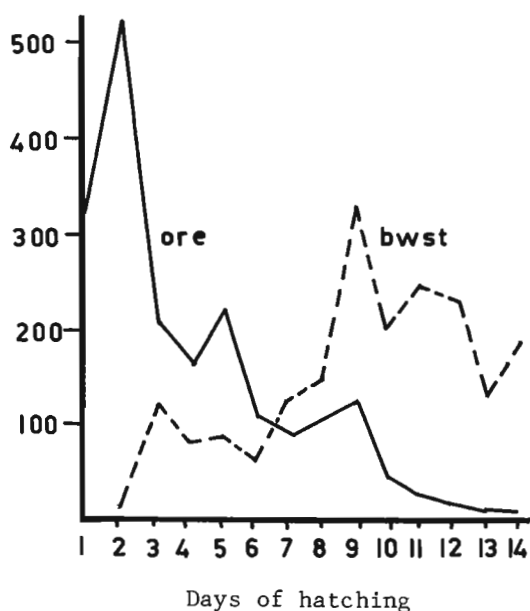
References: 1. Kuroda, Y., 1969 Japan. J. Genetics 44 Suppl. 1: 42; 2. Kuroda, Y., 1970 Exp. Cell Res. 59: 429; 3. Grace, T.D.C., 1962 Nature 195: 788; 4. Grace, T.D.C., 1966 Nature 216: 613.

Chita, O. Institut für allgemeine Biologie, Vienna, Austria and University of Craiova, Rumania. Larval-larval or larval-female interaction in *D. melanogaster*.

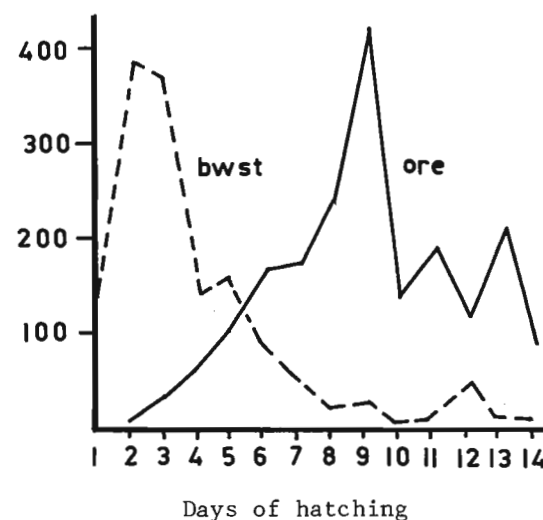
In order to find out the effect of earlier egg laying in a culture bottle with abundance of food a small experiment was carried out as described underneath. For this purpose 100 fertilized normal females (Oregon-strain) five days old (kept

during this period with males) were then transferred to a vial containing normal corn meal food and allowed to lay eggs for 24 hours. These females were then discarded and replaced by 50 fertilized females and 50 males of the mutant strain bw,st. The mutant flies were allowed to lay eggs for 10 days continuously and then removed before the first flies hatched. In another set of experiments the same procedure was followed with the exception that at first 100 mutant bw,st females were used and then replaced by normal Oregon females and males. The bottles were repeatedly seeded with yeast solution so that the food supply was abundant. Fourteen days after the first females were put into the vials, the offspring started to hatch. From that moment the hatching flies were counted daily for a period of two weeks. In total, the offspring from 14 vials, seven for each experiment, were counted. The results are shown in Figure 1. The diagram on the left corresponds to the cultures with Oregon ♀♀ first, the right to those with bw,st ♀♀ first. It can be clearly seen that whichever genotype was used first has its peak for hatching flies on the second day with steady reduction on the days following. The genotype which had a delay of a day in egg laying, however, has its peak on the 9th day.

Number of  
individuals  
hatched



Number of  
individuals  
hatched



From the data presented it can be concluded that in competition the genotype of the eggs laid first is at an advantage. The period between the hatching peaks for the two groups is much longer than the period of delay in egg laying. This finding can be interpreted in two ways. Either there is an interaction between larvae in the sense that the older larvae can suppress the development of the younger ones, or females which are ready to lay eggs do not do so if there are already eggs deposited on the food surface. Egg laying, however, cannot be hindered by the females longer than a certain period. The hatching behaviour of the strains used, irrespective of the strain allowed to lay eggs first, is practically identical.

Wong, P.T., W.D. Kaplan, and W.E. Trout III.  
City of Hope National Medical Center, Duarte,  
California. Alteration of response to a  
visual stimulus by a cholinesterase inhibitor.

The behavior mutant,  $Hk^1$  (hyperkinetic),  
discovered in our laboratory, jumps and  
falls over when presented with a light  
stimulus in the form of a burst of strobe  
flashes. The sensitivity of these flies  
to such a stimulus depends on the intensi-

ty and frequency of the strobe light used. We are currently investigating the effect of various cholinesterase inhibitors on this response. The present communication provides data on the effect of eserine.

Flies were starved overnight and fed for three hours on eserine dissolved in a one % sucrose solution. Control flies were fed sucrose solution only. The mean % response of  $Hk^1$  flies fed different concentrations of eserine and tested at three different light intensities is shown in figure 1. Each point on the graph represents a sample of ten flies

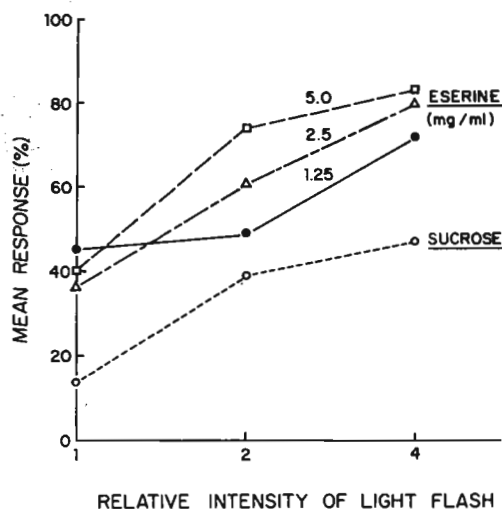


Figure 1

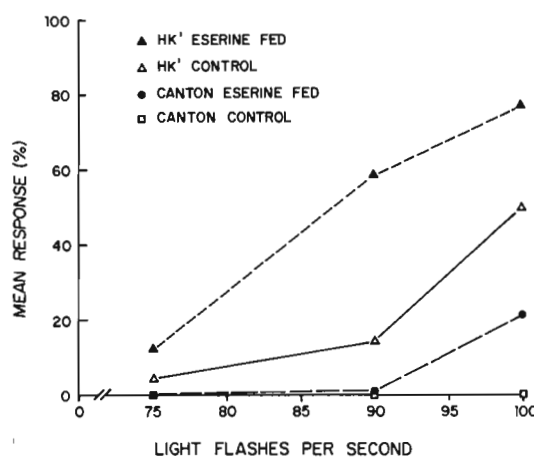


Figure 2

and each fly was tested 30 times. Each stimulus consisted of a burst of 25 light flashes at a frequency of 100 flashes per second. The intensity of the strobe light corresponding to a relative scale of four on the graph is approximately 350,000 candlepower. The eserine fed flies showed an increase in sensitivity to the light stimulus; at the two higher intensities, the mean % response appears to be directly proportional to the concentration of eserine. The increase in sensitivity of eserine-fed  $Hk^1$  flies is also evident when the mean response is measured against frequency of the strobe flashes (figure 2); the strobe was set to give 25 flashes at a relative intensity scale of four.

Even Canton-S (wild type) flies, which do not normally respond to the light stimulus, can be made sensitive by feeding eserine. The type of response appears to be identical to that of  $Hk^1$ , so in this sense eserine produces  $Hk^1$  phenocopies. However, eserine fed flies do not shake their legs while etherized, as does  $Hk^1$ . Thus the change in behavior may be primarily an increased sensitivity of the sensory system rather than the motor system. Dewhurst et al (1970) have suggested that acetylcholine may be a sensory transmitter in *Drosophila*. Therefore eserine, by blocking the degradation of acetylcholine, may increase the sensitivity of the sensory system which when stimulated sufficiently produces a startle response.

Reference: Dewhurst, S.A., McCaman, R.E., and Kaplan, W.D. *Biochemical Genetics* 4: 499-508 (1970).

Supported by NIH grant No. NS08014.

Thompson, S.R. and J.E. Putnam. Ithaca College, Ithaca, New York. Alteration of the rudimentary wing phenotype with Minutes and temperature.

temperatures on wing development in  $r^{39k}$  was studied. Arbitrary rating of the Minute, and the effect of rudimentary wing phenotypes were arbitrarily rated

Besides female sterility, the syndrome of abnormalities associated with rudimentary mutants includes modification of the size and morphology of the wing. The effects of second and third chromosomal Minutes in combination with  $r^{39k}$  and various temperatures on wing development in  $r^{39k}$  wings. Table 1 lists the Minute allele, the Minute upon the rating of  $r^{39k}$  wings. on a scale of 3, a wing having an extreme defect was given a rating of 1, a moderate (intermediate) defect a rating of 2, and a normal wing a rating of 3. The Minutes were classified on a scale of 4, with 4 indicating extremely reduced bristle size and 0 indicating normal bristle size. In general, it seems that the stronger the Minute the more nearly normal the rudimentary wings appear. However, the Minutes appeared to have no modifying effect on the sterility associated with homozygous rudimentary females.

Since Schultz (1979) demonstrated that Minutes delay the length of time required for normal development, rudimentary wing development was studied at different temperatures, which would alter the developmental rate. White prepupae were collected, placed in shell vials and allowed to undergo metamorphosis in constant temperature water baths at three different temperatures ( $18^{\circ}$ ,  $23^{\circ}$ , and  $28^{\circ}$  C.). Table 2 lists the effect of these temperatures on pupal development time and on wing size in  $r^{39k}$ .

Because the slower development, produced by the lower temperature ( $18^{\circ}$  C), yielded wing sizes which were more normal than controls ( $23^{\circ}$  C), and approximated those found with strong Minutes, it seems likely that the Minute effect was due to the slowing down of development. Because the two extreme temperatures,  $18^{\circ}$  and  $23^{\circ}$

C, have no overlap in mean wing size it was possible to use this difference to determine the

Table 1. Minute Interactions with  $r^{39k}$ .

Minute Allele	Mean Minute Rating	Mean $r^{39k}$ Rating
Control (no Minute)	----	1.75
M(2)1 <sup>2</sup>	3.75	3.00
M(2)S7	3.25	1.80
M(2)S6	3.25	2.00
M(2)B	3.00	2.00
M(2)173	3.00	2.50
M(2)S5	2.75	2.50
M(2)S3	2.50	1.75
M(2)S4	2.25	1.75
M(2)S10	1.75	1.75
M(2)S11	1.00	1.80
M(2)S9	1.00	1.75
M(2)S8	0.50	1.75
M(3)w	4.00	3.00
M(3)B <sup>2</sup>	3.50	2.00
M(3)1	3.00	1.75
M(3)S31	2.50	2.00
M(3)y	2.00	2.00
M(3)S34	1.75	2.50
M(3)36e	1.25	2.00
M(3)S36	1.00	1.75
M(3)B	0.00	1.75

Table 2. Effect of temperature on  $r^{39k}$ .

Temperature $^{\circ}$ C.	Number of Flies	Mean Pupal Development Time in Hrs.	Mean Wing Rating
18	60	192.5	2.47
23	81	127	1.75
28	44	91.5	1.40

period in which the wing development was affected by the temperature (Temperature Effective Period). Prepupae were maintained at one temperature,  $18^{\circ}$  C, for a period of time and then later transferred to the second temperature,  $28^{\circ}$  C, to complete their development. The results of these transfers demonstrated that the temperature effective period lies somewhere between 15 and 70 hours in pupal development. Further experimentation would be necessary to more closely define this critical period.

Reference: Schultz, J. 1979. Genetics 14: 366-419.

Grossfield, J. and W.L. Pak. Purdue University, Lafayette, Indiana. Localization of electroretinogram mutants.

We have reported the isolation and physiological characterization of visual mutants which show an abnormal electroretinogram (ERG). Mutants x-7 and x-14 lack the "on" and "off" transients present in the wild type ERG and thus give the

appearance of an isolated receptor response. Mutants x-12, 13, 16 and 24 show a gradation of response from none at all to a slow small amplitude response. These mutants may represent altered visual pigment. Allelism tests show that at least three distinct cistrons have been functionally identified. (See Pak, Grossfield and Arnold, Nature 227: 518 (1970) for pictures of electrophysiological phenotypes and the allelism tests.)

Here we present data for the map positions of the three loci. x-7 was mapped using the multiply marked X chromosomes sc cv v f and sc cv m f; x-14 with sc cv v f and m f car; x-12 with sc cv v f and y cv v f. The remaining mutants were just mapped a single time: x-13 and x-16 using sc ec cv ptg and x-24 with y w cv m f.

Table I. Position of Visual Mutants on X-chromosome.

Mutant	Number of Crossovers Between	Recombinational Fraction Between Mutant and Nearest Marker	Map Position *
x-12	sc and x-12	39	$6.8 \pm 1.4$ $1.5$
	x-12 and cv	40	
x-12	y and x-12	44	$6.5 \pm 1.3$ $1.4$
	x-12 and cv	49	
x-13	ec and X-13	12	$6.8 \pm .8$ $.6$
	x-13 and cv	62	
x-16	ec and x-16	6	$6.9 \pm 1.4$ $.9$
	x-16 and cv	28	
x-24	w and x-24	36	$5.5 \pm 1.0$
	x-24 and cv	111	
x-7	cv and x-7	41	$26.1 \pm 2.1$ $2.7$
	x-7 and v	23	
x-7	cv and x-7	60	$26.3 \pm 2.1$ $2.7$
	x-7 and m	47	
x-14	m and x-14	10	$53.5 \pm 1.7$ $2.1$
	x-14 and f	57	
x-14	v and x-14	58	$51.1 \pm 2.1$ $2.7$
	x-14 and f	18	

\* Confidence limits correspond to 5% significance level. W.L. Stevens J. Genetics 43: 301-307 (1942)

Although these mutants were generated by our phototaxis assay procedure we did not rely on phototaxis for characterization of the mutant phenotype for mapping. F<sub>2</sub> males of all recombinant classes from F<sub>1</sub> heterozygous females were individually tested for their electrophysiological phenotype. Recording of the ERG gives unequivocal identification of each visual mutant and visual inspection of each male provided identification of the usual morphological markers associated with each all-or-none ERG response. The use of markers on either side of each ERG mutant provided unambiguous localization. Recombination rate in the interval containing each mutant was not significantly different from standard values.

The recombination data agree with our earlier allelism tests in delimiting three loci, x-7, which we have shown to be allelic with tan, x-14 and x-(12, 13, 16, 24). It is interesting that x-7 and x-14, which have the same phenotype are more than 20 map units apart. This indicates that there may be more than one way of blocking "on" and "off" transients. Supported by grant GB-8140 to Dr. W.L. Pak and an NIH Health Sciences Advancement Award to Purdue University.

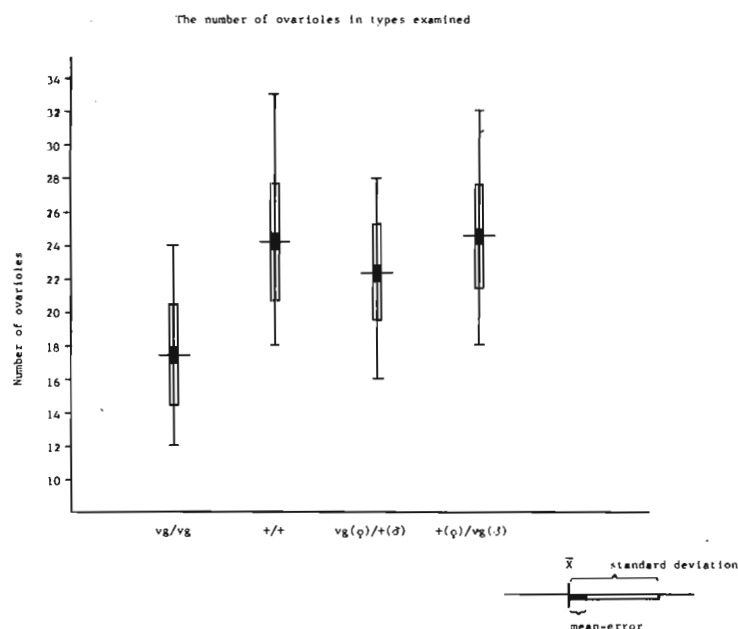
Mikulska, Grygoń. Department of Zoology, University of Nicholas Copernicus, Institute of Biology, Toruń, Poland. The number of ovarioles in the ovaries of females of reciprocal crosses of the types wild of Toruń and *vg/vg* *D. melanogaster* Meig.

The fecundity of females heterozygous for *vg* originating from the cross of wild type of Toruń and *vg/vg* was examined by Grygoń (1970). The result of crossing was always advantageous when compared with *vg/vg*, and compared with *+/+*, it was advantageous if the female used was *+/+*. With reference to these studies, the anatomical

verification is presented here.

The numbers of ovarioles were counted in the following types: 1. *+/+* of Toruń, which has been bred since 1960 in the Genetic Laboratory of Zoology Department in Toruń, 2. *vg/vg*, originating from the laboratory ETH in Zürich and bred in Toruń since 1966 and 3. females originating from reciprocal crosses of these types.

The number of ovarioles in types examined



All the flies were raised on cornmeal medium (500 g of water, 4 g agar, 34 g sugar, 68 g cornmeal and yeast suspension - for 10 bottles) in excess of the medium and uniform conditions. Ten pairs of virgin flies were used to start each culture. The ovaries were removed on the fifth day of life and they were stained with methylene blue in Ringer's fluid with a few drops of alcohol to harden the tissues (Melou 1961). The statistic analysis of results was based on a sample of 50 individuals and from each one ovary was prepared (Melou's method). The results are presented in the table and illustrated in the diagram. As they show, the type *+/+* of Toruń has an average number of ovarioles of  $24.2 \pm 0.98$ .

In comparison to the French type Banyuls-Union and Oregon (Melou op.cit.) it is a little more fecund. The type *vg/vg* indicated  $17.44 \pm 0.44$ .

The number of ovarioles in types examined

Number of ovarioles in:	<i>vg/vg</i>	<i>+/+</i>	<i>+(♂)/vg(♀)</i>	<i>+(♀)/vg(♂)</i>
Range	12 - 24	18 - 33	16 - 28	18 - 32
M (n=50)	$17.44 \pm 0.44$	$24.2 \pm 0.98$	$22.36 \pm 0.39$	$24.5 \pm 0.43$

Females of the reciprocal cross in which the *+/+* female was used, showed a slightly higher number of ovarioles than the type *+/+*, and a slightly lower number when the female used for crossing was *vg/vg*.

As in the initial crosses the females were *vg/vg* both combinations seemed to be expedient.

These anatomical relations explain the experimental data of Grygoń (op. cit.) and are in accord with them.

References: Grygoń, B., 1970, Zesz. Nauk. UMK Toruń, Nauki Mat. Przyr. 22, Biologia 12: 27-36; Melou, J.P., 1961, Ann. de Génétique 3: 25-28.

Hedrick, P.W. University of Kansas, Lawrence, Kansas. Competition experiments between *D. melanogaster* and *D. simulans*.

J.S.F. Barker tested a number of *D. melanogaster* strains against his ver strain of *D. simulans* in order to find a *D. melanogaster* strain of approximately equal competitive ability. He selected a yw strain that produced 45% *D. melanogaster* progeny when 50% of the parents were *D.*

*melanogaster*.

In order to test the competitive ability of these two strains over a longer period of time, I initiated a series of serial transfer experiments. In these experiments all freshly emerged adults from several different age bottles during a certain time period were combined in a fresh bottle at the end of the time period. This procedure allows a semi-continuous population to be maintained.

Samples of the yw *D. melanogaster* and the ver *D. simulans* were obtained from Barker. An initial experiment was set up with parental *D. melanogaster* percentages of 10%, 50% and 90%. Unexpectedly, the *D. melanogaster* (Barker 1) eliminated the *D. simulans* (*simulans* 1) almost immediately (Table 1). To check these results a sample (Chicago 1) of the original yw stock was obtained from W.G. Baker. A second experiment was performed which repeated the 10% level of the first and all three levels for the second strain. Again Barker 1 eliminated *simulans* 1, but surprisingly Chicago 1 was eliminated by *simulans* 1 at 10% and 50% and greatly reduced at 90%. New samples (Barker 2, *simulans* 2, and Chicago 2) were obtained from both sources and a third experiment was carried out. In this experiment both the Barker 2 and Chicago 2 samples remained the same or slightly decreased in frequency from parental frequencies over a seven week period.

Table 1. Per cent *D. melanogaster* (yw) observed from competition with *D. simulans* (ver) using a serial transfer method. Values are the means of three replicates. Several initial percentages were not conducted (N.C.) each experiment.

Experiment 1	Bottle begun on Day	Initial percentage yw Barker 1, <i>simulans</i> 1			Initial percentage yw		
		10%	50%	90%	10%	50%	90%
	0	45.6	97.0	98.4	N.C.	N.C.	N.C.
	3	49.8	93.5	100.0			
	7	59.2	95.9	100.0			
	10	86.2	98.6	-			
	14	93.4	-	-			
Experiment 2		Barker 1, <i>simulans</i> 1			Chicago 1, <i>simulans</i> 1		
		10%	50%	90%	10%	50%	90%
	0	67.4	N.C.	N.C.	8.9	31.8	100.0
	7	59.7			1.4	5.7	38.4
	14	95.8			4.8	5.6	89.7
	21	95.8			0.0	0.0	12.5
	28	98.8			0.0	0.3	22.0
Experiment 3		Barker 2, <i>simulans</i> 2			Chicago 2, <i>simulans</i> 2		
		10%	50%	90%	10%	50%	90%
	0	7.1	36.8	N.C.	N.C.	58.4	93.9
	7	11.4	57.5			80.2	94.7
	14	10.3	47.5			52.5	92.1
	21	16.7	44.4			79.4	80.6
	28	13.9	61.1			63.2	90.8
	35	26.8	27.7			60.4	86.2
	42	11.2	16.5			32.6	85.5
	49	9.2	21.4			29.6	78.8

The following hypothesis explains these results and is presently being tested. The

Barker 1 sample became adapted to the different conditions in my laboratory (a different food and the change from vials to bottles are probably the most dramatic differences) before experiment 1 was conducted. This adaptation increased the "competitive ability" of Barker 1 so that it quickly eliminated simulans 1. Simulans 1 also adapted in some manner so that it could outcompete the fresh Chicago 1 sample in experiment 2. The fact that experiment 3 results were as initially predicted was because all samples were obtained simultaneously and the experiment was run immediately after they were obtained. Further support of this hypothesis comes from Barker (pers. comm.) who has found in a second one-generation test that his stock in a recent retest produced approximately 40% *D. melanogaster* from 50% *D. melanogaster* parents, a value very close to his earlier results and quite unlike mine in experiments 1 and 2.

Grossfield, J. Purdue University, Lafayette, Indiana. A non-heuristic attribute of the ERG.

In conversation with several colleagues the question of electroretinograms (ERG's) being used to trace evolutionary patterns in the genus has arisen. I wish to point out that all species of *Drosophila* tested to date have the

same waveform and time course when ERG's are recorded under comparable conditions. This has held true over the past few years when ERG's have been cursorily checked in this laboratory for members of the *melanogaster*, *obscura*, *virilis*, *quinaria*, *robusta*, and *annulimana* species groups. Indeed, on the basis of available information, the same is true for all Diptera, with flies such as *Calliphora* and *Musca* showing larger amplitude responses. The presence of screening or accessory pigments in the eyes of various species may change the sensitivity of the response somewhat but that would be the maximal effect expected.

Hall, J.C. University of Washington, Seattle, Washington. The failure of two alleles of *c(3)G* to increase frequencies of X-linked lethals.

Green (Mut. Research 10:353, 1970) has discovered a putative allele of the recombination-deficient mutant, *c(3)G*, picked up as a mutator gene. It is a third chromosome semidominant whose locus is absent from *Df(3R) sbd*<sup>105</sup> -- as is true

for *c(3)G*. The frequencies of mutations at certain X chromosome loci are increased in the presence of the mutator gene in females. In addition, recombination is somewhat reduced by this mutant (M. Green, personal communication). Green's preliminary allelism tests show that females bearing the mutator and *c(3)G* in heterozygous condition do not generate mutations in the relatively high frequency found for the mutator in homozygous condition. This means that a) *c(3)G* and the mutator are alleles, but *c(3)G* is not a mutator; or b) *c(3)G* may or may not be a mutator, but it is not an allele of Green's mutant.

Both *c(3)G* and *c(3)G*<sup>68</sup> -- a newly arisen allele of this meiotic mutant (mei-W22 of Sandler, DIS 47, 1971, in press) -- have been directly assayed for possible mutator properties. Parry (Ph.D. Dissertation, University of Washington, 1970) found that a meiotic mutant which lowers recombination and increases nondisjunction generates increased frequencies of sex-linked lethals. Such sex-linked lethal tests were carried out for the two alleles of *c(3)G*, in which the treatment of X chromosomes consisted of passage of these X chromosomes through females homozygous for either meiotic mutant. For *c(3)G* only one of 473 treated X chromosomes carried a lethal. And for *c(3)G*<sup>68</sup> none of 553 X's had a lethal induced. In a control, seven of 931 X's recovered from *c(3)G*<sup>+</sup>/*c(3)G*<sup>+</sup> females bore a lethal. Four of these lethals arose from one female, and three from another, so the seven lethals probably represent only two mutations, each of which occurred at an oögonial stage and was proliferated. In any event, *c(3)G* does not appear to be a mutator gene.

Samples of the X chromosomes passed through these three kinds of females were examined for the presence of half chromatid (mosaic) lethals (produced in high frequency by chemical mutagens -- e.g. Carlson and Southin, Genetics 48:663, 1963). If *c(3)G* were generating increased frequencies of such half chromatid lethals in meiosis, after chromosome replication, they would go undetected among the F<sub>1</sub> males in a sex-linked lethal test (defining the P generation mothers as those bearing an X chromosome balancer and an X from a *c(3)G* female). However, of 87 X's from *c(3)G*, 70 from *c(3)G*<sup>68</sup>, and 292 from *c(3)G*<sup>+</sup>, none was found (in the F<sub>2</sub>) to have originally carried a half chromatid lethal.



Stockert, J.C. Centro de Investigaciones sobre Reproducción, Facultad de Medicina, Buenos Aires, Argentina. Alteration in the spermatocyte nuclei of a testis of *Drosophila hydei*.

Hess 1969), in one testis (Fig. 1-a), while the other one showed a very heterogeneous aspect in which normal spermatocytes and spermatocytes with an abnormal organization of the nuclear structure participated (Fig. 1-b, c, d).

The most prominent characteristics of these cells were the following:

1) A variable reduction in the size of the peripheral (attached to the nuclear envelope) nucleolus.

2) The occurrence of 1 or more (sometimes 4 or 5) free spherical bodies without the appearance of any normal loop, but quite similar in optic density and structure to the peripheral nucleolar mass. The size of these free nucleolus-like bodies was variable, and

Figure 1-a: normal spermatocyte; b,c: abnormal nuclei of spermatocytes, with free nucleolus-like bodies (arrows). Living spermatocytes, phase contrast. -d: schematic representation of one of these altered nuclei. Th: compact threads; PN: pseudo-nucleolus; Co: cones; N: nucleolus; FN: free nucleolus-like bodies; C: clubs; T: tubular ribbons.

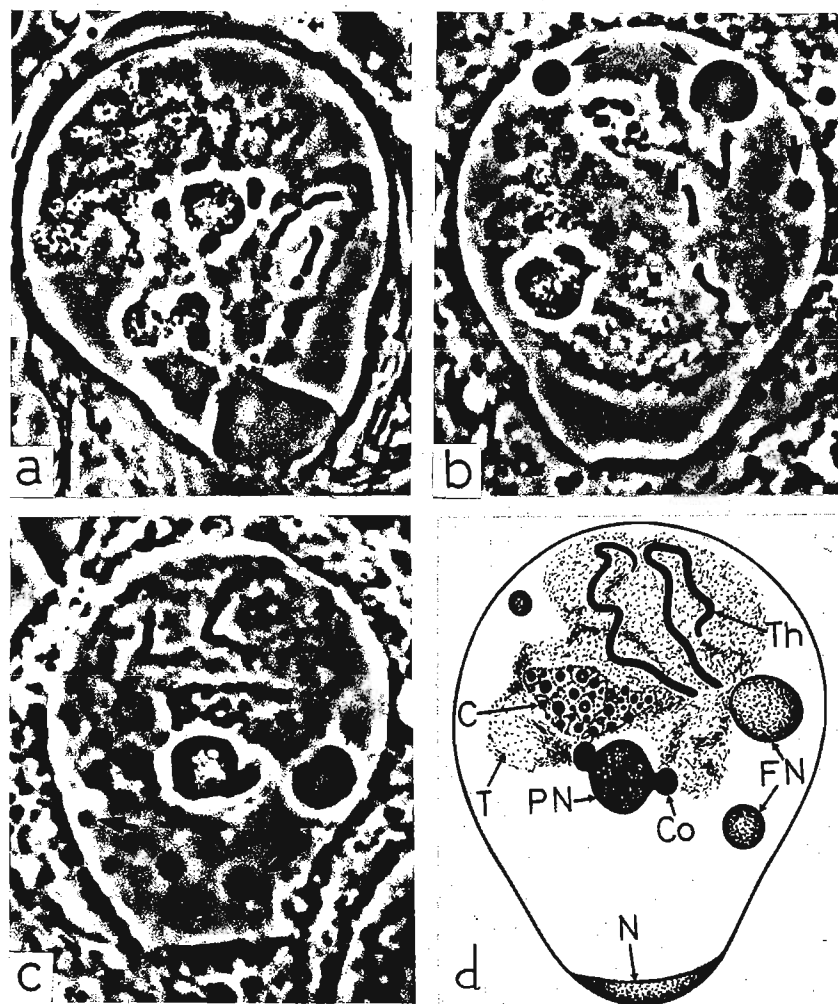
when it did not appear, the peripheral nucleolus was normal in size.

3) Frequently altered orientation of some loops, i.e. the compact threads or the clubs that were bound to one or more of the free nucleolus-like bodies, instead of the normal attachment to the peripheral nucleolus. However, in other cells we also observed a normal

orientation of these loops in spite of the presence of free nucleolus-like bodies.

4) The total absence of loops in some nuclei of fully grown spermatocytes, and the presence of duplicate loops (i.e. two pseudonucleoli) in a few other nuclei.

Unfortunately we could not obtain more data about this testis. A possible interpretation of the cytologic abnormalities we have found could be based on the existence of a fragmentation of the Y chromosome in some cells, in early stages of development of this testis. If



each nucleolar mass has its own organizer region (belonging to chromosomes, see Hay 1968, or free and segregated from the chromosomes, see Miller and Beatty 1969), it seems logical to assume that the free nucleoli to which certain loops were attached could contain portions of the nucleolar organizer near the loci of the loop forming sites in the Y chromosome.

References: Hay, E.D., 1968, Structure and function of the nucleolus in developing cells. In: Ultrastructure in Biological Systems. The Nucleus. (A.J. Dalton and F. Haguenau, Eds.) Acad. Press, N.Y. and London; Hess, O., 1969, Genetic activities of the Y chromosome in *Drosophila*. Ann. Embryol. Morphog., suppl. 1: 165-176; Hess, O. and G.F. Meyer, 1968, Genetic activities of the Y chromosome in *Drosophila* during spermatogenesis. Adv. Genetics 14: 171-223; Miller, O.L. and B.R. Beatty, 1969, Visualization of nucleolar genes. Science 164: 955-957.

Laughnan, J.R., S.J. Gabay and I.N. Montgomery. University of Illinois, Urbana, Illinois. Genetic basis for the exceptional events in Dp(1;1)MNB-8 *Drosophila melanogaster* males.

Males carrying Dp(1;1)MNB-8, designated the modified long duplication (MLD), and having the genotype  $f(B^+ os^+)(B os)car$  (duplicated members in parentheses), are  $B os^+$  in phenotype. When mated with attached-X females, patroclinous sons are mainly of the  $B os^+$  (parental) class. Not infrequently, however,

three classes of exceptional sons are produced. The distribution of exceptional offspring among progeny of single pair matings of the above type indicates that the exceptional event occurs almost exclusively in germinal tissues of the MLD parent and that it often takes place at a relatively early stage in the development of germinal elements. Analysis of salivary gland chromosomes reveals that the exceptional events involve a loss of portions of the duplication.-----Genetic analyses (Gabay and Laughnan, 1970) of progenies of MLD male parents from six different strains indicate a striking variation in overall frequency of exceptional events, and in the relative frequencies of the different kinds of exceptions. There have also been instances of stabilization within sublines of strains characterized by a high frequency of exceptional events, and of changes from a relatively stable to relatively unstable or active condition.-----The existence of stable and unstable MLD strains, and the strong tendency for these traits to be inherited through many generations, suggest a genetic control over the exceptional event. The particular mating system we employ, and the fact that, except for sudden changes of the type noted above, the various strains have, over many generations, retained the differences in frequency of exceptional events which they exhibit, make it unlikely that genetic control resides in either the autosomes or in the Y chromosome. On the other hand, since the duplication-bearing X chromosome of an MLD male parent is passed from father to son in each mating cycle, it seems most likely that if exceptional events are under the control of a chromosomal gene, the latter is located in the X chromosome. This hypothesis was tested using marked females that derive one duplication-X chromosome from an MLD stock characterized by a relatively high frequency of exceptional events, and another duplication-X chromosome from an MLD strain that is stable in this regard. These  $f$  "unstable" /  $f^+$  "stable" females were mated with wild-type males and  $f$  and  $f^+$  MLD sons were test mated with attached-X females to search for patroclinous exceptional sons. Among the 550 progenies from matings involving the  $f$  MLD sons, 205, or 37 per cent, had one or more exceptions, while, in similar matings, 458  $f^+$  MLD sons produced only seven, or 1.5 per cent, progenies with exceptions. These results indicate that genetic control of the exceptional event is carried in the X chromosome. Since forked, the marker used here to screen for sons carrying the X chromosomes from the unstable and stable MLD sources, is close to the distal end of Dp(1;1)MNB-8, and since the screen proved to be highly effective in identifying unstable and stable X chromosomes in the test matings, it appears that the genetic element in control of exceptional events is at a site in, or not far removed from the duplication itself. As noted above, the exceptional events involve a loss of chromosomal material from the duplication; moreover, the array of deficiency types among exceptions from the MLD strains suggests that there are characteristic hot spots for breakage in these strains. Hence there is no reason to assume a separate, closely-linked controlling element in the X-chromosome. For the time being it is sufficient to consider that the X chromosome of an unstable MLD strain differs from that of a stable strain in carrying within the duplication two or more sites that are highly susceptible to breakage and consequential loss of specific chromosomal segments.

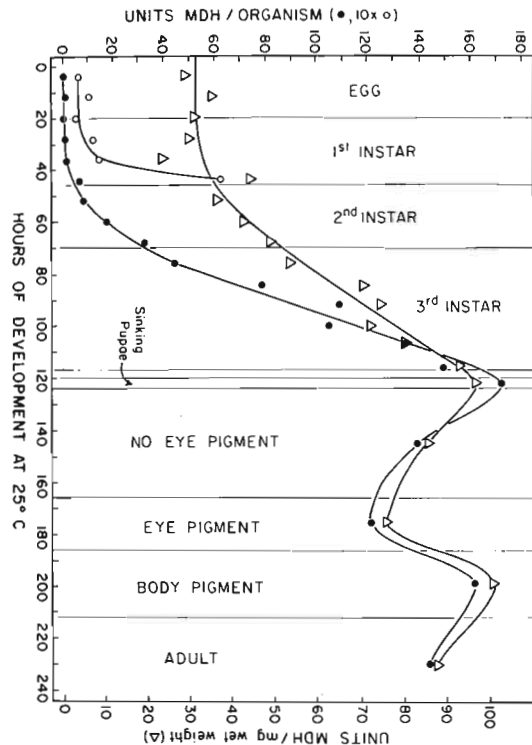
This work was supported by National Science Foundation Grant GB-7635.

Reference: Gabay, S.J. and J.R. Laughnan, 1970, Genetics 65: 249-265.

Anderson, M.\* The Johns Hopkins University, Baltimore, Maryland. Variations in the level of malate dehydrogenase during development.

pupae, have been described (Rechsteiner, 1970). At various times, eggs and/or larvae were collected by washing them into fine mesh nylon net with distilled water. An aliquot was counted and weighed; the remainder of the organisms was weighed, and the number estimated from the weight of the counted sample. They were homogenized at 100 mg/ml in 0.2 M sodium phosphate, pH 6.4, 0.1% in phenylthiourea. The homogenates were centrifuged for 20 minutes at 17,000 rpm, and the MDH activity of the supernatant solution was immediately determined. For pupal points, pupae were washed from the walls of an established bottle, staged by morphological characteristics, and counted, weighed, homogenized, and assayed. The pupae were found to be significantly lighter than the oldest larvae, presumably because they were raised under more crowded conditions on the standard food. To correct for this in determining enzyme units/organism, the experimentally determined enzymatic activity for the pupal points was multiplied by the ratio of the weight/organism of the oldest larvae to that of the youngest pupae. That the difference in body weight is not merely a reflection of differences in body water, and that some sort of correction factor need be applied, is shown by the fact that enzyme activity/mg. wet weight and enzyme activity/mg. extracted protein were similar for the oldest larvae and youngest pupae, while enzyme activity/organism and wet weight/organism in the pupae were about half of the larval figure.

MDH was assayed at 30°C on a Zeiss monochromator equipped with a Gilford Absorbance Recorder. Two to 25  $\mu$ l. of homogenate were added to 3 ml. of an assay mixture containing 16.7 mM malate, 2 mM NAD<sup>+</sup>, 50 mM glycine, pH 10.0, and the initial rate of change in O.D. at 340 nm. was determined.



The variation in the level of malate dehydrogenase (L-malate: NAD oxidoreductase, EC 1.1.1.37) during development has been studied in Oregon RCH flies. The techniques used to obtain eggs of known age, to raise the eggs and larvae, and to stage

It is apparent that the level of MDH in *Drosophila* undergoes stage-specific variations during development which are regulated independently of total soluble protein. Other *Drosophila* dehydrogenases show variations which are similar, but not identical. The levels of MDH are approximately constant up to hatching. This has been observed in alcohol dehydrogenase (unpublished observations), isocitrate dehydrogenase (Fox, 1971),  $\alpha$ -glycerolphosphate dehydrogenase (Rechsteiner, 1970), and glucose-6-phosphate dehydrogenase and 6-phosphogluconate dehydrogenase (Wright and Shaw, 1970). However,  $\beta$ -hydroxybutyrate dehydrogenase (unpublished observations) and lactate dehydrogenase (Rechsteiner, 1970) show seven- and twenty-fold increases, respectively, in enzyme units per organism during embryogenesis. The drop in MDH activity per organism during metamorphosis (30%) is much less than that observed for alcohol dehydrogenase (500%),  $\beta$ -hydroxybutyrate dehydrogenase (900%) (unpublished observations), and  $\alpha$ -glycerolphosphate dehydrogenase (300%, Rechsteiner, 1970), but similar to that seen in NADP<sup>+</sup>-dependent isocitrate dehydrogenase (40%, Fox, 1971). The measured activity of MDH represents the sum of s-MDH and m-MDH. Separate measurement of the two during development would provide interesting data on the control of the levels of the forms of MDH. If the two

are not regulated coordinately, this could bear on the significance of the existence of two forms of the enzyme. E.g., different cell types, or a single cell type at different times

in the life cycle of the cell or organism, could require a different balance between the mitochondrial and supernatant forms of the enzyme. Experiments in other systems (Kitto, 1967; Kitto and Lewis, 1967), including *D. virilis* (McReynolds and Kitto, 1970), indicate that antibodies to the two forms of MDH do not crossreact; hence immunologic techniques should allow one to measure the activities of the two forms separately during development.

References: Fox, D.J., 1971, *Biochem. Genet.* 5:69-80; Kitto, G.B., 1967, *Biochem. Biophys. Acta* 139:16-23. Kitto, G.B., and R.G. Lewis, 1967, *Biochem. Biophys. Acta* 139: 1-15; McReynolds, M.S., and G.B. Kitto, 1970, *Biochem. Biophys. Acta* 198:165-175; Rechsteiner, M.C., 1970, *J. Ins. Physiol.* 16:1179-1197; Wright, D.A., and C.R. Shaw, 1970, *Biochem. Genet.* 4:385-394.

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Parzen, S.D., M.J. Kessenich, and A.S. Fox.  
University of Wisconsin, Madison, Wisconsin.  
A method for the preparation on high molecular weight DNA from adult *D. melanogaster*.

Twenty-five grams of flies (wet weight) are ground with 120 ml of cold absolute methanol in an all-glass homogenizer in an ice bucket. The homogenate is centrifuged at 12,000 Xg for 10 minutes at 4° C, the supernatant discarded, and the pellet

reground in 120 ml of a solution containing 0.15 M NaCl, 0.015 M sodium citrate, and 0.05 M EDTA at pH 7.0 (Solution No. 1). This is centrifuged as before and the pellet is ground again in 120 ml of Solution No. 1, followed by another centrifugation. After this centrifugation the pellet is suspended in 40 ml of 0.1 M NaCl, and 40 ml of 5% Aerosol O.T. in 0.1 M NaCl is added slowly with gentle stirring. The suspension is placed in a water bath at 50° C for one hour. It is then allowed to cool to room temperature and sufficient solid NaCl is added to raise the salt molarity to 1.0 M. After 10 minutes the preparation is centrifuged at 12,000 Xg for 10 minutes at 4° C. The precipitate is now discarded and one volume of cold 2-ethoxyethanol is added to the supernatant. This is placed in a freezer for 15 minutes, and is then centrifuged for 10 minutes at 10,000 Xg.

The precipitate is dissolved in 20 ml of 0.15 M NaCl, and is deproteinized by shaking vigorously for 10 minutes with an equal volume of a solution containing 24 parts chloroform to 1 of isoamyl alcohol. It is then briefly centrifuged to separate the phases, the aqueous phase is removed, and to the aqueous phase is added two volumes of cold ethanol. The DNA is now spooled out on a glass rod and dissolved in 9 ml of 0.015 M NaCl. When it is completely dissolved, 1 ml of 1.5 M NaCl is added.

Removal of contaminating RNA is now carried out by the addition of 1.0 ml of a 0.2% RNase solution (prepared in 0.1 M TRIS, pH 7.6, and heated for 10 minutes at 80° C to inactivate contaminating DNase). This is allowed to incubate in a 37° C water bath for one to three hours. After this, the solution is deproteinized as before. The DNA is then precipitated with either ethanol or 2-ethoxyethanol, deproteinized again, and precipitated once more with either ethanol or 2-ethoxyethanol. Following the last precipitation it is dissolved in 9 ml of 0.015 M NaCl + 1.0 ml 3.0 M NaAcetate containing 0.001 M EDTA, pH 7.0. It is then precipitated by the dropwise addition of 0.60 volumes of cold isopropanol. The DNA is spooled out and dissolved in 9 ml of 0.015 M NaCl + 1.0 ml 3.0 M NaAcetate containing 0.001 M EDTA, pH 7.0, and precipitated by the addition of 0.60 volumes of cold isopropanol. This final precipitate is dissolved in 10 ml of 0.15 M NaCl and is then typically chromatographed on a Sepharose-4B column, eluting with 0.15 M NaCl, collecting and pooling those fractions coming off the column immediately after the void volume.

Orcinol tests for RNA are negative. Melting point determinations show a  $T_m = 83.6^\circ \text{C}$  with a hyperchromic effect of 40-45% in 0.15 M NaCl.  $A_{260 \text{ m}\mu} / A_{280 \text{ m}\mu} = 1.8 - 2.0$ . Sedimentation equilibrium studies in CsCl show a single narrow main band at  $\rho = 1.702$  with a small shoulder at  $\rho = 1.687$ . Prior to Sepharose-4B chromatography an additional smaller but wider band is present at  $\rho = 1.675$ . Yield has been as great as 2.5 mg DNA from twenty-five g of flies.

Supported by the following grants from the National Institutes of Health, USPHS: GM-11777, GM-15422, and GM-00398.

Miklos, G. L. G. University of California, San Diego, California. SD distributions and the measurement of distortion.

In Segregation-Distorter (SD) experiments, K value distributions are often encountered which have seemingly unusual shapes, and a distribution can appear to be a composite of two different histograms.

It is sometimes found in such an experiment that many or most SD/SD<sup>+</sup> males exhibit very high distortion, whilst others show greatly reduced distortion or none at all. The interpretation of the shape of SD histograms can be approached using the make analysis of Miklos and Smith-White (Genetics 67:305-317, 1971). This communication presents the results of an experiment in which an unusual SD distribution is interpreted as having a high variance on the make scale. It further gives other odd shapes which can be generated by the make method, and the following examples extend the spectrum of histogram shapes initiated by the two high variance examples of Miklos and Smith-White (1971). The problems associated with the use of K value as a measure of distortion will also be clear from the examples.

A. During a series of experiments designed to investigate the phenomenon of conditional distortion, (Sandler and Hiraizumi, Genetics 46:585-604, 1961) an SD-72 chromosome was passed through females, and the SD-72/cn bw sons of the SD-72/cn bw mothers were tested for distortion in the standard way. Instead of obtaining the characteristic high K value distribution, in which all males yield values near 1.00, it was found that approximately 25 per cent of the males exhibited markedly reduced distortion. These results are shown in histogram A, in which the 65 tested males yielded 9,509 progeny. This type of histogram is frequently encountered in the SD literature, and it is due to a high level of distortion, together with large between-male variability.

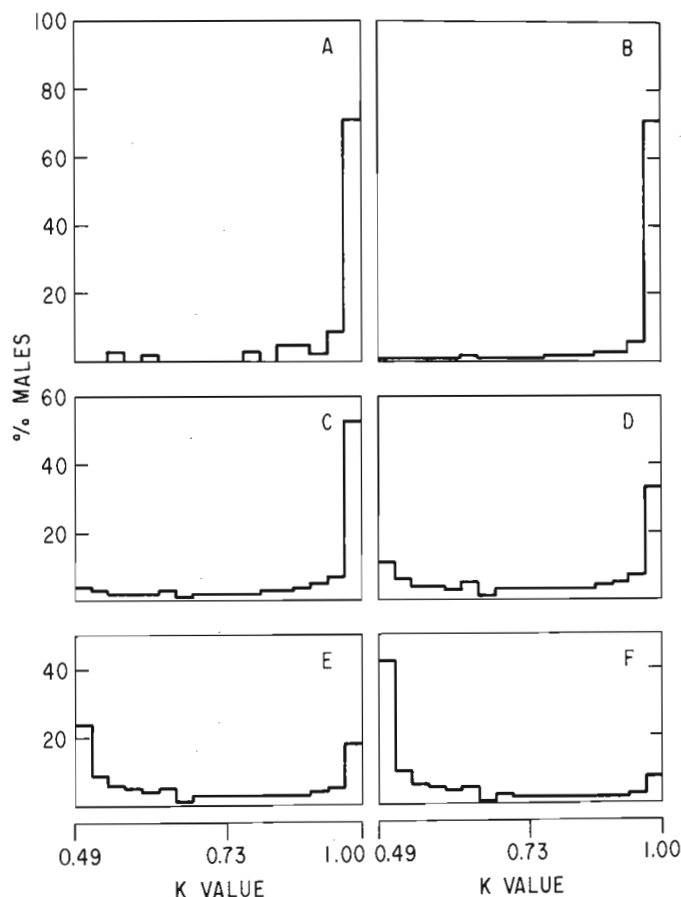
The theoretical histograms B,C,D,E and F have been derived from normal distributions on the make scale. They all have the

same high variance ( $V_M = 4.0$ ), and differ only in their means; B,C,D,E and F have means of 8,7,6,5 and 4 Probits respectively, corresponding to K values of 0.999, 0.98, 0.86, 0.67 and 0.54. It can be seen that the experimental histogram A corresponds to a theoretical one such as B, which has a high mean and a large underlying variance. The level of distortion of the SD-72 stock has thus remained unaltered during the passage of the SD-72 chromosome through females, however the between-male variance has increased. If the level of distortion is lowered, but a high between-male variance is retained, then SD distributions such as B,C,D,E and F are obtained. Histograms such as these are relatively abundant in the SD literature, and most probably occur when some form of heterogeneity is introduced into the SD stock.

It should be pointed out that if an SD distribution is highly skewed, its mean and median can differ by a large amount, and the mean will provide an inappropriate measure of distortion because it is influenced by the skewness. The median is unaffected and should be utilized in such cases.

B. These results demonstrate the deceptive nature of K value distributions. Another property is that K is

not uniform over its range of measurement. The difference in distortion between K's of 0.95 and 0.90 is not the same as between 0.70 and 0.65 for example. It takes the same amount of "work" to go from K = 0.67 to K = 0.86 (ie, 5 Probits to 6 Probits), as it does to go from



$K = 0.98$  to  $K = 0.999$  (7 Probits to 8 Probits). Thus changes in distortion in terms of  $K$  value can be misleading, and an unambiguous description of an SD experiment involves describing means, variances and changes in Probits.

The difficulties are increased further when tests of significance are involved. Owing to the nature of the  $K$  scale, two distributions may appear to be significantly different under standard statistical tests, but in reality, are not significantly different when compared in Probits. This is not due to a deficiency in the test itself, but rather to the unsuitable analytical properties of the  $K$  scale.

These results on the shape of SD histograms and the nonuniform changes in  $K$  over the range of distortion, further show that the phenomenon of Segregation-Distortion is best considered in terms of the make analysis.

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Sandler, L. University of Washington, Seattle, Washington. Induction of autosomal meiotic mutants by EMS in *D. melanogaster*.

A scheme for the detection and isolation of autosomal meiotic mutants (i.e. mutants on either chromosome 2 or 3 that affect disjunction of either the sex chromosomes or chromosome 4 in either

sex or X chromosome recombination in females) has been described by Sandler et al. (Genetics 60:525-558, 1968). They examined autosomes collected from natural populations.

This scheme has now been applied to mutagenized major autosomes. Chromosomes 2 and 3 were recovered from Canton-S males treated with EMS according to the method of Lewis and Bacher (DIS 43:193, 1968) using a treatment that produced about 10% (54/545) sex-linked recessive lethals after one additional backcross generation to resolve mosaics. The scheme of Sandler et al. for examining autosomes was also modified to resolve mosaics.

There were 35 lethal-free 2-3 complements examined for meiotic mutants -- 24 in both sexes, 8 in females only (the males were sterile) and 3 in males only (the females were sterile). Among these, two meiotic mutants were recovered: (1) mei-W5, a second chromosome recessive that causes the production, in homozygous males, of sperm lacking paternal chromosomes and has no obvious effects in females, and (2) mei-W22, a third chromosome recessive that eliminates recombination and increases nondisjunction in homozygous females and is sterile in males (for reasons not yet investigated).

From the cross,  $\text{In}(1\text{LR})\text{sc}^{\text{VI}}$ ,  $y \text{ pn } v \cdot y^+/y$ ;  $\text{spa}^{\text{pol}}/\text{spa}^{\text{pol}}$  females homozygous for the indicated meiotic mutant by  $Y^{\text{S}}X \cdot Y^{\text{L}}$ ,  $\text{In}(1)\text{EN}$ ,  $v \text{ f } B/O$ ;  $C(4) \text{ RM}$ ,  $ci \text{ ey } R/O$  males, there were observed:

mei-W5				mei-W22			
X chromosome	chromosome 4			X chromosome	chromosome 4		
	+	pol	ey		+	pol	ey
(v,v <sup>+</sup> )B/+♀♀	97	0	0	(v,v <sup>+</sup> )B/+♀♀	51	2	5
B <sup>+</sup> ♀♀	0	0	0	B <sup>+</sup> ♀♀	17	4	3
v f B ♂♂	0	0	0	v f B ♂♂	14	7	9
pn v ♂♂	26	0	0	pn v ♂♂	32	3	8
y ♂♂	20	0	0	y ♂♂	37	2	3
y pn ♂♂	6	0	0	y pn ♂♂	0	0	0
v ♂♂	9	0	0	v ♂♂	0	0	0
y pn v ♂♂	11	0	0	y pn v ♂♂	0	0	0
+ ♂♂	11	0	0	+ ♂♂	0	0	0
pn ♂♂	2	0	0	pn ♂♂	0	0	0
y v ♂♂	2	0	0	y v ♂♂	0	0	0

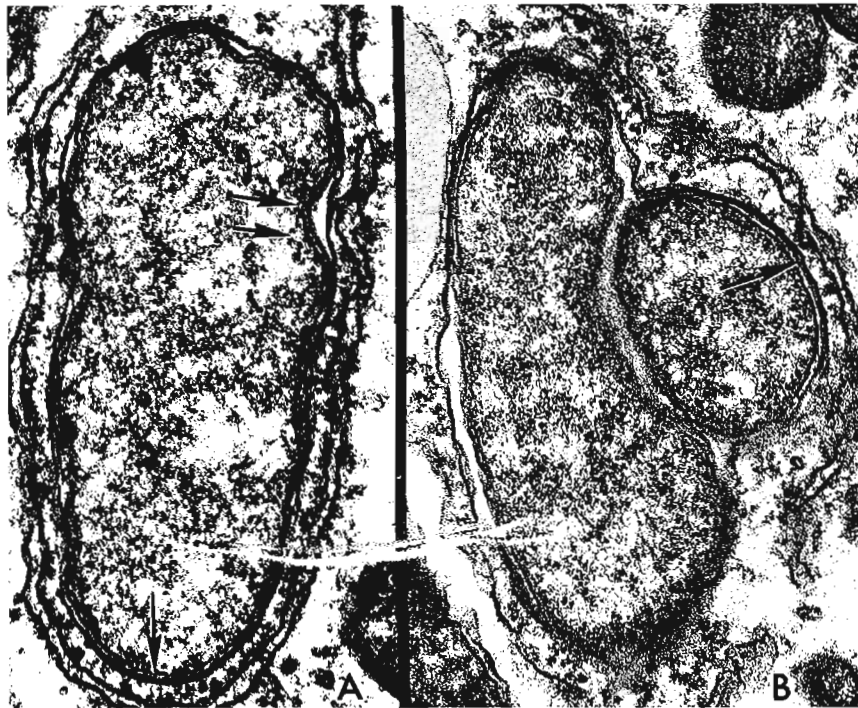
From the cross,  $\text{In}(1)\text{FM6}$ ,  $y^{3\text{ld}} \text{ sc}^8 \text{ dm } B/y^+Y$ ;  $\text{spa}^{\text{pol}}/\text{spa}^{\text{pol}}$  males homozygous for mei-W5 by  $y \text{ pn}/y \text{ pn}$ ;  $C(4) \text{ RM}$ ,  $ci \text{ ey } R/O$  females, there were observed:  $y^2 B \text{ ♀♀} = 213$ ,  $\text{pn } \text{ ♂♂} = 217$ ,  $B \text{ ♀♀} = 0$ ,  $y \text{ pn } \text{ ♂♂} = 9$ ,  $y^2 B$ ;  $\text{pol } \text{ ♀♀} = 0$ ,  $y^2 B$ ;  $ci \text{ ey } \text{ ♀♀} = 33$ ,  $\text{pn}$ ;  $\text{pol } \text{ ♂♂} = 0$ ,  $\text{pn}$ ;  $ci \text{ ey } \text{ ♂♂} = 28$ ,  $B$ ;  $\text{pol } \text{ ♀♀} = 0$ ,  $B$ ;  $ci$ ,  $ey \text{ ♀♀} = 0$ ,  $y \text{ pn}$ ;  $\text{pol } \text{ ♂♂} = 0$ , and  $y \text{ pn}$ ;  $ci \text{ ey } \text{ ♂♂} = 0$ .

Kernaghan, R.P. SUNY at Stony Brook, New York. The ultrastructure of the organism associated with hybrid sterility in *D. paulistorum*.

Crosses between some semispecies of *D. paulistorum* produce sterile  $F_1$  male offspring. Usually a cross between Mesitas females X Santa Marta males produce fertile  $F_1$  male progeny but in recent years the male progeny are sterile as if the ability

to produce sterile offspring has been acquired. At the electron microscope level, the testes of these males are congested with degenerating sperm and contain a microorganism or symbiont similar to that previously described by Kernaghan and Ehrman (1970). In addition, extracts prepared from such infected tissue are potent in inducing sterility in the sons of recipient females (Williamson, Ehrman and Kernaghan 1971).

High resolution analysis of testes of these sterile  $F_1$  hybrid males, as well as developing eggs of their fertile sisters and mothers shows a cytoplasmic membrane limited vacuole enclosing one or more of the symbionts. In addition, each microorganism is limited by two membranes. The outer membrane may be juxtaposed to the vacuolar membrane to produce a more electron dense region (Figure A & B). The internal structure of the organism is described as a reticulate network of fibers and may or may not be accompanied by a rough peripheral granulation of ribosome-like material. Pleomorphic forms are not unusual ranging from a smaller dense granular form .1  $\mu$  in diameter to the larger reticular form .3 to .5  $\mu$  in diameter.



Electron micrograph of the reticulate form of the microorganism in the sterile  $F_1$  male testis from a cross Mesitas females X Santa Marta males. Figure A, (upper arrows) show the duplicate membranes of the symbiont while Figure A, (lower arrow) and Figure B demonstrate the juxtaposition of the external membrane to the vacuolar membrane. 90,000X

Penicillin has no detectable effect on the ultrastructure of the symbiont. Both external membranes remain intact when such sterile  $F_1$  hybrid males are raised on drug treated media. Large reticulate forms are common in these treated individuals. In adult tissue the symbiont has been detected only in the gonads while in some cases larval and pupal gut muscle exhibit the microorganism.

On the ultrastructural level alone, no clear assignment may be made as to the type of organism involved. Although a general similarity to mycoplasma-like morphology exists, the ultrastructure of a Rickettsia described by Briton and Burgdorfer (1971) or the fine structure of Chlamydia by Tamura et al (1971) is equally applicable.



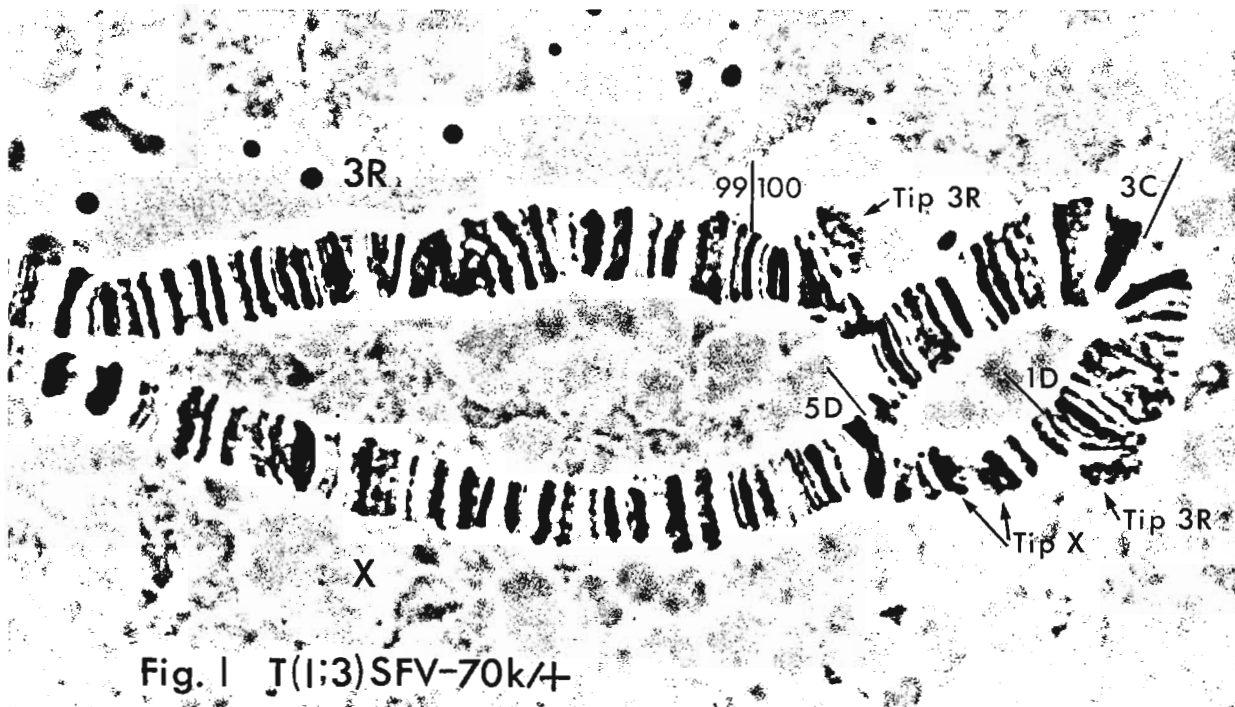
References: Kernaghan, P. and L. Ehrman 1970 Chromosoma 29:291-304; Williamson, D., L. Ehrman and P. Kernaghan 1971 P.N.A.S. (in press); Brinton, L.P. and W. Burgdorfer 1971 J. Bacteriol. 105:1149-1159; Tamara, A., Matsumoto, A., G.P. Manire and N. Higashi, 1971 J. Bacteriol. 105:355-360.

P. Kernaghan acknowledges support of Nih Grant #AI09945.

Lefevre, G., Jr. San Fernando Valley State College, Northridge, California. Crossing over in an insertional translocation.

A cytogenetic analysis of some EMS-induced sex-linked lethals unexpectedly revealed an insertional translocation in which a segment of X extending from 1D1-2 to approximately 5C5-6, i.e., from

su ( $w^a$ ) through cv, was inserted in direct order near the tip of 3R, just before 100E1. This translocation, designated as T(1;3)SFV-70k, is illustrated in Fig. 1. The aneuploid deficiency segregant is lethal as a heterozygous female; the duplication segregant survives as a fertile female, but is lethal as a male.



Because of the favorable orientation and location of the inserted material, an attempt was made to recover a single crossover between it and a normal, marked X. Although only a portion of such crossovers should be identifiable, a total of 4 were found among 2,551 daughters of T(1;3)/y<sup>2</sup> w<sup>a</sup> ec cv ct f females. Each of these recombinant daughters carried one T(1;3) chromosome in which the original insertional translocation had been converted by the single crossover into a reciprocal translocation. However, only a half-translocation was recovered in each recombinant fly. (Although the full reciprocal translocation can be recovered in a single individual, it should not be recognizable as a recombinant.)

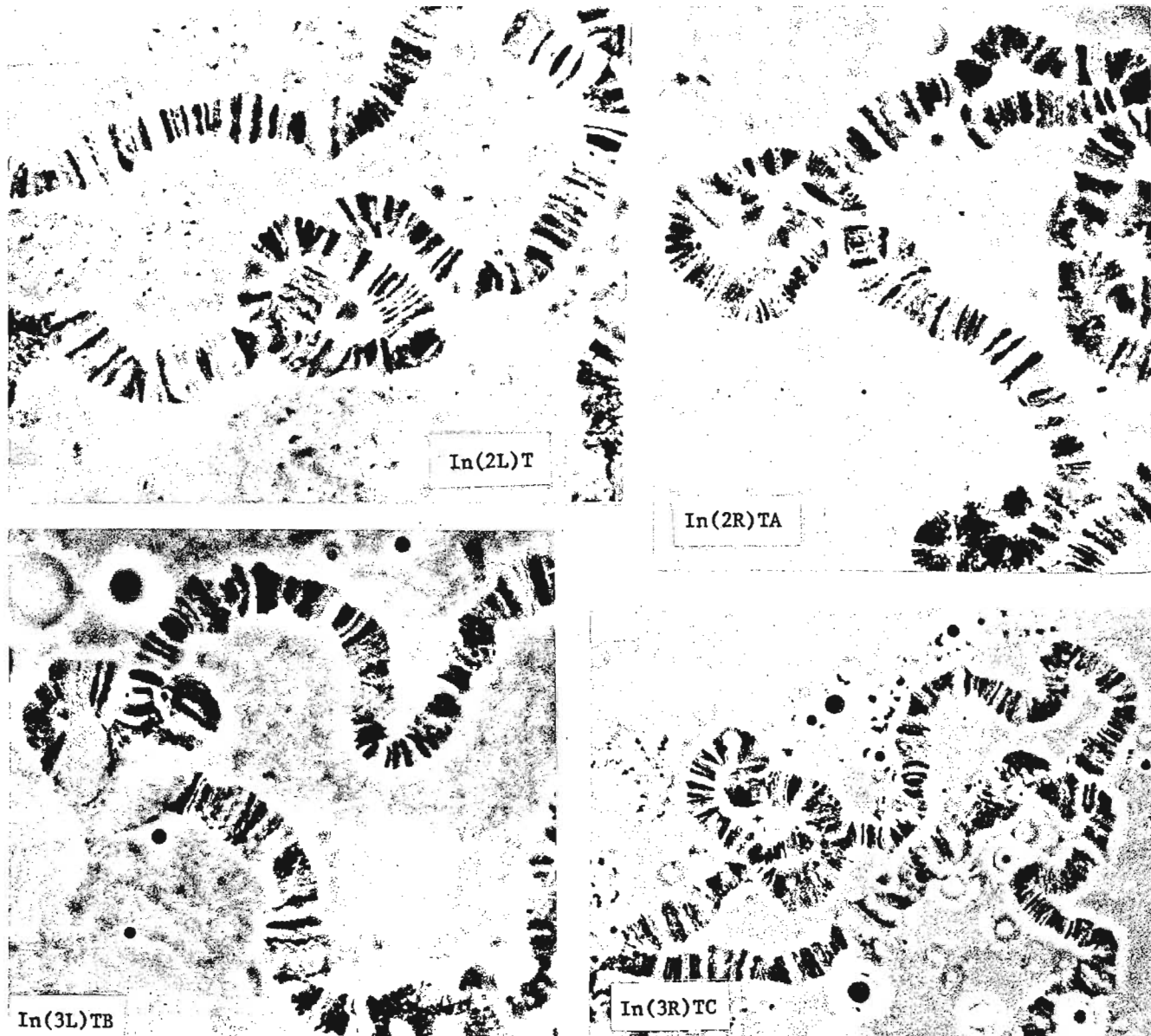
The successful recovery of these crossovers demonstrates that effective synapsis does not require a zipperlike action initiated only at the telomere or centromere, but is compatible with the view of von Wettstein (PNAS 68:851-855, 1971) that precise synapsis between homologous elements can be initiated at any point.



Yang, H., K. Kojima and A. Kovarik. The University of Texas, Austin, Texas. Inversions in a Southwest Texas *D. melanogaster* population.

Four new and eight cosmopolitan chromosomal inversions on Chromosomes II and III were found in a year-round population of *D. melanogaster* in Southwest Texas. The flies were collected near Brownsville, Texas in 1970.

Break-points in these inversions were identified on the basis of the salivary gland chromosome map by Bridges (1935). The eight cosmopolitan inversions identified were: In(2L)t, In(2R)NS, In(3L)P, In(3L)M, In(3R)C, In(3R)K and In(3R)MO.



The new inversions (Fig. 1) were found in each arm of the second and third chromosomes. In(2L)T is a small single paracentric inversion between 30F and 36D. In(2R)TA is another small paracentric inversion between 50A and 53A. In(3L)TB is a small inversion between 70B

and 75A. In(3R)TC is a large paracentric one which is located between 84D and 91E. The break-points of this In(3R)TC are similar to those of In(3R)Antp<sup>LC</sup> which were induced by neutrons.

The relative frequencies of occurrence of these inversions are as follows, in percentages:

In(2L)t = 14.5	In(2L)T = 0.3	In(2R)NS = 20.9	In(2R)TA = 0.3
In(3L)P = 19.1	In(3L)M = 1.0	In(3L)TB = 0.3	In(3R)TC = 0.3
In(3R)K = 0.3	In(3R)P = 38.2	In(3R)C = 3.9	In(3R)MO = 0.3

The total of the genomes extracted by the Cy/Pm; Ubx/Sb method was 233.

Benner, D.B.\* University of California, Riverside, California. Some evidence against the presence of suppressors of variegation on the Y chromosome.

Brosseau (1964) reports that the localized regions of the Y chromosome near the kl-2 fertility factor on Y<sup>L</sup> and proximal to ks-1 on Y<sup>S</sup> act as position-effect suppressors. I would like to present some evidence that suggests that there is no suppression of variegation by these localized regions.

The first evidence comes from an analysis of the Y-4R fragments reported by Parker (1965, 1967). The Dubinin effect (a position-effect variegation of the cubitus interruptus gene on the fourth chromosome) is observed only in those cases where 4R is located distal to kl-2. In these cases Y<sup>S</sup> is intact and the break in Y<sup>L</sup> is distal to kl-2 and therefore the presumed suppressor. The second evidence is of like nature and comes from a similar analysis of twenty Y-4R fragments that were produced using an unmarked Y obtained from a wild population. Sixteen of the fragments do not show the Dubinin effect and show no evidence of any of the KL fertility factors. The four remaining fragments have at least kl-1 and kl-2 present, and all four show the Dubinin effect.

The third bit of evidence that these specific regions of the Y may not be responsible for variegation suppression comes from an analysis of X detachments in which 4R and some portion of the Y from Parker's fragments have been attached to the X chromosome. C(1)RM, y v bb; ci ey<sup>R</sup> ♀♀ bearing the Y-4R, y<sup>+</sup> ci<sup>+</sup> ey<sup>+</sup> fragment were irradiated with 3Kr of X-rays within twelve hours of eclosion, mated to In(1)sc<sup>SIL</sup> sc<sup>8R+S</sup>, sc<sup>S1</sup> sc<sup>8</sup> w<sup>a</sup> B/Y; ci ey<sup>R</sup> ♂♂, and allowed to lay eggs for four days. y v; ci<sup>+</sup> ey<sup>+</sup> ♂♂ were recovered and put into stock by mating to C(1)RM, y v bb/Y; ci ey<sup>R</sup> ♀♀. In all cases where the fertility factors that were present in the fragment have been lost the Dubinin effect has been lost. This suggests that the loss of variegation is accompanied by loss of the region adjacent to kl-2 and the region proximal to ks-1. In those cases where the proximal region of Y<sup>L</sup>, and therefore the regions adjacent to kl-2, have not been disrupted the Dubinin effect persists. Likewise, in those cases where neither Y<sup>S</sup> nor Y<sup>L</sup> have been disrupted the variegation persists.

These results suggest that there is no region of the Y that specifically acts to suppress the Dubinin effect. The effect is absent when all or most of the Y is missing. It has been previously reported that the Y chromosome does not always act as a suppressor of the Dubinin effect and in fact may act as an enhancer of the effect (Panshin, 1938; Altofer, 1967). The results reported by Brosseau were obtained using a variegating B<sup>S</sup>.

The results reported here are consistent with Brosseau's conclusion that the heterochromatic Y does not suppress variegation. The discrepancy in the results concerning the action of specific sites on the Y as variegation suppressors may mean that suppression of a particular variegation is the result of an effect on that specific rearrangement by the Y and not the peculiar property of a specific region. In other words, the Y may have different effects on different rearrangements because of modifications in spatial associations within the nucleus (as suggested by Muller, 1935) not because of the special influences of specific regions.

References: Altofer, N. 1967 Genetics 55:755-767; Brosseau, G.E. 1964 Genetics 50: 237; Muller, H.J. 1935 Fifteenth Int. Physiol. Congr., Leningrad; Panshin, I.B. 1938 Biol. Zh., Mosk. 7:837-868; Parker, D.R. 1965 Mutation Res. 2:523-529, 1967 Mutation Res. 4:333-337.

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Nash, W.G. The George Washington University, Washington, D.C. Deep orange and carnation: Another lethal gene combination in D.m.

The dor car lethal gene combination resembles both dor ry (Lucchesi, DIS 39: 127) and ey eyg (Hunt, Gen. Res. Camb. 15: 29-34) in that the genes involved are all recessive whose combined action results in death of the pupa. Unlike the latter two lethal combinations it is rare for

dor car zygotes to develop beyond the early stages of the pupal period.

With the exception of having colorless malpighian tubules dor car larvae appear to develop in a completely normal manner. Shortly after pupating, however, normal development stops abruptly and is signaled by the formation of a large air bubble in the center of the pupa. Since larval development appears normal it seems likely that the lethal biochemical lesion is affecting some or all of the adult imaginal disks.

This hypothesis is tested using a special stock which produces gynandromorphs with a high frequency. Certain female zygotes in this stock have an unstable ring X chromosome ( $X^{c2} \text{In}(1)w^{VC}, w^{VC}, f$ ) which may be lost at the first meiotic division in the egg or at any later nuclear division. The genes on the remaining X chromosome are thus unmasked in a hemizygous condition. For example, dor car/ $X^{c2} \text{In}(1)w^{VC}, w^{VC} f$  female zygotes will express the dor car phenotype in any tissues which are derived from a cell line having lost the unstable ring X chromosome.

The mating scheme used to produce the type of female zygote in the above example follows.

Balanced lethal stock  
 FM-6,  $y^{3ld} \text{dm B/y dor car} \times \text{FM-6, } y^{3ld} \text{dm B}$   
 FM-6,  $y^{3ld} \text{dm B/y dor car} \times X^{c2} \text{In}(1)w^{VC} w^{VC} f/Y$   
 Type A ♀  $y \text{ dor car}/X^{c2} \text{In}(1)w^{VC}$   
 Type B ♀ FM-6,  $y^{3ld} \text{dm B}/X^{c2} \text{In}(1)w^{VC} w^{VC} f$

The yellow alleles in type A and B females makes it possible to recognize any male cuticular tissues in the resulting gynandromorphs. In type A zygotes dor car male tissue should result

Table 1. A comparison of gynandromorphic tissue patterns between type A and type B female zygotes.

	Total No. of flies	Total No. of gynandromorphs	Hemizygous tissue found in gynandromorphs		
			Head	Thorax	Abdomen
Type A zygotes	624	24	14	6	9
Type B zygotes	640	69	36	42	38

only from imaginal disks in which this gene combination is viable. Type B zygotes act as controls in the sense that any gynandromorphic patterns possible should be unrestricted in their appearance among these zygotes. The results of this experiment are shown in Table 1.

Of the 69 gynandromorphs which appear among the type B adult progeny, the hemizygous FM-6 chromosomal markers express themselves randomly in the head, thorax and abdominal tissues. Only 24 gynandromorphs appeared among the type A adult progeny. This might result from a random killing tendency among all kinds of type A gynandromorphs or from the specific elimination of certain kinds of gynandromorphic patterns. The latter explanation appears to be correct in that complete bilateral gynandromorphs are common in type B adults but absent among type A adults. It is also interesting that in the six cases where the whole head of type A adults is dor car no part of the thorax or abdomen is dor car. If only half of the head is dor car then parts of the thorax are sometimes also dor car. The kinds of gynandromorphic patterns among type A adults suggest that the dor car gene combination is lethal due to an abnormal interaction between the different regions of the fly and not due to the lack of development of any single region.

Bennett, J. and J.F. Hughes. Northern Illinois University, DeKalb, Illinois. Behavioral correlates of the  $w$ ,  $w^+$  gene substitution, observations without ether.

dish type observation chambers. In this study ether was dispensed with and smaller more convenient observation chambers were utilized (Figure 1).

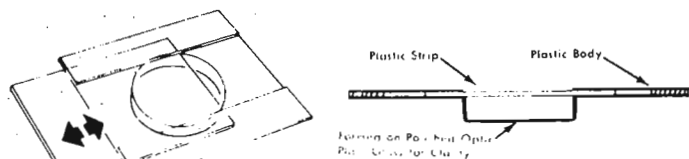


FIGURE 1

This procedure made anesthesia unnecessary and allowed direct observations without the ether trauma involved in the earlier study. A preliminary effort with Nitrogen anesthetization indicated that it, too, was traumatic in that many flies emerged with wings locked over their backs and did not recover for several hours. Thus complete avoidance of anesthesia seemed preferable.

Preliminary observations revealed a somewhat different set of characteristic behavioral patterns. Fourteen, many similar to the earlier study, were selected to record. One hundred flies of each sex and of each line were observed for 10 minutes each. An observed behavior pattern was recorded only once in each observation period. Flies were generally from two to four days of age and from stock cultures.

Of the observed traits five displayed significant differences in frequency between lines, or sexes, or both. Rubbing Forelegs: The fly stands on rear four legs and rubs front legs together rapidly. Rubbing Antenna: The fly used one front leg to brush antenna from base to tip many times in rapid succession. Rubbing Head and Eyes: The fly turns head and brushes top of head and eye and the neck region with one foreleg. Combs Wings: The fly stands on five legs, depresses tip of abdomen, combs top of wing with rear leg on the same side. Pulls Anus: The fly stands on front four legs uses rear pair of legs to pull anal region posteriorly.

The table indicates the observed numbers and the  $\chi^2$  probabilities for the differences, either between lines  $w^+$  &  $w$ , or between sexes (across lines). In addition an activity value

Line & Sex	Rub Forelegs	Rub Antenna	Rub Head	Combs Wing	Pulls Anus
$w^+$ ♂	98	87	83	75	1
$w^+$ ♀	100	77	78	78	1
$w$ ♂	92	77	80	72	1
$w$ ♀	87	59	55	62	14
$w$ vs $w^+$	<0.001	<0.004	<0.002	<0.04	<0.004
♂ vs ♀		<0.001	<0.002		

was obtained by summing all recorded activities. This score showed significant difference between the sexes but not between lines.

The wing combing behavior is closely similar to the earlier study. There does not seem to be any other parallelism with the behavior following etherization. In either case it appears that the substitution of the  $w$  gene for its normal allele does result in a number of measurable changes in behavioral tendencies. In several of the measures it appears to result in a lessening of the probability of the activity, and usually a more extreme reduction among males than females.

A pair of isogenic, inbred, Oregon-R lines, differing only at the white locus, were examined for behavioral differences. The lines were described more fully in DIS 45: 140-141. In the earlier study ether was used to separate the flies and place them in the small petri-dish type observation chambers. These chambers (#BGS4, 16mm dia. x 3mm deep, \$3.75/100, The Blister Co., 845 3rd Ave. East, Kalispell, Montana 59901) known as Blister™ slides, have sliding covers over a circular, flat bottomed cavity. It is possible to place a chamber at the end of a vial, shake the fly into the cavity and slide the cover in place before removing the vial. When turned over under a stereoscope the chamber is easily visible and fills the field at 10X magnification.

Hoenigsberg, H.F. Universidad de los Andes, Bogotá, Colombia. A new environmental variable that changes the rate of development of some members of the willistoni group of species.

The usual laboratory conditions for *Drosophila paulistorum* cluster of species (Dobzhansky and Spassky 1959), *D. willistoni*, *D. tropicalis* and *D. equinoxialis*, have been up to now cultivated in the same banana-agar medium utilized to grow and study other species of the genus *Drosophila*. In our laboratory as elsewhere little attention

has been paid (see recent issues of *Evolution and Genetics*) to culturing conditions. Most geneticists and zoologists in general think that to speak of the adaptive optimum or more specifically of Darwinian fitness (any component can be drastically affected by different culturing conditions as one can see in what follows) do not implicate the judicious testing of those laboratory conditions that will show the most abundant egg laying and larval survival the species or the simispecies have. Thus testing how natural selection changes the paths of gene frequencies under environmentally poor conditions brings forth very biased results and conclusions, in some cases producing even contradictory data of what the gene pool can do to approximate the average phenotype to a particular ecological exigency. Very often while studying the environmental variance in *D. willistoni* (experiments on the genetic load) we were puzzled by the apparent impossibility of reducing it under our laboratory conditions. After testing various ways of doing our crosses, changing incubating practices, rotating our culture media so that each tray could have the same amount of exposure to each condition, and after following the usual statistical recommendations, the environmental variance was still relatively prominent. By simply changing our laboratory culturing medium from bananas to guayaba (*Psidium guajaba*) we were able to reduce the variance to approximately one fourth of what it was before, and as expected the statistical analysis became twice as effective as before.

The following table shows the rate of development in *D. paulistorum*, *D. willistoni*, *D. tropicalis* and *D. equinoxialis*, in both banana - agar and guayaba - agar media. The reader can see how the rate of development from egg laying to adult emergence as imagoes were reduced considerably in our new laboratory medium.

Table 1. Rate of development of different species of the willistoni group in different laboratory media. The numbers represent days of development from egg laying to adult emergence from pupae.

	Banana - agar		Guayaba - agar	
	Pupa	Adult	Pupa	Adult
<u><i>D. paulistorum</i></u>				
Andean	10	16	8	12
Caripe	12	18	8	12
Llanos 13A	10	18	8	12
Transitional	10	16	8	12
Amazonian	11	16	8	12
Central American	11	16	9	12
<u><i>D. willistoni</i></u>				
Yaguaracaca	10	17	7	12
Valparaiso	10	17	7	13
<u><i>D. Tropicalis</i></u>				
Valparaiso	10	17	8	13
<u><i>D. equinoxialis</i></u>				
Mitú 2A	10	18	8	12

Acknowledgment: We wish to thank the Colombian National Science Foundation (Colciencias) for their support.

Reference: Dobzhansky, Th. and B. Spassky, 1959, *Proc. Natl. Acad. Sc.* 45: 419-428.

McCrady, E. University of North Carolina at Greensboro, North Carolina. Wing disc tracheotomy prior to pupation in *D. virilis*.

In order to establish the relative roles of the tracheal supply and the hypodermal stalk in the development of the wing disc during metamorphosis, we have compared the results of three distinct types of operations with mock operated controls. Mature third instar larvae of *D.*

*virilis* were etherized and dissected with a sterile Aloe #115 ultra-micro dissecting hook. The tip of the hook was inserted through the dorso-lateral body wall in the meta-thoracic segment so that four distinct results were obtained:

- a) unilateral cutting of the ventral branch of the dorsal segmental trachea, the only tracheal supply to the disc,
- b) severance of the hypodermal stalk, including its internal tracheal branch,
- c) operations in which both "a" and "b" were accomplished on the same side in one animal,
- d) mock operations consisting of insertion of the dissecting needle without the cutting of either the tracheal trunk or the hypodermal stalk.

The preliminary results of these operations are summarized below:

Operation Category	Number	Survivors (%)	Normal Wings (%)	Abnormal Wings (%)
a	117	77(66)	66(86)	11(14)
b	108	37(34)	35(94)	2(06)
c	46	19(41)	12(63)	7(37)
mock	44	41(93)	40(98)	1(02)

A majority of the wing abnormalities found following category "a" and "b" operations consisted of wings which were not properly expanded following emergence. Five operated animals in category "b" which had normal wings lacked one or more macrochaetae, most often the anterior notopleural. The relatively large number of wing abnormalities observed after category "c" operations, in which the mortality was also the highest, consisted of wings in which a major failure of cell differentiation had occurred, similar to earlier results obtained in the transplantation of whole discs into mature larvae<sup>1,2</sup>. Since the majority of operated discs were able to develop normally, it appears likely that the previously observed failure in wing development in transplanted discs could be associated with the metamorphosis of such discs in the absence of the normal association of their thoracic areas with the other discs ordinarily contiguous with them. The nature of this association and its influence are being investigated further. (Supported by grant 153 from the North Carolina Board of Science and Industry).

References: 1) Glancy, E.A. and R.B. Howland; 1938, Bio. Bull. 75: 99-105; 2) McCrady, E., 1970, Amer. Zoologist 10: 320 (Abst.)

Tarantul, V.Z., V.T. Kakpakov and V.A. Gvozdev. Kurchatov Institute of Atomic Energy, Moscow, U.S.S.R. Protein, RNA and DNA synthesis in the established line of diploid cells of *Drosophila melanogaster* in vitro.

The synthesis and intracellular content of macromolecules were determined in the established diploid line of embryonic cells of *Drosophila melanogaster* (Genetika, Russ., 1969, 5, 12, 67; DIS 1970 45: 110). The quantity of DNA per diploid cell measured chemically by diphenylamine reaction according to Burton was  $1.9 \times 10^{-12}$ g.

The quantity of RNA and protein per cell was  $7-14 \times 10^{-12}$ g and  $5-10 \times 10^{-11}$ g correspondingly. Actinomycin D ( $3 \mu\text{g/ml}$ ) inhibited incorporation of  $\text{C}^{14}$ -uracil or  $\text{C}^{14}$ -uridine to 5-10% of the control. The protein synthesis measured by the  $\text{C}^{14}$ -lysine incorporation stops in 8-9 hours after the actinomycin addition. The half life of messenger RNA evaluated by the actinomycin induced inhibition of  $\text{C}^{14}$ -lysine incorporation is about 3-3.5 hours. The RNA synthesis is resistant to  $\alpha$ -amanitin ( $20 \mu\text{g/ml}$ ), although the same sample of the drug effectively inhibits the RNA synthesis in the isolated rat liver nuclei. The RNA synthesis in presence of rifampicin ( $100 \mu\text{g/ml}$ ) decreases by 30%. Puromycin ( $100 \mu\text{g/ml}$ ) inhibits  $\text{C}^{14}$ -lysine incorporation by 85%. After 6-12 hours of puromycin treatment the increase of the number of cells accompanied by the decrease of cell size was observed although the normal mitotic figures were absent. The presence of purumycin leads probably to the abnormal cell division or cell fragmentation. Hydroxyurea in concentrations of  $100 \mu\text{g/ml}$  and  $1 \text{ mg/ml}$  inhibits the  $\text{H}^3$ -thymidine incorporation to 15% and 3% of the control respectively.

Hoenigsberg, H.F. Universidad de los Andes, Bogotá, Colombia. New culturing conditions for several *Drosophila* species.

Cheap, safe and easily prepared culturing media is one of the necessary instruments to do field work (experimental stations, collecting work, field research etc.) of population genetics of *Drosophila* in the tropics. Often the usual

banana-agar medium appears contaminated with fungi which impedes the normal growth of first generation larvae. Furthermore, banana-agar, although a good laboratory medium, if periodically seeded with dry yeast, proves to have a hard surface for *Drosophila* eggs to survive casual mechanical contact when not seeded with yeast. The reason is that timely fermentation propitiates the structure of a soft surface.

The author has made and tested three new culture media where difficulties such as those mentioned above are obviated. Moreover, outside of ground Agar and Tegocept, easily found tropical fruits are the only important ingredients. Common fruits such as those prescribed here are necessary in order to avoid bringing from long distances baskets with bananas to prepare adequate culturing conditions for *Drosophila*. Bananas are not as easily found in the tropical forest as thought in various genetic laboratories up north.

Table 1 presents the kitchen formulae for those new culturing conditions where papaya (*Carica papaja*) piña (*Ananas sativus*) and guayaba (*Psidium guajaba*) are used instead of bananas. Table 2 presents the list of species tested for adequate growth in such new culturing conditions.

Table 1. Kitchen formulae for three new culturing conditions for several *Drosophila* species.

	PAPAYA - AGAR	GUAYABA - AGAR	PIÑA - AGAR
Water	2000 cc.	2000 cc.	2000 cc.
Agar	87.5 gm.	87.5 gm.	87.5 gm.
Tegocept	40 cc.	40 cc.	40 cc.
	3 medium sized papayas liquified with 100 cc of water.	30 large sized quayabas liquefied in 250 cc. of water.	2 small sized piñas liquefied in 45 cc. of water.

Table 2. A list of species which grow well in the above mentioned agar media.

1. Saltans Group

*D. saltans*  
*D. prosaltans*  
*D. sturlevanti*  
*D. cordata*  
*D. neosaltans*  
*D. pseudosaltans*  
*D. parasaltans*  
*D. subsaltans*  
*D. neocordata*

2. Willistoni Group

*D. willistoni*  
*D. tropicalis*  
*D. insularis*  
*D. paulistorum*  
*D. equinoxialis*

3. Melanogaster Group

*D. melanogaster*  
*D. simulans*

Acknowledgment: We wish to thank the Colombian National Science Foundation (Colciencias) for their support.

Oshima, C. and T.K. Watanabe. National Institute of Genetics, Misima, Japan. Sterility genes in natural summer and autumn populations of *D. melanogaster*.

Frequencies of sterile male and female flies among several hundred flies collected simultaneously from the summer and autumn populations of 1970 in Katsunuma locality were found to be 4.3, 4.8 per cent and 3.4, 7.8 per cent re-

spectively. Among 342 second chromosomes extracted from each male fly in the summer and autumn populations, 69 chromosomes were found to be sterility chromosomes. The results are shown in Table 1.

Table 1. Frequency of sterility chromosomes and frequencies of male, female and both sexual sterility chromosomes

Collection time: 1970 (July, October)				
Class of viability	Semilethal	Subvital	Normal	Total
No. of tested chromosomes	46	41	255	342
No. of sterility chromosomes	13	12	44	69
Frequency (%)	(28.3)	(29.3)	(17.3)	(20.2)
No. of male sterility chromosomes (%)		32	(46.4)	
No. of female sterility chromosomes (%)		27	(39.1)	
No. of both sexual sterility chromosomes (%)		10	(14.5)	

The frequency of sterility chromosomes was higher than 12.6% in 1968.

By half diallel crosses between sterility lines, which have been maintained by the Cy-sterility balanced system, the frequency of allelism was determined in the summer and autumn populations as Table 2 shows.

Table 2. Results of allelism between sterility genes

Collection time		July 31		October 12	
Sex		♀	♂	♀	♂
No. of sterility chromosomes		11	19	24	23
No. of crosses		55	171	276	253
No. of allelic crosses		10	1	14	26
Frequency of allelism (%)		(18.2)	(0.6)	(5.1)	(10.3)
Frequency of finding		No. of sterility genes			
1		6	17	14	10
2		-	1	1	-
3		-	-	1	2**
5		1	-	1*	-
7		-	-	-	1

\*\*, \* persistent gene

Table 3. Persistent and allelic sterility genes during two years in Katsunuma locality

Collection date	Oct. 2, 1968	July 31, 1970	Oct. 12, 1970
Female sterility gene	FS 801 A - F (6 chromosomes)	FS 102 (1 chromosome)	FS 201 A - E *
Male sterility gene	MS 801 A - M (13 chromosomes)	MS 113 (1 chromosome)	MS 202 A ** MS 202 B • FS 206 MS 202 C • FS 217 (3 chromosomes)
		MS 114	MSS 203 A
		MS 103	MSS 203 B
		MS 102	MSS 203 C

allellic relationship

\*\*, \* persistent gene



The sterility strains, which were extracted from a natural population in Katsunuma locality in 1968 and have been maintained in our laboratory, were crossed diallelly with new sterility strains extracted from the summer population of 1970. On the other hand, diallel crosses between the sterility strains of the summer and autumn populations were performed.

One male sterility gene (MS 801) and one female sterility gene (FS 801) have persisted for two years. The male sterility genes (MS 202, B, C) in the autumn population of 1970 were linked with different female sterility genes (FS 206, FS 217). Three different male sterility genes in the summer population of 1970 were found to be combined in double sterility chromosomes (MSS 203 A - C). On the other hand, a female sterility gene (FS 102) has increased in the autumn population. The results are shown in Table 3.

The breeding pattern of *D. melanogaster* in Katsunuma locality has been scarcely known, but the places of hibernation and the dispersion of flies were probably localized in the district.

Ehrie, M.G. and R.C. King. Northwestern University, Evanston, Illinois. The anatomy of the larval ring gland of *Drosophila melanogaster* and its associated organs.

During histological processing *Drosophila* larvae are generally punctured to insure penetration of the fixative. The drop in hydrostatic pressure caused by puncturing often alters the three dimensional interrelations of adjacent organs. In 1966 F.G. Gottlieb published the details of a procedure that avoids

puncturing (J.Roy.Mic.Soc. 85: 369-373). We have adapted this technique for plastic embeddings. Larvae were relaxed, fixed, and dehydrated following Gottlieb's directions. The Dioxane used as the dehydrating solvent was replaced by propylene oxide, and the larvae were then infiltrated with the following resin mixture: Araldite 502 monomer, 8 ml; dodecenyl succinic anhydride hardener, 8.5 ml; and DMP-30 accelerator, 0.32 ml. The monomer and hardener must be mixed thoroughly before the accelerator is added. All components were obtained from Polysciences, Inc., Paul Valley Industrial Park, Warrington, Pa. 18976. Larvae were infiltrated with resin using the following schedule: 12 hours each in (1) 3 parts propylene oxide (P): 1 part resin mixture (R), (2) 1 P:1 R, (3) 1 P:3 R, and (4) 100% R. The specimens were transferred to embedding capsules containing fresh resin and were left to polymerize at 60°C, for 24 hours.

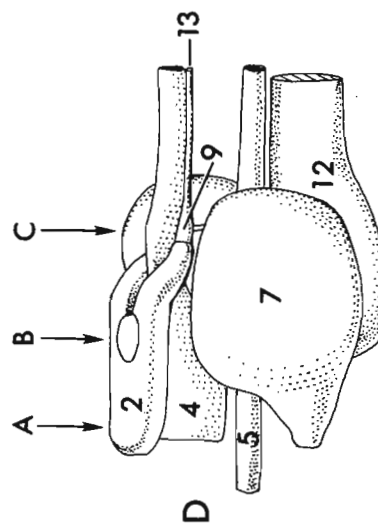
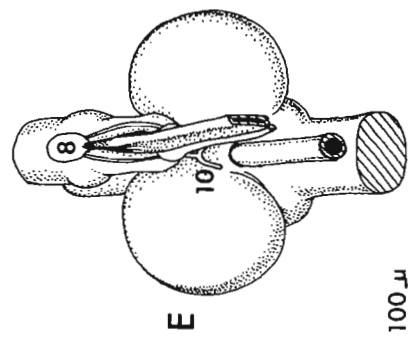
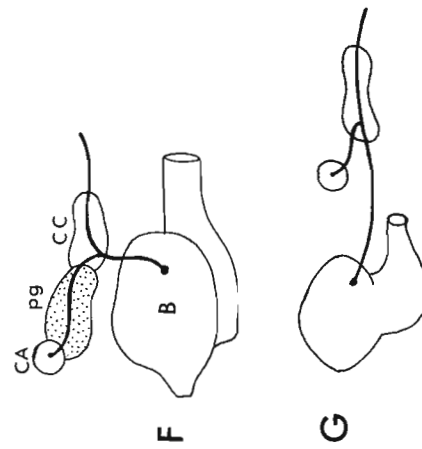
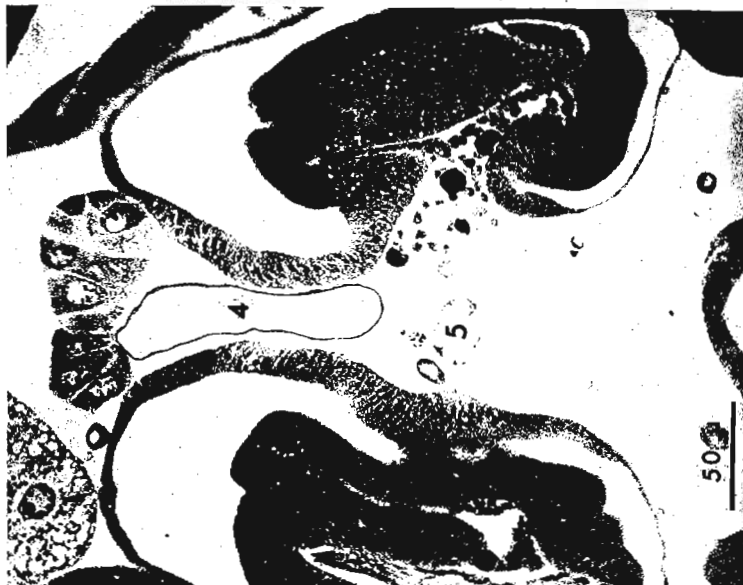
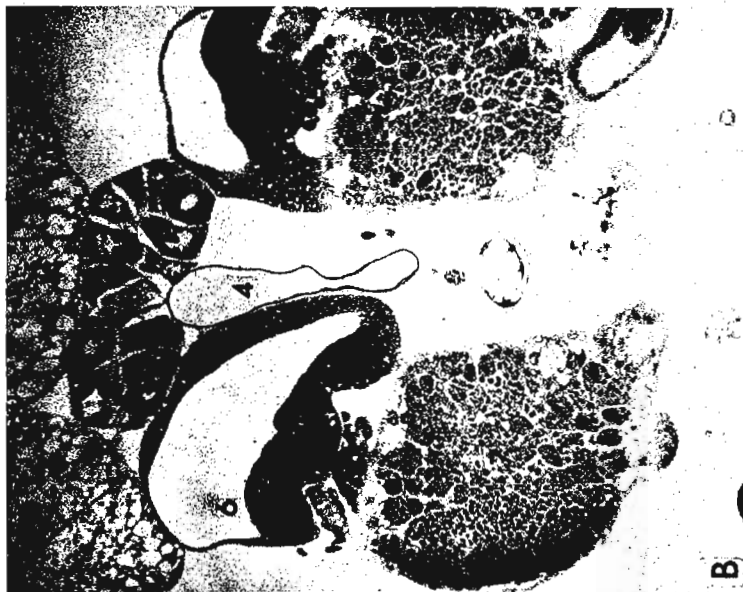
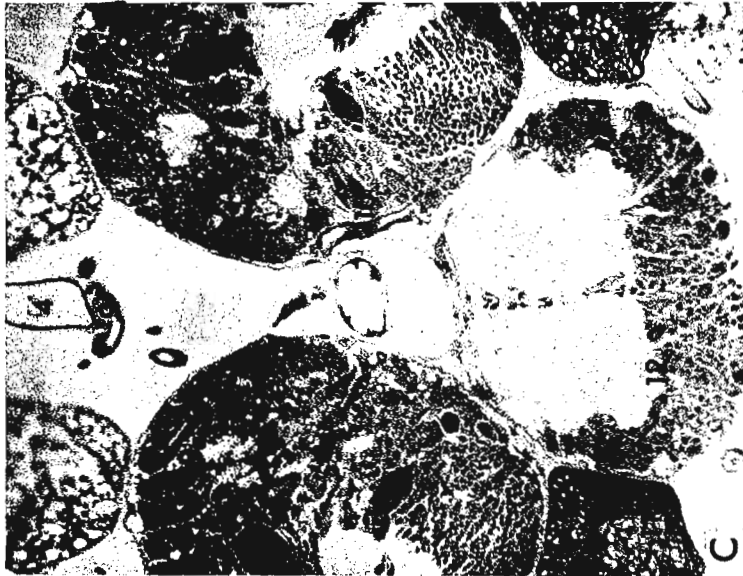
The female sectioned was in the terminal portion of the third instar (92 hrs. after hatching). One micron sections were cut with glass knives mounted in a Leitz Fernandez-Moran microtome. The serial transverse sections were stained with Azure B. Outline drawings of the sectioned brain, ring gland, aorta, and oesophagus were made on sheets of cardboard using a Wild M20 microscope equipped with a drawing tube. The tracings were cut out and glued together to form a three dimensional model of these organs at 350X.

The accompanying photomicrographs and drawings illustrate the results. The following labeling system is used: 1. fat body cell; 2. prothoracic gland cell; 3. tracheal cell; 4. aorta; 5. oesophagus; 6. antennal imaginal disc; 7. brain hemisphere; 8. corpus allatum cell; 9. corpus cardiacum; 10. afferent nerve to corpus cardiacum; 11. salivary gland cell; 12. ventral ganglion; 13. efferent nerve from corpus cardiacum. (See next page.)

The ring gland lies above the brain, straddling the aorta (Figs. D and E). Fig. A shows a section through the anterior portion of the ring which contains prothoracic gland cells clustered in paired longitudinal strips along the dorsal surface of the aorta. Two tracheae enter the prothoracic gland to the left and right where its ventral lateral surfaces rest upon the paired antennal discs (Figs. A and B). The anterior tips of the brain hemispheres and the corpus allatum are included in the section shown in Fig. B. The ventral ganglion and the corpus cardiacum are included in section shown in Fig. C. An afferent nerve to the corpus cardiacum is sectioned where it leaves the right brain hemisphere. It is joined by a nerve from the left hemisphere (Fig. E). A left and right nerve leave the corpus cardiacum and pass through the arms of prothoracic gland to the corpus allatum.

The positions of the brain (B), corpus allatum (CA), and corpus cardiacum (CC) are contrasted for the larva and adult in diagrams F and G. During metamorphosis the prothoracic gland (pg) degenerates and the corpus allatum and corpus cardiacum move posteriorly and ventrally relative to the brain.

(M.G. Ehrie held an Undergraduate Research Participation Award from the N.S.F. during the summers of 1968 and 1969.)



Baird, M.B., H.V. Samis and H.R. Massie.  
Masonic Medical Research Laboratory, Utica,  
New York. Changes in *Drosophila* catalase  
activity associated with preadult develop-  
ment.

Catalase (EC 1.11.1.6) activity was deter-  
mined in preadult *D. melanogaster* at vari-  
ous time intervals after oviposition at  
25°C. Ore-R females were permitted to  
lay eggs on enriched yeast plates for  
four-hour intervals, following which the  
eggs were transferred to standard corn

meal-agar-molasses medium (1). This technique results in cultures of developing flies which  
are temporally synchronous within  $\pm 2$  hours.

Samples of eggs were collected directly from the yeast plates at the indicated times  
after oviposition and washed thoroughly with insect saline to remove yeast and media contam-  
inants. Samples of larvae were collected by flotation in 1M NaCl, followed by thorough  
rinsing with insect saline. Samples of pupae were manually collected from the sides of the  
culture bottles.

All samples were homogenized in water, and dechitinized by a filtration technique des-  
cribed elsewhere (2). Catalase assays were performed at 22.5°C by modification of the  
spectrophotometric technique of Price et al. (3), utilizing a Perkin-Elmer 139 spectrophoto-  
meter equipped with an externally thermostated constant temperature cuvette chamber. 0.050ml  
of sample was added to a cuvette containing 3.0ml of 0.02M phosphate buffer, pH 6.8. The  
cuvette was blanked to zero absorbance, and 0.030ml of 0.98M hydrogen peroxide was added to  
the cuvette. Helium gas was immediately bubbled through the contents of the cuvette for  
five seconds, and the disappearance of hydrogen peroxide was recorded at 230 m $\mu$  for an  
additional 30 seconds with a chart recorder. Units of catalase were calculated as described  
by Lück (4), and protein was determined by the method of Lowry et al. (5).

No appreciable catalase activity was found in *Drosophila* embryos [Fig. 1]. However,

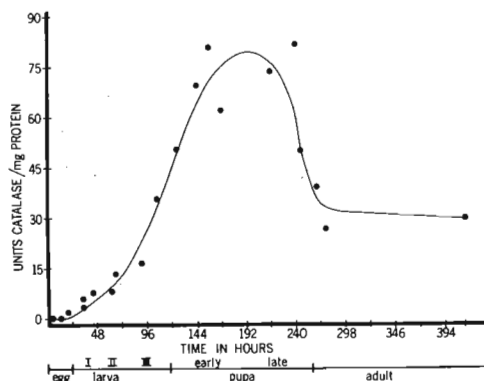


Figure 1. Catalase activity in preadult *D. melanogaster* at various times after oviposition at 25°C. Data expressed as units/mg. protein, where one unit is that amount of enzyme necessary to liberate half the peroxide oxygen from a hydrogen peroxide solution of given concentration in 100 seconds at 22.5°C.

appreciable enzyme activity appeared during the early larval stages of development, in-  
creasing 70-fold to maximal activity during mid-pupal development. This increase was fol-  
lowed by a sharp decline in enzyme activity prior to eclosion.

These preliminary results indicate a stage specific activation (e.g. differential gene  
action) of those genes in *D. melanogaster* which code for the normally ubiquitous catalase.

References: 1. Ursprung, H. 1967. In *Methods in Developmental Biology* (F.H. Wilt  
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F.C. Erk. 1969. DIS 44:132; 3. Price, V.E., Sterling, W.R., Tarantola, V.A., Hartley,  
R.W., Jr., and Rechcigl, M., Jr. 1962. J. Biol. Chem. 237:3468; 4. Lück, H. 1965.  
In *Methods of Enzymatic Analysis* (H. Bergemeyer, ed), Academic Press, New York, p885;  
5. Lowry, O.H., Rosebrough, N.J., Farr, A.L., and Randal, R.J. 1951. J. Biol. Chem. 193:  
265.

Minamori, S. and K. Ito. Hiroshima Univer-  
sity, Hiroshima, Japan. Effects of delta on  
fertility in *D. melanogaster*.

The ID<sup>b</sup>-45 chromosome line usually car-  
ries an appreciable amount of delta b,  
but it is not susceptible to the killing  
action of this delta (Minamori et. al.  
1970). The productivity of this line

was examined when it carried various amounts of delta b. More than one-third of Cy/ID<sup>b</sup>-45

males and females tested became sterile when they carried cytoplasm of Cy/Pm stock which is considered to carry no delta b. The number of progeny was smaller when flies of this line were raised at 25° C than raised at 28° C at which temperature the multiplication of delta is accelerated. The progeny number was appreciably reduced when the flies were raised at 18° C at which temperature the multiplication of delta is suppressed. This line could not be maintained at that temperature, since both males and females became sterile (Table 1).

Table 1. The number of progeny (average) recovered from Cy/ID<sup>b</sup>-45 flies which were raised for successive generations at 18° C

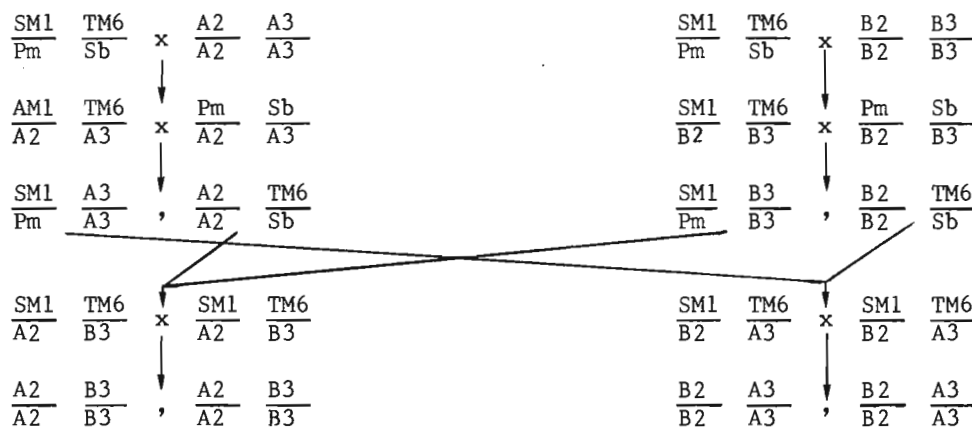
Subline	Raising temperature for progeny (C)	Generations raised at 18° C			
		1	2	3	4
0-9	18°	0	-	-	-
	25°	0	-	-	-
y-9	18°	5.9	15.0	3.4	0
	25°	33.5	0	-	-

Thus, the conclusion drawn may be that the presence of an appreciable amount of delta b is necessary for the gametogenesis of the Cy/ID<sup>b</sup>-45 flies.

Reference: Minamori, S., Fujioka, N., Ito, K., and Ikebuchi, M. 1970. *Evolution* 24: 735-744.

Moree, Ray. Washington State University, Pullman, Washington. A method for the construction chromosomal interchange lines.

The following scheme has been found useful for the construction of chromosomal interchange lines used in heterozygosity studies, where only the 2nd and 3rd chromosomes are interchanged.



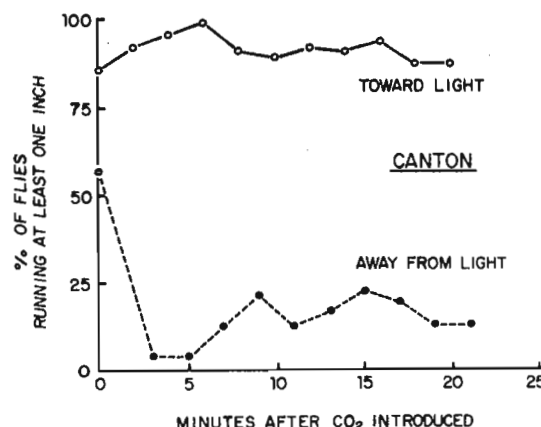
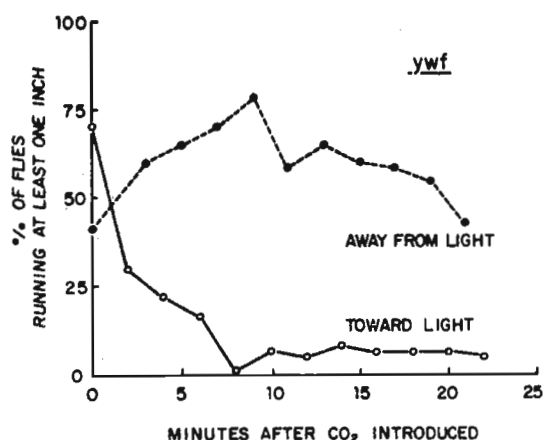
A and B designate different wild type stocks; 2 and 3 designate chromosomes 2 and 3. All other chromosomes are those described in Lindsley and Grell (Carnegie Institution of Washington Publication No. 677, 1968) except that TM6, obtained from E.B. Lewis, has a new marker, Ubx<sup>P15</sup>. Males used in the fourth cross can of course carry Pm instead of SM1 and Sb instead of TM6, which sometimes makes this cross easier to set up. The X chromosomes consist of material from the double balancer line, from line A, and from line B in the approximate ratio of 4:1:1, respectively. If lines A and B are made isogenic prior to making the interchanges, then maximum heterozygosity contrasts are possible. (Aided by funds from the State of Washington Initiative Measure No. 171 for the Support of Biological and Medical Research.)

Kaplan, W.D., and B. Hanstein. City of Hope National Medical Center, Duarte, California. A mutant stock showing negative phototaxis in the presence of  $\text{CO}_2$ .

Flies of a ywf stock carried in this laboratory for the past several years show a normal response to light. Placed in a 5-7/8 inch test tube containing air, and given one minute to run horizontally toward a fluorescent light, 70% move at

least one inch from the bottom of the tube toward the source of the light in an otherwise darkened room. 85% of the flies of a Canton-S stock exhibited this positive phototactic response.

When placed in an atmosphere of 20% carbon dioxide, however, the ywf stock responds by moving away from the light source. Placed at the end of the tube closest to the light, they begin to move away from the light. At nine minutes after the introduction of  $\text{CO}_2$ , 78% move at least one inch away from the light in a minute. At 8 minutes less than 10% run one inch toward the light. This is in contrast to the Canton-S stock which in the presence of 20%  $\text{CO}_2$  continues to run toward the light and not run away from it. (see graphs). The zero



minute point on the graph is before  $\text{CO}_2$  was introduced and the flies are in air. The curves are for the same groups of flies run on alternate minutes toward and away from light, and given one minute to run. Total number of flies in the four ywf groups was 60, and in the four Canton-S groups, 61.

White-eyed flies of a bw st stock, and just w by itself did not show this reversal. The reason for this behavior is currently under investigation.

Supported by N.I.H. grant No. NS08014.

Lefevre, G., Jr. San Fernando Valley State College, Northridge, California. A cytological analysis of X-ray-induced recessive sex-linked lethals.

Salivary chromosome preparations of sex-linked lethals recovered following exposure of mature sperm from wild-type males to 2000r and 3000r doses of X rays were analyzed for the presence of detectable rearrangements in the euchro-

matic portion of the X chromosome (1A through 19F). Among 190 lethal X chromosomes recovered in the 2000r experiment, 70 (36.8%) carried rearrangements; among 125 lethals from the 3000r experiment, 55 (44.0%) were detectably abnormal. These values are not significantly different. It would appear that, insofar as mature sperm are concerned, the proportion of recovered X-ray-induced lethal effects that are associated with rearrangements is not dose dependent.

Ouweneel, W.J. Hubrecht Laboratory, Utrecht, Netherlands. Homoeotic mutants in *Drosophila*: interaction during development.

A given mutation not only affects a specific type of normal tissue, but also the same kind of tissue whenever this arises at an abnormal location in the fly under the influence of a homoeotic

mutation. For instance, *dachs* not only shortens the normal legs but also the legs formed instead of the arista in *aristapedia* (*ss<sup>a</sup>*);<sup>1</sup> multiple-wing-hairs not only effectuates a characteristic hair pattern on the normal wing but also on the wing outgrowths appearing in the eyes under the influence of *ophthalmoptera* (*opht*).<sup>2</sup> It turns out that also homoeotic mutations themselves affect allotypic tissues produced by other homoeotic mutations. Three classes of genetic combinations are suitable to demonstrate this type of interaction. First, *ss<sup>a</sup>* was found to change not only the normal arista, but also the arista produced from the proboscis by *proboscipedia* (*pb*) (at low temperature) into a tarsus, whereas *pb* has no influence on the antenna. This confirms earlier results of Vogt,<sup>3</sup> although I feel her explanation (sequential activity of *ss<sup>a</sup>* and *pb*) is irrelevant. Secondly, *bithorax* (*bx*) changes the haltere into a wing-like structure, while *tetraltera* (*tet*) strongly reduces the wing and produces a hypodermal (leg-like?) protrusion along with, or instead of the wing. In *bx tet* combinations *tet* seems to reduce not only the normal wings but also the wing-like structures produced by *bx*; however, no leg-like structures replacing the halteres were observed so far. The same class of interactions is exemplified by the effect of *Contrabithorax*, which heterotopically changes the posterior mesothorax produced by *bithoraxoid* from the first abdominal segment into a posterior metathorax.<sup>4</sup> Thirdly, in combinations of *tet* with either *eyeless-opht* or *loboid-opht* it was found that *tet* not only affects the normal wings but also changes the wing-like outgrowths from the eye area produced by *opht* into hypodermal bristle-bearing protrusions. Therefore, homoeotic mutations act not only in one specific imaginal disc, but also at any other place in the developing organism where the genome allotypically determines a tissue to follow a given developmental pathway with which the mutations concerned can interfere.

References: 1. Braun, W. 1940, *Genetics* 25: 143-149. 2. Ouweneel, W.J. 1970, *Genetica* 41: 1-20. 3. Vogt, M. 1946, *Z. Naturforsch.* 1: 469-475. 4. Lewis, E.B. 1963, *Am. Zoologist* 3: 33-56.

Krimbas, C.B. Agricultural College of Athens, Athens (Votanikos), Greece. Gene arrangement frequencies in Pindos population of *D. subobscura*.

A small sample of *D. subobscura* taken in the summer of 1968 in a Quercus forest near the village of Korydallos, 34 km from Kalambaka, on the road of Kalambaka-Metsovon, in Pindos Mt. was analysed for the gene arrangement frequencies

in all five chromosomes, by crossing wild males and sons of wild females to a stand-

Table I

A <sub>st</sub>	A <sub>I</sub>	A <sub>2</sub>	N					
.33	.38	.29	58					
J <sub>I</sub>	J <sub>3+4</sub>	J <sub>St</sub>	N					
.74	.01	.25	85					
E <sub>1+2+9</sub>	E <sub>st</sub>	E <sub>8</sub>	E <sub>1+2</sub>	E <sub>1+2+9+12</sub>	N			
.58	.19	.19	.03	.01	86			
U <sub>1+2+6</sub> *	U <sub>1+2</sub>	U <sub>st</sub>	U <sub>1+2+8</sub>	U <sub>1+2+7</sub>	N			
.54	.29	.13	.02	.02	84			
O <sub>3+4</sub>	O <sub>3+4+1</sub>	O <sub>st</sub>	O <sub>3+4+22</sub>	O <sub>3+4+7</sub>	O <sub>3+4+2</sub>	O <sub>3+4+2</sub>	N	
.48	.21	.12	.06	.02	.06	.05	87	

\*U<sub>1+2+4</sub> included

ard strain. Table I reports these frequencies as well as the numbers of chromosomes studied. This is the first sample reported from the West of North Greece and does not differ strikingly from the mainland (North and South) Greek samples. It seems that local differentiation between not very far remote populations of *D. subobscura* is not great.

Khishin, A.F. and M.M. Megaheid. University of Assuit, U.A.R. Storage of germ cells and process of mutation in *Drosophila melanogaster*.

This research is designed to study the effects of storing *Drosophila melanogaster* male germ cells on the spontaneous and induced sex-linked and second chromosome lethal mutations. Three different storage periods of 3, 6 and 9 days were used. Adult males 3 days old were irradi-

ated with 2352 r of X-rays.

The Muller-5 (M-5) and the Curly Lobe (Cy/L) methods were used for the determination of the mutation rates for sex-linked and second chromosome lethals respectively. The result obtained suggests that:

The frequency of the spontaneous recessive sex-linked lethal mutations are not different statistically for different storage periods, or when compared with the unstored.

X-ray induced sex-linked lethal mutations may increase slightly after storing male germ cells in untreated females for 3, 6 and 9 days; the difference, however, is not statistically significant.

The percentage of spontaneous second chromosome lethals increases by storage. The difference is significant when storage continues for 6 and 9 days.

Storage of irradiated sperm for different periods increases the rates of second chromosome lethals over the rates obtained from the unstored irradiated sperm.

The effect of storage is more pronounced in the case of irradiated than in the case of untreated sperm.

The present study shows that the ratio between the induced sex-linked and second chromosome lethals increases by sperm storage.

Rosenfeld, A., A. Carpenter, and L. Sandler. University of Washington, Seattle, Washington. A nonchromosomal factor causing factor causing female sterility in *D. melanogaster*.

A homozygous *pr cn* stock in our laboratory appears to carry a nonchromosomal factor which will sterilize females that carry specific chromosomes contributed by the male parent. This system has features similar to the case of CO<sub>2</sub> sen-

sitivity studied by L'Héritier and his collaborators and to the delta-factor-induced lethality studied by Minamori, and is perhaps the same as the "maternally inherited factor" reported by Picard and L'Héritier (DIS 46:54, 1971).

The standard experiment, here, is to cross the two stocks to be examined such that each serves as female parent. *F*<sub>1</sub> females are tested for fertility by crossing to Canton-S ♂♂. *F*<sub>1</sub> daughters of *pr cn* mothers are fertile when the male is *pr cn*, Canton-S, or Muller-5; sterile when the male is *y*<sup>+</sup>; *abo/Cy* or Muller-5<sub>A</sub> (=Muller-5/*y*<sup>+</sup>; *+/+*; *+/+*; *spa*<sup>pol</sup>/*spa*<sup>pol</sup> ♂♂ from a stock kept as Muller-5/Muller-5; *Sml*/*+*; *+/+*; *spa*<sup>pol</sup>/*spa*<sup>pol</sup> x Muller-5/*y*<sup>+</sup>; *Sml*/*+*; *Ly Pr*/*+*; *spa*<sup>pol</sup>/*spa*<sup>pol</sup>) and *Cy* daughters are fertile, but *Cy*<sup>+</sup> daughters are sterile, when the male parent is *y*; *abo/Cy*. All other pairwise crosses gave fertile *F*<sub>1</sub> females (Muller-5<sub>A</sub> was not tested with *y*<sup>+</sup>; *abo/Cy*, *y*; *abo/Cy*, or Muller-5).

These data indicate: (1) that both parents much contribute something to the female-sterility phenotype; (2) that chromosome 2 may be of especial importance (from the results of *pr cn* ♀♀ x *y*; *abo/Cy* ♂♂), and (3) that *abo* (Sandler, Genetics 64:481-493, 1970) is not specifically involved (from the results of *pr cn* ♀♀ x Muller-5<sub>A</sub> ♂♂).

To examine the nature of the maternal contribution from the *pr cn* ♀♀, *F*<sub>1</sub> ♀♀ from the crosses: (A) *pr cn* ♀♀ x Muller-5 ♂♂ and (B) *pr cn* ♀♀ x Canton-S ♂♂ were crossed to *y*<sup>+</sup>; *abo/Cy* ♂♂ and *F*<sub>2</sub> ♀♀ tested for fertility (by mating with Canton-S males). In cross A, 52 B; *Cy*<sup>+</sup> and 50 B<sup>+</sup>; *Cy*<sup>+</sup> *F*<sub>2</sub> females were tested and all were sterile; in cross B, 59 *Cy*<sup>+</sup> *F*<sub>2</sub> females were tested and all were sterile. The *pr cn* maternal contribution, therefore, appears to be non-chromosomal since 1/16 of the sterile females should have received no chromosomes from the *pr cn* stock (except for the B; *Cy*<sup>+</sup> ♀♀ from cross B, where 1/8 would lack such chromosomes).

Further evidence on the nature of the non-chromosomal element in the *pr cn* stock comes from the results of the following experiment. *F*<sub>1</sub> females from a cross of *pr cn* ♂♂ by Canton-S ♀♀ were backcrossed to *pr cn* males. From this cross, 173 *F*<sub>2</sub> females were crossed to *y*<sup>+</sup>; *abo/Cy* males and the *F*<sub>3</sub> *Cy*<sup>+</sup> female progeny tested for fertility. In these crosses, *pr cn* ♀♀ were not involved; nevertheless 21 females were sterile, 81 were semisterile (producing one or occasionally two larvae), and only 71 were normally fertile. These data strongly suggest some transmission of the nonchromosomal element through the sperm and the existence of quantitative effects; the parallel with the unstable state of sigma in the CO<sub>2</sub> sensitivity system is striking.

Band, H.T. Michigan State University, East Lansing, Michigan. Four decades of natural selection.

Darwinian natural selection typically implies directional selection. To date reported instances of genetic changes brought about by directional selection pressures exerted by climatic changes

have been rare. In biology, man is still considered to be the primary agent effecting environmental changes (Crow, 1971). As noted by Lamb (1966) between World War I and World War II it was widely believed that climate was static except on the geological time scale. This was the period of classical genetics and the development of classical mathematical population genetics, which was incorporated into population ecology.

The study of the South Amherst, Mass. *D. melanogaster* natural population now emerges as the study of a population in a slowly changing climate. Regular shifts in decade averages for daily temperature range in summers (Band, 1971) reflect a trend toward more days in the wider range categories.

	1930's	1940's	1950's	1960's
Narrow (2-20°F)	334	289	261	225
Intermediate (21-25°F)	243	254	248	234
Wide (26-44°F)	343	377	411	461

Periods of genetic changes reflecting changes in lethal and semilethal frequencies are: 1938-1946, 1947-1961, 1962-1966, post-1966. In the first, second chromosome lethal and semilethal frequencies fluctuated around 48.8%, in the second around 33-34%; the third was marked by increased developmental homeostasis and resistance of *le + sle* frequency to decline followed by plunge to 16-17%. Post-1966 *le + sle* frequency has been rising. The number of days in the different temperature range categories in the different periods in summers are given in Table 2.

Table 2. Mean number of days in different temperature range categories in the different periods of genetic changes in the Amherst *D. melanogaster* population.

	1930-1946	1947-1961	1962-1966	1966-1969
Narrow	32	26	19	25
Intermediate	25	25	21	24
Wide	35	41	52*	43

\* $P < 0.05$  that significantly more days are in the wide range category. This is the case for all 5 summer.

Heterozygotes containing drastics have been found to have higher viability in narrow temperature ranges (Band, 1963; Oshima, 1969), those free of drastics to have higher viability in wide range conditions (Band, 1963, 1969). The behavior of the population 1962-1966 has provided an example of genetic homeostasis and population adaptation to a more severe climate (Band, 1971). The environmental data thus give support to the hypothesis that selection can be disruptive within summers, directional over the longer term (Band, 1971). They also provide further evidence that such recessive deleterious variants may actually be adaptive in heterozygous condition, hence are maintained in the population in response to the dynamic environment.

References: Band, H.T. 1963. *Evolution* 17:307-319; Band, H.T. 1969. *Japan. J. Genet.* 44, Suppl. 1: 200-208; Band, H.T. 1971. *American Naturalist* (in press). Crow, J.F. 1971. *BioScience* 21:107. Lamb, H.H. 1966. *The Changing Climate* (Methuen and Co., Ltd, London). Oshima, C. 1969. *Japan. J. Genet.* 44, Suppl. 1:209-216.



Band, H.T. Michigan State University, East Lansing. On the negative relation between summer rainfall and average daily temperature range.

Genetics studies on recessive lethal and semilethal frequencies in the S. Amherst D. melanogaster population have usually been made in the Fall until the discovery of a breeding site enabled investigations throughout the breeding season, May-

October (Ives, 1970). A negative significant relationship between lethal and semilethal frequencies and average daily temperature range of the week prior to collection (Band and Ives, 1961) and a positive significant relation between these frequencies in Fall and total summer rainfall (Band and Ives, 1968) has been observed. Studies on the relation between climate and genetic changes further indicated changes in general level of precipitation and average daily temperature range had occurred several times in the past 40 years, May-October, although decade averages for summer temperature range increased regularly (Band, 1971). To explore the rainfall-average daily temperature range relation in summer, data back to 1889 have been analyzed.

Table 1 shows a negative relation between these two climatic variables. Rainfall declines spanning 4 or more summers are noted in the 1890's, 1910's, and 1960's. To a lesser extent rainfall in the 1930's was also reduced. Rowan (1954) reports that although the Bruckner cycle has a periodicity of approximately every 35 years, variations from 20 to 50 years are noted. Average maximum daily temperature can be judged to be about 1°F higher at the Amherst College Weather station (data past Nov. 1948) than at the old U. Mass. weather station site (1889- Nov. 1948) which was moved in 1960. Although low rainfall in summers was most prolonged in 1907-1913, Ives' father recalled that Spring rains and cooler summers mitigated against drought. The number of days with 90° maxima or above have increased since the 1930's.

Table 1. Summer rainfall, average daily temperature range and average daily maximum temperatures in the same season, Amherst, Mass., 1889-1970

Interval	no. years	Rainfall in inches	Daily T° range	Max T°
1889-1892	4	14.71	22.5	79.0
1893-1896	4	8.96	24.7	80.9
1897-1906	10	14.23	23.0	79.2
1907-1913	7	8.15	25.4	80.7
1914-1922	9	13.09	22.6	79.7
1923-1924	2	6.35	26.3	81.4
1925-1928	4	13.45	21.9	78.2
1929-1936	8	10.11	23.3	80.7
1937-1946	10	12.74	22.7	80.6
1947-1961	15	11.40	24.0	82.0
1962-1966	5	8.60	26.1	82.1
1967-1970	4	12.60	4.0	82.2

Table 2 shows decade averages for daily temperature range in summers since the 1890's. Singer (1970) has commented that nothing is known about climatic stability. Although an apparent stability of temperature range is noted in the first few decades of recorded observations in that area, the data in Table 1 indicate there have been fluctuations in temperature range in association with rainfall levels. We may only speculate that had the tools for Drosophila genetics research been available then, genetic changes in the population in relation to climatic shifts might have been observed in the population as have been witnessed since studies began on recessive lethal and semilethal frequencies in 1938.

Table 2. Average daily temperature range per summer in the past 8 decades in Amherst

1890's	1900's	1910's	1920's	1930's	1940's	1950's	1960's	1970's(?)
24.0	23.9	23.8	23.3	22.5	23.3	24.1	24.9	25.7(?)

References: Band, H.T. 1971. Amer. Nat. (in press); Band, H.T. and P.T. Ives. 1961 P.N.A.S., Wash. 47:180-185; Band, H.T. and P.T. Ives. 1968. Evolution 22:633-641; Ives, P.T. 1970. Evolution 24:507-527; Rowan, W. 1954. J. Wildl. Mgt. 18:52-60; Singer, E.F. 1970. Science 170:125 (editorial).

Band, H. T. Michigan State University, East Lansing, Michigan. Was there a drought in the northern U.S. in the 1960's?

Genetic changes in the South Amherst, Mass. D. melanogaster population in the 1960's appear to have been initiated after the onset of a severe decline in rainfall, July, 1961, which lasted to

June, 1966. In a study of meteorological drought in Michigan, Strommen, van den Brink and Kidder (1969) commented that drought had been an increasing problem in the Northeast in the 1960's. Their study over the past 4 decades indicated prolonged drought effects in both the 1930's and 1960's in many areas of the state. The 1930's drought appears more evident in Michigan data than in Amherst, Mass. data.

Kendeigh (1961) indicates a low rainfall cycle has been observed and effects on duck numbers noted. Bruckner (1890) detected a cycle approximately every 35 years in data back to the 1700's. Rowan (1954) correlated low duck numbers in the 1820's, and 1860's, and 1890's and 1930's with this cycle. Rowan had been at the University of Alberta, Edmonton. A study recently compiled by the U.S.D.I. (1971) indicates that drought was widespread in the prairie provinces of Manitoba, Saskatchewan and Alberta in Canada and upper Great Plains states of Montana, the Dakotas and western Minnesota in the late 1950's and early 1960's, and again in the summer and fall of 1967. About 50-75% of the important game ducks come from this region. Duck numbers, 1955-1970, appear to follow the effects of the drought on numbers of suitable prairie pothole habitats. The following table has been compiled from the survey of breeding size of populations reported annually for the past 15 years and included in the report.

Mean breeding size of duck populations for the past 15 years. Numbers in millions.

1955-1960	1961-1965	1966-1968	1969-1970
42462.3	31254.6	34605.3	42355.0

Duck breeds surveyed include mallards, gadwall, American widgeon, green winged teal, blue winged teal, shoveler, pintail, redhead, canvasback and scaup.

The continuing relationship between drought cycle and duck numbers in that area indicates that population numbers do not fluctuate at random, somewhat at variance with the conclusions of Cole

(1954). It also raises the question, was there a widespread reduction in rainfall throughout the northern United States in the 1960's?

References: Bruckner, E. 1890. Klimaschwankungen seit 1700. Vienna; Cole, L.C. 1954. J. Wildl. Mgt. 18:2-24; Kendeigh, S.C. 1961. Animal Ecology (Prentice-Hall, Inc., Englewood Cliffs, N.J.); Rowan, Wm. 1954. J. Wildl. Mgt. 18:52-60; U.S. Department of the Interior, Bureau of Sport Fisheries and Wildlife. Migratory Game Bird Briefing Book. Prepared by the Division of Management and Enforcement. Jan. 1971.

Gabay, S.J. University of Illinois, Urbana, Illinois. Recombination at the bar locus in a reverse attached-X system in *D. melanogaster*.

Recombination at the bar locus in *Drosophila melanogaster* was studied in a reverse attached-X stock synthesized by Dr. E. Novitski. The males of this stock were of the constitution  $XY^{S.Y^L}$ . The females of the stock carried a compound reversed metacentric X

chromosome. They were homozygous for y and B, and heterozygous for v, f and  $os^0$ . The X chromosomes were attached at the "yellow" end rather than at the usual centromere end. This arrangement facilitates a higher frequency of homozygosis for the bar region and thus permits analysis of the marker constitution of exceptional females, that is, females which are double-bar or half-bar in phenotype. The likelihood of recovery of sister chromatids involved in exceptional events is increased as well, a prerequisite to the study of the reciprocity of intrachromosomal events.-----Exceptional females were analyzed to determine the bar genotype and the combination of marker genes carried by each of their X chromosomes. The results were interpreted in regard to exchange between obliquely synapsed members of the duplication, which is associated with exchange of outside markers and is presumed to be reciprocal. Results were also interpreted in regard to the hypothesis of intrachromosomal exchange (Laughnan, 1961) in which the markers are expected to be nonrecombinant and which proposes a reciprocal sister chromatid event.-----A total of 59,836 females, and therefore 119,672 X chromosomes, were examined for changes at the bar locus. A total of 193 recombinant exceptions was observed. Fifty-seven of these carried bar changes in both X chromosomes and were  $B^+/BB$  in genotype with the exchange of outside markers which is predicted on the basis of reciprocal products of crossing over between obliquely synapsed members of the bar duplication. The reciprocity of the exchange between obliquely synapsed duplication members was thus demonstrated. One hundred of the recombinant exceptions were of the genotype  $B^+/B$  and thirty-six were  $BB/B$ . The three classes of recombinant exceptions are expected to be equal in frequency and the discrepancy is ascribed to the reduced viability of the  $BB$  females.-----Eleven females carrying nonrecombinant exceptional strands were observed. Eight of these were half-bar in phenotype, one had round eyes and two were double-bar in phenotype. One of the  $BB$  nonrecombinant strands was recovered along with its sister strand and offered an opportunity to test the prediction that intrachromosomal exchanges between sister strands produce  $BB$  and  $B^+$  strands as reciprocal types. Analysis showed this exceptional female to carry bar on one strand and double-bar on the other. While females of this type are not expected on the hypothesis of intrachromosomal exchange, the number of  $BB$  nonrecombinant females is insufficient to refute the hypothesis. More cases in which the nonrecombinant  $BB$  exceptional strand is recovered along with its sister are needed before any final conclusions concerning the model can be drawn.-----Four females carrying aberrations associated with bar changes were also observed. Three of these carried deletions and the fourth is still in the process of analysis.-----All 193 recombinant-exceptional females obtained from these studies exhibited normal genetic behavior in that all exceptional strands were homozygous viable. The frequency of these recombinant bar changes was one per 480 X chromosomes. The eleven females carrying nonrecombinant exceptional strands were analyzed cytologically as well as genetically and the frequency of nonaberrant changes was one per 10,000 X chromosomes. The frequency of X chromosomes carrying bar changes associated with aberrations was one per 24,000 X chromosomes.

This work submitted in partial fulfillment of the requirements for the degree of Doctor of Philosophy in Botany in the Graduate College of the University of Illinois, 1969.

This work was supported by National Science Foundation Grant GB-7635.

Reference: Laughnan, J.R., 1961, Mutation and Plant Breeding, NAS-NRC 891: 3-29.

Würgler, F.E. Swiss Federal Institute of Technology, Zürich, Switzerland. Synthetic female sterile factors in two combinations of X-chromosomes with  $dp\ bw$ ;  $st\ p^D$  autosomes in *D. melanogaster*.

During the last few years we tried to construct some new multipurpose stocks with marked sex-chromosomes and the autosomal markers  $dp\ bw$ ;  $st\ p^D$ . The autosomes were always derived from the Inscy;  $dp\ bw$ ;  $st\ p^D$  stock obtained from I.I. Oster, Bowling Green, Ohio, USA (stock number j

419 in the 1971 stock list). In a first set of experiments (U. PETERMANN, Mutation Res. 5, 397-410, 1968) we tried to replace the Inscy chromosome by a crossover product with the markers  $y\ sc^{S1}\ B\ f\ In-49\ v\ w^a\ sc^8$  ( $=X^*$ ) obtained from females heterozygous for the following

chromosomes:  $y^{sc^{S1}} B^{In-49} v^w sc^8 / sc^{S1} f^{In-49} v^w sc^8$ . It turned out that none of the crossover chromosomes obtained gave fertile females in combination with the  $dp\ bw; st\ p^P$  autosomes. A similar result was recently found in an attempt to replace the Inscy chromosome by the  $XY(Parker\ 110-8)\ y^2\ su(w^a)\ w^a\ KS.KL\ y^+$  chromosome. The following table summarises our results:

Inscy/Inscy ; $dp\ bw; st\ p^P$	fertile
$X^* / Inscy; dp\ bw; st\ p^P$	fertile
$X^* / X^*; dp\ bw; st\ p^P$	sterile
$XY\ (Parker\ 110-8)/Inscy; dp\ bw; st\ p^P$	fertile
$XY\ (Parker\ 110-8)/XY(Parker\ 110-8); dp\ bw; st\ p^P$	sterile

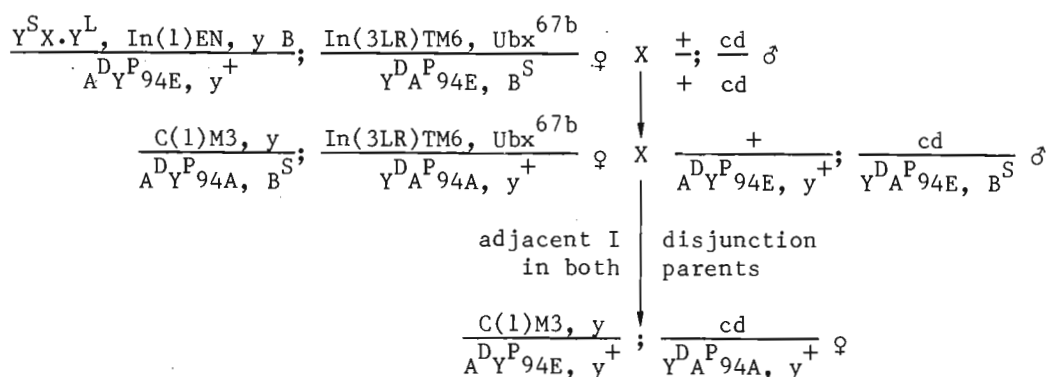
Since the  $X^*$  and the  $XY(Parker\ 110-8)$  in homozygous condition in combination with other autosomes (including heterozygosity for  $dp\ bw; st\ p^P$ ) give fertile females "synthetic sterility factors" seem to result in the particular combinations listed in the table. Male fertility was not affected.

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Jones, A.M. University of California, La Jolla, California. The cytological localization of  $cd$  and  $wo$  by means of deficiency mapping.

$X/T(Y;3)/cd$  males were produced using translocations B93, D100, B27, and H173 with autosomal breakpoints in 93F-94A, 94A, 94E, and 95E, respectively. These males were crossed to stock females carrying translocations with adjacent autosomal breakpoints. In this manner

it was possible to produce interstitial deficiencies for the segments between the autosomal breakpoints of the two translocations in combination with the normal third chromosome 3 carrying  $cd$ . These heterozygotes are recognizable on the basis of the phenotype with respect to  $y$ ,  $Hw$ ,  $B^S$ , and  $Ubx$  as outlined below:



The  $y^+y^+$  (extreme hairy wing) non  $B$  phenotype of this female shows unambiguously that she carries the  $A^{DYP94E}$  and the  $Y^{DAP95E}$  elements and is therefore deficient for 94A to 94E. The fact that she has cardinal eyes places the  $cd$  locus between 94A and 94E. Similar crosses placed  $wo$  (white ocelli) in the same cytological interval. The exact status of  $wo$  is uncertain, however, as both  $cd/cd$  and  $cd/wo$  flies have white ocelli. By the same procedures  $obt$  (obtuse) and  $bar-3$  were found to lie outside the 94A-95E interval.

The sterility of some  $X/T(Y;A)$  males impairs the general utility of the above method; this difficulty can be circumvented by first constructing males of constitution  $Y^{SX.Y^L}/X$ ; autosomal recessive/autosomal balancer for use in the first generation indicated above to produce  $Y^{SX.Y^L}/T(Y;A)$ /autosomal recessive in place of  $X/T(Y;A)$ /autosomal recessive in the second generation of the crossing scheme.

Wargent, J.M. University of Sheffield, Sheffield, England. Position-effect variegation in wings of *D. melanogaster*.

The wings of the mutant miniature of Krivshenko<sup>1</sup> ( $m^K$ ) at normal temperatures (25°C) are sometimes crumpled and may vary in size from fully wild type to fully 'miniature'. The 'miniature' wings show the distinctive morphology of the

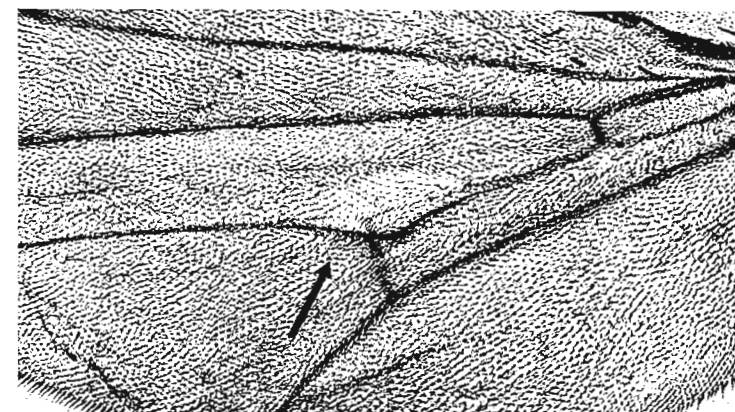
miniature ( $m$ ) mutant.

Reciprocal crosses,  $m^K/m^K \times m/Y$  and  $m/m \times m^K/Y$  were set up, and the  $F_1$  bred at 14°C. Microscopic examination of the wings of the female heterozygote,  $m^K/m$ , showed that they contain patches in which the hairs appeared closer together than in surrounding areas. Measurements were made of the distances between hairs in patches (p) and also in surrounding areas (s). For comparative purposes measurements were also made of the inter-hair distances in the Amherst wild type and in a homozygous  $m$  strain, both bred at 14°C. The results are shown in the table below.

Strain	Number of Measurements	Mean inter-hair distances $\pm$ standard error
$m^K/m$ (p)	50	62.19 $\pm$ 0.93
$m^K/m$ (s)	50	88.55 $\pm$ 0.98
+/+	50	92.08 $\pm$ 1.51
$m/m$	50	53.40 $\pm$ 0.90

wild type. The difference between the means for  $m^K/m$  (s) and  $m^K/m$  (p) is significant at the 1% level. These results indicate that the wings contain both wild type and  $m$ -like cells.

There is considerable variation in the size of the patches observed; some wings contain predominantly wild type cells while others consist mainly of  $m$ -like cells. The photograph



Statistical comparison of the difference between means shows that the wild type and the  $m$  mutant are significantly different at the 1% level. The difference between  $m^K/m$  (s) and +/+ is not significant and although a significant difference was obtained between  $m^K/m$  (p) and  $m/m$  the inter-hair distances of the former resemble those of the  $m$  strain rather than those of the wild type. The difference between the means for  $m^K/m$  (s) and  $m^K/m$  (p) is significant at the 1% level. These results indicate that the wings contain both wild type and  $m$ -like cells.

The photograph shows a predominantly 'miniature' wing with an area of wild type cells. It is assumed that the crumpled phenotype is caused by the presence of patches, and that the number and size of the patches determine the size of the wing.

The strain contains an inversion with breakpoints in section 10E4-5 and section 20B of the salivary X chromosome. The  $m$  locus, in section 10E1-2, is relocated next to broken heterochromatin in the rearranged chromosome. The presence of mutant cells in the wing is therefore probably due to a variegation-type position effect<sup>2</sup> at the  $m$  locus; this conclusion is supported by the observation that the mottling is enhanced by low temperature.

References: 1. Krivshenko, J., 1956, DIS 30: 75; 2. Lewis, E.B., 1950, Advances in Genetics 3: 73-115. This work was supported by Grant No. 68/1317 from the Science Research Council of Great Britain.

Novitski, E., E. Ehrlich and H. Becker\*. University of Oregon, Eugene, Oregon and University of Munich, Germany\*. A terminal attachment region on 2L.

Compounds involving the X and Y chromosomes or several X-chromosomes have been put together in virtually every combination, but compounds involving the autosomes have been limited to the five cases where homologous arms of an autosome have

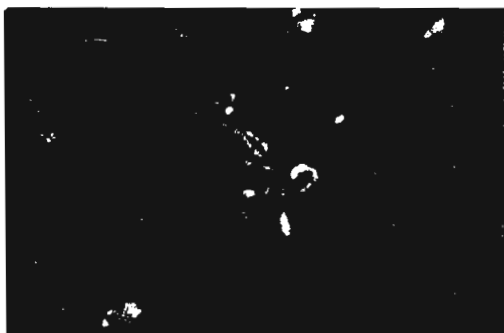
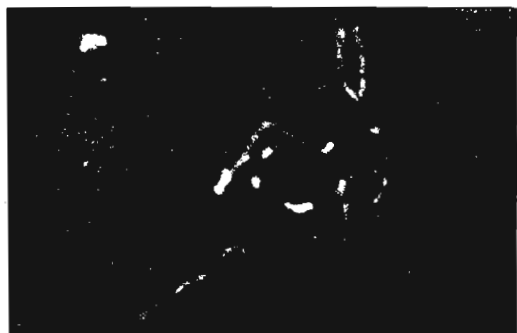
been attached to the same centromere as reversed metacentrics. There are two reasons for this difference in versatility of the sex chromosomes as opposed to the autosomes.

1. The sex chromosomes are much simpler to handle in single and compound forms because of the manner of their transmission and because of the prior existence of certain specialized chromosome types.

2. The occurrence of a complete X-chromosome with a terminal heterochromatic (and dispensable) section, first found in In(1)EN, made it possible to tack X-chromosomes together in serial order into compounds of all possible combinations. Since the construction of compound autosomes is limited by the absence at any of the ends of any useful terminal heterochromatic piece with an associated dispensable marker, it was decided to synthesize such chromosomes as a first step in making more useful autosomal compounds.

The hope that such an attempt might be successful stems from the suspicion that on one or more of the chromosome tips there may be small regions that are essentially heterochromatic (or a few loci which can be readily dispensed with in the heterozygote). This thought is based in part on the occurrence of the two ring chromosomes, R(1) and R(1)2, from attached X-chromosomes. The simplest and perhaps only explanation for their origin is that one of the tips of the attached X's underwent a rare "exchange" with the base of the other arm of the attached X. It seemed a good possibility that a similar situation might exist at the tips of one or more autosomes and that a search for a similar exchange that would add a larger piece of heterochromatin along with a good marker to an autosome would give positive results.

The procedure involved irradiating (3,000r) females carrying a doubly marked Y-chromosome along with an attached X, and looking for cases among the  $F_1$  where the two markers,  $y^+$  and Bar, have been separated, indicating that some kind of "exchange" had taken place, but where the attached X was not involved. Almost without exception, each female so treated produced one or more progeny showing such a separation and it was necessary to limit the analysis to one exceptional progeny per parental female, in order to avoid duplication. The exceptions selected were tested for segregation of the X-chromosome markers from the major autosomes. Two cases were found of attachment of Bar to the left end of the second chromosome. In both cases, results of the tests of this chromosome against a normal second showed for it to be .01 units to the left of the locus of  $al$ . Tests for the presence of Y-chromosome fertility factors at the left end of X have showed that in the first case (B3) none of the fertility factors is present and in the second case (B5) two (S1 and L5, and possibly L4) are. B3 is viable and fertile when homozygous; tests are not yet complete for B5.



Ganglion metaphases stained with quinacrine hydrochloride confirm the presence of some of the Y-chromosome at the tip of the second chromosome of B5. As can be seen on the photographs, there are two bright fourth chromosomes and XY chromosome with two bright regions at one end (undoubtedly corresponding to the short arm of the Y) and three at the other (the long arm of the Y). One of the autosomes has two bright blobs at one end, with a less well stained piece distal to them. This must represent the tip of 2L to which part of Y long, along with Bar, have been attached.

With these chromosomes, it should now be possible to make up certain additional compounds, by hooking another autosomal arm (or an X, or a Y) to the tip of 2L.

Additional runs are being made to try to synthesize chromosomes in which a marker is similarly placed on the tips of the other autosomal arms. If no other cases occur, an attempt will be made to use certain Y-autosome translocations (see the Washington-La Jolla report in this issue).

Bremner, T.A., W.L. Douglas and G.O. Ogonji.  
Howard University, Washington, D.C. Substrate-specific differences of alcohol and octanol dehydrogenases in eight species of *Drosophilidae*.

Of the ten cathodally migrating isozymes of alcohol dehydrogenase (ADH) detected by Ursprung and Leone (1965) in *D. melanogaster*, the slowest three showed stronger formazan staining with n-octanol than with ethanol. These three bands were shown, on the basis of linkage relation-

ships, substrate specificity, and differential elution from DEAE cellulose columns, to belong to a separate enzyme system, octanol dehydrogenase (ODH) which shows strong formazan staining with n-hexanol, n-heptanol, and n-octanol (Courtright, et al., 1966). Isopropanol and sec-butanol are equally good substrates or better than ethanol for ADH in *D. melanogaster* (Johnson and Denniston, 1964; Grell et al., 1965).

In an attempt to differentiate these two enzyme systems further and to ascertain whether there might be species-related differences in substrate specificity within each system, a comparison of the substrate requirements of both ADH and ODH in eight members of the family *Drosophilidae* was undertaken. The eight species belong to two genera, *Drosophila* and *Zaprionus*. Of the seven *Drosophila* species four, *robusta*, *camargoi*, *metzii*, and *unipunctata* belong to the subgenus *Drosophila*, while *D. lebanonensis casteeli* belongs to the subgenus *Pholadoris* (primitive), *D. busckii* to the subgenus *Dorsiphola*, and *D. melanogaster* to the subgenus *Sophophora*. The single *Zaprionus* species is *Z. multistriata*.

The substrates fall into four categories, primary unbranched alcohols, secondary alcohols, branched primary alcohols, and a cyclic alcohol, cyclohexanol. The method of agar gel electrophoresis and formazan staining of Ursprung and Leone (1965) as modified by Pipkin (1968) was used to assay crude homogenates of single female flies cultured on an enriched medium, and aged according to the following schema: *D. busckii*, 4-6 days; *D. melanogaster*, 5-6 days; *D. metzii*, 5-8 days; *D. unipunctata*, 7-9 days; *D. camargoi*, *D. robusta*, *D. l. casteeli*, and *Z. multistriata*, 9-11 days. These were the ages at which the respective species attained their optimum levels of enzyme activity as measured by the intensity of formazan staining. To compensate for the very small size of *D. busckii* 2-4 females were homogenized in a drop of distilled water.

From Tables I-IV it can be seen that both the ADH and ODH of *D. unipunctata* show more intense staining when secondary alcohols are used as substrates. The ADH of *D. busckii* is aberrant in that it shows a preference for the short chain unbranched primary alcohols while its ODH activity is quite low. Both ADH and ODH show a moderate preference for the long chain, unbranched primary alcohols in five of the species assayed. Although the two enzymes have overlapping substrate specificities, that of ODH is distinctly narrower and comprises

Table I Unbranched Primary Alcohols

Substrate	Enzyme Activity	mel.	metz.	busc.	rob.	uni.	car.	Dlc.	Z.mult.
Methanol	ODH	-	*	-	-	-	**	-	*
	ADH	***	-	***	***	**	***	***	***
Ethanol	ODH	-	-	-	-	-	*	-	*
	ADH	**	-	*	***	*	***	***	***
N-propanol	ODH	**	-	-	*	***	**	**	**
	ADH	***	-	***	***	*	***	***	***
N-butanol	ODH	*	*	-	**	**	**	**	*
	ADH	***	-	*	***	*	***	***	***
N-amyl	ODH	**	**	*	*	-	***	***	***
	ADH	***	-	**	***	*	***	***	**
N-hexanol	ODH	**	***	**	**	-		***	***
	ADH	***	-	**	**	*		***	**
N-heptanol	ODH	**	***	*	***	***	***	***	***
	ADH	***	-	*	***	*	***	***	***
N-octanol	ODH	***	***	**	**	-	***	***	***
	ADH	***	-	-	***	*	**	***	***
Nonyl alc.	ODH	*	*	*	**	*	**	***	***
	ADH	***	-	*	***	*	***	***	***
Decyl alc.	ODH	*	*	*	-	-	**	**	**
	ADH	***	-	*	***	*	***	***	**

Table II Secondary Alcohols

Substrate	Enzyme Activity	mel.	metz.	busc.	rob.	uni.	car.	Dlc.	Z.mult.
2-butanol	ODH	-	*	*	*	**	**	-	***
	ADH	***	-	*	**	**	***	***	***
2-hexanol	ODH	-	-	-	-	***	*	-	***
	ADH	***	*	**	***	*	***	***	***
4-heptanol	ODH	-	-	-	-	**	***	-	-
	ADH	***	-	*	***	**	***	*	*
2-octanol	ODH	-	-	*	***	***	***	***	***
	ADH	**	-	*	*	**	**	***	**

Table III Branched Primary Alcohols

Substrate	Enzyme Activity	mel.	metz.	busc.	rob.	uni.	car.	Dlc.	Z.mult.
Iso-propanol	ODH	-	*	-	*	-	-	-	-
	ADH	***	*	*	***	**	***	***	***
Iso-butanol	ODH	*	*	-	***	*	***	**	**
	ADH	***	*	*	**	***	*	***	***
Iso-amyl alcohol	ODH	*	**	*	*	**	*	**	**
	ADH	***	-	*	*	**	**	**	**
Tert-butanol	ODH	*	-	*	-	-	**	-	-
	ADH	**	*	-	***	*	**	***	**
Tert-amyl alcohol	ODH	*	-	-	*	-	-	*	*
	ADH	***	-	***	*	***	*	***	**

Table IV Cyclic Alcohol

Substrate	Enzyme Activity	mel.	metz.	busc.	rob.	uni.	car.	Dlc.	Z.mult.
Cyclohexanol	ODH	*	-	-	*	**	*	-	*
	ADH	***	-	**	***	*	***	*	*

Legend, Tables I - IV.

Species: mel. = *D. melanogaster*; metz. = *D. metzii*; busc. = *D. busckii*;  
 rob. = *D. robusta*; uni. = *D. unipunctata*; car. = *D. camargoi*;  
 Dlc. = *D. l. casteeli*; Z. mult. = *Z. multistriata*.

Intensity of formazan staining:

\*\*\* = strong; \*\* = moderate; \* = trace; - = negative

primarily the 5 to 8-carbon alcohols, with n-heptanol giving the highest intensity of staining. The ADH of *D. unipunctata* shows no staining with n-hexanol, but moderate staining with cyclohexanol; its ADH uses all the substrates tested. The ADH of *D. metzii* does not use any of the unbranched primary alcohols, while that of *D. melanogaster* uses all of them.

The above findings suggest that the same enzyme in different species shows small but significant differences in substrate specificity which may be related to minor evolutionary differences in the structure of the molecule resulting in different stereochemical requirements for enzyme activity.

This work was supported by National Science Foundation Grant GB 8779, and NIH Grant 1 R 101 GM 18409-01.

References: Courtwright et al, 1966, *Genetics* 54:1251-1260; Grell et al, 1965, *Science* 149:80-82; Johnson and Denniston, 1964, *Nature* 204:906-907; Pipkin, 1968, *Genetics* 60: 81-82; Ursprung and Leone, 1965, *J. Exptl. Zool.* 160:147-154.



Forman, M. and S.K. Majumdar. Lafayette College, Easton, Penn. Studies on the effects of monosodium glutamate on development and productivity of *D. melanogaster*.

Monosodium glutamate (MSG) is a widely used food additive. The first indication of possible ill effects from consumption of MSG was recorded in what has come to be known as the Chinese Restaurant Syndrome (Schaumberg, 1968) in man. Admin-

istrating high doses of MSG subcutaneously Olney (1969) and Olney and Sharpe (1969) produced brain lesions in the hypothalamus of mice and Rhesus monkey. As a result of these findings baby food manufacturers were asked to remove MSG from their products. Recent studies by Bazzano et al (1970) found no clinical or pathological changes in adult humans and adult gerbils when MSG was administered orally. Similarly Turner and Wright (1971) reported that 1% and 3% solutions of MSG caused no change in the development of *D. melanogaster*. Because of the conflicting reports the present investigation was undertaken to study the effects of MSG on the development and productivity of *D. melanogaster*. After initial work, the possibility of a lethal recessive sex-linked mutation was studied.

Eight female and eight male two-day-old Oregon-R flies were placed in vials containing Carolina Biological instant *Drosophila* medium. To the medium, one of the three solutions was added, with each solution being used in eight vials except for the sucrose where only four vials were used. One solution was pure distilled water, used as a control. A second control was a 0.7 M solution of sucrose. The third solution contained 0.7 M MSG. The  $F_1$  flies were removed after seven days and the  $F_1$  adults were counted on the seventeenth day according to sex and total number.

To determine the effect of MSG on the productivity, two-day-old Oregon-R male and female flies were allowed to drink 0.7 M solution of MSG for 24 hours. The drinking procedure involved soaking lens paper in the MSG solution and placing it in a bottle. Both sexes were kept together in order to insure mass culturing during consumption of the chemical. The control flies drank distilled water. After 24 hours the flies were transferred to vials containing instant medium and distilled water--two males and one female in each. The five-day brood system was used and four broods were obtained. Seventeen days after a brood was set up, the offspring were counted and the sex ratio was recorded.

It is apparent from Table 1 that 0.7 M MSG had an inhibitory effect on the development of the flies; as a result fewer  $F_1$  offspring were produced. Table 1 shows that the average number of  $F_1$  adults in the control (494.4) was greater than twice the average number of  $F_1$  flies in the MSG (215.1). However, the numbers of the  $F_1$  adult flies developed in the sucrose (519.5) and in the control were quite similar. The results of Turner and Wright (1971) seem to be contrary to the findings presented here. However, since they used only 1% and 3% solutions, it is possible that there is a threshold point somewhere between 3% and 10% solutions (0.7 M MSG is approximately 10%) at which the chemical produces an effect on the flies. An abnormal sex ratio occurred in the MSG cultures. There were 60.1% females in the MSG cultures compared with 49.95% in the control and 50.05% in the sucrose. A chi-square performed for determining the probability of the sex ratio obtained in the MSG cultures showed a significant difference ( $p < 0.001$ ).

Table 1. Effects of monosodium glutamate (MSG) on the development of *D. melanogaster*.

Treatment	No. of vials	Total no. of $F_1$ flies produced	Average no. of $F_1$ flies produced per vial	% female produced
Control (Water)	8	3955	494.4	49.95
Control (0.7 M Sugar)	4	2078	519.5	50.05
MSG (0.7 M)	8	1724	215.1	60.1

The results of the brood experiment are summarized in Table 2. In brood 1 (0-5 days), there was a control:MSG ratio of 2:1 for the average number of  $F_1$  progeny. In the second and third broods, the ratio fell to approximately 1.75:1.00 and 1.50:1.00 respectively, and in the last brood, the ratio was almost 1.0:1.2 with the MSG flies producing a greater number of offspring in the last brood. The total average number of flies in the control (423.52) was greater than the total in the MSG (261.94) by nearly 60%. It appears that MSG has more noticeable effect on the productivity in the first and second broods. The sex ratio in the MSG culture was not significantly altered.

Table 2. Five-day brood showing comparative productivity between control and 0.7 M MSG treated *D. melanogaster*

Treatment	Brood 1 0-5	Brood 2 5-10	Brood 3 10-15	Brood 4 15-20	Total	Sex Ratio
<u>Control</u>						
Total	3982	2887	1068	267	8204	
No. of vials	21	21	16	9		
Average	189.62	137.48	66.75	29.97	423.52	
Percent of total	44.77	32.46	15.77	7.00		
<u>Monosodium Glutamate</u>						
Male	1035	890	302	135	2362	1:1.017
Female	1058	834	287	144	2323	
Total	2093	1724	589	279	4685	
No. of vials	21	21	13	8		
Average	99.67	82.10	45.31	34.88	261.94	
Percent of total	38.05	31.34	17.30	13.31		

The Muller-5 method was used to determine the mutagenic activity of MSG. The male Oregon-R flies drank 0.7 M MSG for 24 hours and the flies were tested for sex-linked recessive lethal mutations. This test on 474  $F_1$  females yielded no recessive mutations.

References: Bazzano, G., J.A. Delia, and R.E. Olson 1970 Science 169: 1208-1209; Olney, J.W. 1969. Science 164: 719-721; Olney, J.W. and L.G. Sharpe 1969. Science 166: 386-388; Schaumburg, H.H. 1968. New Eng. J. Med. 278: 1122-1124; Turner, D.C. and C.P. Wright 1971. DIS 46: 118.

Thomas-Orillard, M. Faculté des Sciences de Paris VI, France. Influence of the culture medium on the number of ovarioles in *D. melanogaster*.

The phenotypic expression of the number of ovarioles is influenced by the environment. The rearing temperature (David et Clavel, 1967) and the food (Saviliev, 1928; David, 1960) have an important effect on this character. Control of

the temperature is always possible: all the experiments are carried on at  $25 \pm 0.5^\circ \text{C}$ . It is more difficult to appreciate the quantity of food which is necessary for a perfect ovary development. A study of the influence of the food on the phenotype expression of the ovarioles number is necessary to establish the experimental conditions for a study of the action of genes. Culture medium nature, density of population, sensibility of each instar larva to feeding were examined on two laboratory strains with very different geographical origins: one from France, the other from Japan and also on crosses between French and Japanese strains.

The phenotypic expression of the number of ovarioles is different on cornmeal medium and on yeast medium: t test with 98 degrees of freedom gives a significant value for t;  $P = .05$ . It depends also on the quantity of culture medium available for each larva. The biometrical characteristics of a strain are stable when all the larvae are well fed during the three instars. When first, or first and second or three instars are not well fed the mean of ovarioles number decrease significantly (F test between effect of the culture medium at different instar and residual variation gives  $F = 4.15$  for 3 and 41 degrees of freedom).

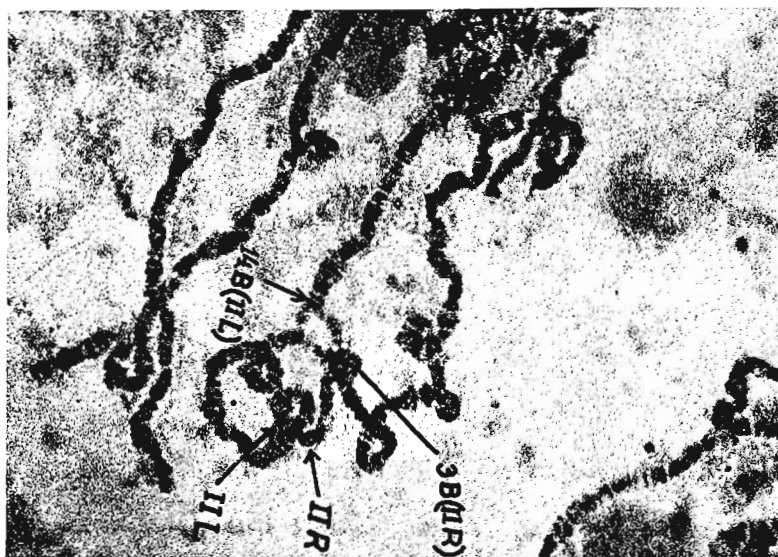
When the density of adult population does not exceed more than 50 animals for each culture bottle, the mean of the strain is stable; but when the density is bigger than 50 the mean decreases significantly from 36 ovarioles per female to 28. The ovarioles number of the females of the first generation is significantly greater than the arithmetic mean when the number of imagos is lower than 50 in the rearing bottles. It is not far from the arithmetic mean value when the density of the population varies from 100 to 150. We conclude, in the first case that there is evidence of heterosis, in the second case that genes have additive effects.

We see that in fact, this orientates the conclusions about the genes action. In all experiments on the genetic control of ovarioles number it is necessary to work with rearing bottles where the population density is maintained around 50. By this way, controlling the effect of feeding and working at constant temperature we can expect to run into the purely genetic problems.

Singh, V.K., M. Mishra and A.P. Jha.  
Bhagalpur University, Bhagalpur-7, Bihar,  
India. A new pericentric inversion in *D.*  
*ananassae*.

*D. ananassae* a member of the *melanogaster*  
species group, is highly polymorphic due  
to inversions in its natural populations  
(Kikkawa 1938; Dobzhansky and Dreyfus  
1943; Freire-Maia 1960; Ray-Chaudhuri and  
Jha 1965 and Futch 1966). We are report-

ing, herewith, for the first time, a new pericentric inversion on the second chromosome of  
*D. ananassae* from its Bahadurpur population. Bahadurpur is a sparsely situated village with-  
in a dense forested area known as Samtha forest in the State of Bihar. Breakage points in  
this inversion were determined from the reference map prepared by Ray-Chaudhuri and Jha  
(1965). One of the breaks has occurred in region 14A of IIL and other in 2B of IIR as shown



above. Freire-Maia (1960) and Futch (1966) respectively reported pericentric inversions on  
the second chromosome of *D. ananassae* from Brazil and South pacific islands. Our report is  
a new one in that its breakage points are located on the regions different from those report-  
ed by them.

Voss, R. Hebrew University of Jerusalem,  
Israel. A common suppressor for a lethal  
mutation and a forked mutation.

Recent experiments with lethal  $1^{3DES}$ ,  
a suppression of which was described in  
DIS#46, had revealed that the suppression  
of forked was not induced simultaneously  
with the suppression of the lethal, but

was associated with the lethal  $3^{DES}$  originally. This is in accord with A. Schalet's findings  
in mapping the proximal X chromosome region (DIS 46, 131). However, the reversion of the  
lethal does not cause reversion of *su-f*. The revertant flies still show suppression of fork-  
ed. Therefore it seems necessary to call the new suppressor  $su-1^{3DES}$  and not  $su-f^V$  as was  
suggested before. All features of the suppressor which were described previously still hold.  
To this may be added that the suppressor seems to be a Y suppressed lethal as XO males are  
inviable and homozygous females  $1^{3DES}$ ,  $su-1^{3DES}$  with a Y chromosome are viable, although  
sterile. The independent reversion may be interpreted to mean that  $1^{3DES}$  covers more than  
the *su-f* locus only, or that there is more than one function associated with *su-f* which can  
be separated from it.

Jha, A.P., M. Mishra and V.K. Singh.  
Bhagalpur University, Bhagalpur-7, Bihar,  
India. Abnormal sex ratio in Darjeeling  
*Drosophila* population.

Attempts were made to collect *Drosophila*  
at Darjeeling at an altitude of 7000 ft.  
in India from October 5 to October 10,  
1970. Our collection of 776 flies com-  
prised seven good species (table below).  
Out of them only 276 were females. The

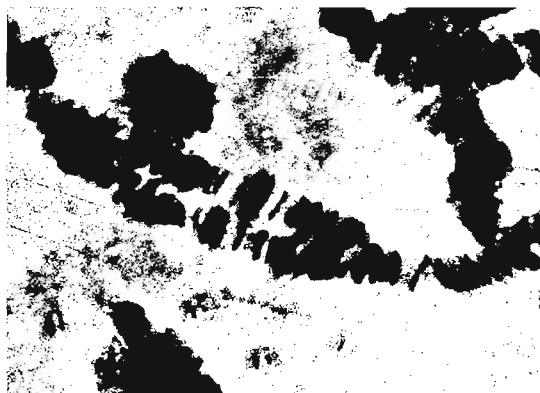
collection was made by using fermented banana, orange and pineapple mesh as baits. Traps  
were suspended in air by hanging them upwardly. Morning and evening collections were made.  
The population was so thin that we could collect only 2 to 3 individuals of different species  
on the 5th and 6th of October from each trap. Probably it was due to extreme winter (50-  
55°F). Then we put some traps in the evening of October 7th inside some bakeries whose  
temperature was higher than outside. From the morning of October 8th the flies started  
pouring into the traps. Specimens were identified in collaboration with Dr. J.P. Gupta,  
*Drosophila* Laboratory, Banaras Hindu University, India. *D. bipectinata* was the dominating  
species. *D. ananassae* and *D. melanogaster* were next in number to visit the traps.

Table 1. Data showing the sex ratio of *Drosophila* flies collected at Darjeeling

Species	Males	Females	Total
<i>D. bipectinata</i>	148	76	224
<i>D. ananassae</i>	129	93	222
<i>D. melanogaster</i>	141	56	197
<i>D. busckii</i>	41	13	54
<i>D. kikkawai</i>	23	18	41
<i>Chaetodrosophila quadrilineata</i>	12	17	29
<i>D. malerkotliana</i>	6	3	9
	500	276	776

An ideal sex ratio in a population is 50M:50F. But this equality of the number of  
sexes has not been favoured in any of the *Drosophila* species, herewith reported in the en-  
vironmental conditions of Darjeeling. If two or more males are necessarily required to  
inseminate a female *Drosophila*, in that case the males may preponderate in the population.  
We do not know if any male producing tendency has acquired in the heredity of Darjeeling  
populations of these *Drosophila* species. The problem is intricate and we leave its solution  
for the future.

Krimbas, C.B. Agricultural College of Athens,  
Athens (Votanikos), Greece. A newly spontane-  
ously formed chromosome arrangement in one  
salivary gland cell of *D. subobscura*.



In one of the salivary gland cells of  
an individual of *D. subobscura* a small  
inversion or deletion has been detected  
in heterozygous condition near the  
centromere end of chromosome O (region  
76 of the map of Kunze-Mühl and E. Müll-  
ler, Chromosoma 1958) shown in the  
photograph. All other cells of the sali-  
vary gland of the same individual did  
not show heterozygosity in that region.  
It is apparently a newly formed chromo-  
somal arrangement in only one salivary  
gland cell of this individual.

Marengo, N.P. and S.H. Vernick. C.W. Post College of Long Island University, Greenvale, New York. Virus-like particles in nuclei of muscle fibers of genetically "rotated" prepupae of *D. melanogaster*.

The gene abdomen rotatum (*ar*) was discovered and named by Beliajeff (1931). Marengo and Howland (1942) described its effect on the development of the fly and established that the actual rotation of the imaginal abdomen coincided with the termination of the prepupal period

and the movements attendant on the change from prepupa to pupa. In addition to the rotation of the pupal abdomen within the completely symmetrical pupa case, abnormalities of the puparium appeared which were clearly attributable to exaggeration of the function of the persisting larval muscles. Robertson (1936) states that these muscles are responsible for the contraction of the larval cuticle in puparium formation and also the movements of the individual as it changes from a prepupa to a pupa, with a clearly defined head, thorax and abdomen. Marengo and Howland (1942) suggested that these muscles were also responsible for the actual rotation of the pupal abdomen since the time of the first recognition of actual rotation coincided with the movements characterizing the change from prepupa to pupa. The most clearly recognizable puparial abnormalities were persistent segmentation of the hardened cuticle and abnormally large lateral bulges of the puparial wall between the attachments of the dorso-ventral muscles.

**MATERIALS AND METHODS:** Since abdominal rotation prevents normal copulation, the stock used in the 1942 study was a balanced lethal with eyeless Dominant (*ar/ey<sup>D</sup>*). This stock apparently has not survived, and the current study was made possible through the courtesy of Pollards Wood Research Station of the Institute of Cancer Research, Royal Cancer Hospital, Buckinghamshire, England, which supplied a new balanced lethal with cubitus interruptus (*ar/ci<sup>D</sup>*).

Since the structural abnormalities of the puparia of "rotated" flies appeared to be a direct consequence of abnormal muscle function, and since paraffin sections for light microscopy revealed no consistent structural muscle abnormality, it was decided to examine the prepupal muscles of "rotated" (*ar/ar*) individuals and normals (*+/ar* and *+/+*) by electron microscopy.

Prepupae with puparial abnormalities identifying them as genetically "rotated" were opened in 3% buffered glutaraldehyde and the muscle fibers dissected out. After three hours these were washed overnight in a rinsing buffer consisting of 10 gm. sucrose in 100ml Sorenson's stock buffer, post-fixed two hours in Palade's buffered osmium fixative, pre-stained three hours in .5% uranyl acetate, dehydrated and embedded in epon. Sections were cut with a diamond knife on a Reichert OMU-2 ultramicrotome, stained in uranyl acetate and lead citrate, and examined with an Hitachi HS-7 electron microscope. Identical treatment was given to muscle fibers of normal (*ar/ci<sup>D</sup>*) individuals and *+/+* individuals of Oregon "R" stock.

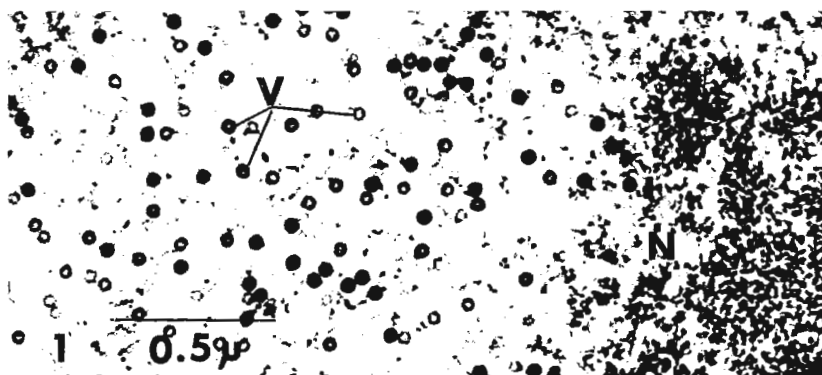


FIG. 1. Section through nucleus of muscle cell from prepupa homozygous for abdomen rotatum (*ar/ar*). Virus-like particles (V) are apparent in the nucleoplasm which has a characteristic empty appearance. Part of the nucleolus (N) is visible. X49,000.

**RESULTS:** At the present stage in this study, no readily identifiable ultrastructural difference was found between the fibrillar organization of the muscles from the "rotated" prepupae and those of the normal individuals. However, a high percentage of the muscle nuclei of genetically "rotated" prepupae contained virus-like particles approximately 48mμ in size (Fig. 1).

In a number of cases, they appear to be budding from a local electron-dense region of the nucleoplasm (Fig. 2). No particles were observed in nuclei from the muscles of either the normal heterozygotes (*ar/+*) or the *+/+* Oregon "R" wild type.

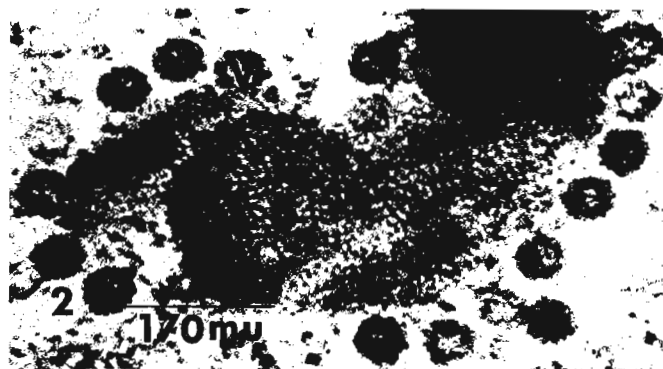


FIG. 2. Section through nucleus showing particles (V) which appear to be budding from a faintly crystalline nucleoplasmic condensation. X145,000.

DISCUSSION: The classification and significance of the observed particles is at present not known. Since abdomen rotatum as originally described by Beliajeff (1931) was a mutant of chromosome IV it might be speculated that the presence of a virus may relate to the carrying of information from the mutant DNA of chromosome IV to the larval muscles. These muscles bring about through exaggerated contraction both the puparial abnormalities described and the rotation of imaginal abdomen which is characteristic of the adult phenotypic expression of this gene.

If these particles are carriers of the *ar* codon, their isolation from genetically "rotated" individuals might

provide a means of experimental transformation of normal larvae capable of metamorphosis into abnormal prepupae, pupae and "rotated" adults.

SUMMARY: Nuclei of muscle fibers of genetically "rotated" (*ar/ar*) prepupae of *D. melanogaster* examined by electron microscopy were found to contain virus-like particles not previously described. These particles were not observed in identical preparations of Oregon "R" prepupae and normal heterozygotes of the *ar/ci*<sup>D</sup> stock.

References: Beliajeff, N.K. 1931 Erbliche Asymmetrie bei *Drosophila*. Ein neues Gen im IV chromosom von *D. melanogaster*. Biol. Zbl. 51:701-709; Marengo, N.P., and R.B. Howland. 1942 The effect of the gene abdomen rotatum on the development of *D. melanogaster*. Genetics 27:604-611; Robertson, C.W. 1936 The metamorphosis of *D. melanogaster*, including an accurately timed account of the principal morphological changes. J. Morph. 59:351-359.

Supported in part by a faculty research grant from C.W. Post College.

Doane, W.W. Yale University, New Haven, Connecticut. Isoamylases in *Drosophila hydei*: a system for the analysis of gene-specific puffing activity.

$\alpha$ -Amylase activity in the larval gut of *D. hydei* is essentially restricted to a small region at the anterior end of the posterior midgut. The region is characterized by large secretory cells with polytene chromosomes and, unlike the situation in *D. melanogaster*, poly-

teny here is of a magnitude readily open to cytological investigation of gene-specific puffing activities. As a secretory protein whose activity may be manipulated by dietary starch, amylase is an ideal subject for such a study (Doane, W.W., 1969, pp. 73-108 in "RNA in Development" E.W. Hanly, ed., Univ. of Utah Press).

A single structural gene for Amylase (*Amy*) is located on chromosome 5 in *hydei*, between *cn* and *vg*. Making use of the latter markers and two electrophoretic *Amy* variants, a screening program was set up to select for potential X-ray induced deficiencies of the *Amy* locus, and so position the locus on the cytological map of the chromosome (Doane, W.W., 1971, Isoz. Bull. 4:46-48). Of the strains selected, one is particularly useful: it contains a fifth chromosome with an inversion near the center. The inverted section is apparently accompanied by deficiencies of the *Amy* and *vg* loci, perhaps at opposite ends. Tentative location of the break points are: proximally, in 107A before the doublet, and, distally, in 109C, between bands 7 and 8 (according to the map of H. Berendes, 1963, Chromosoma 14: 195). The inversion has been examined in polytene chromosomes from salivary glands and midguts (anterior and posterior regions) taken from larvae reared on starch- and/or sugar-yeast diets with promising results. (Supported by grant NSF GB 8607.)

Sunanda Mahajani, Division of Genetics, Indian Agricultural Research Institute, New Delhi-12, India. Crossing over in the inversion carrying second chromosome of a *D. melanogaster* male.

Two facts are known which strongly suppress crossing over in *D. melanogaster*. These are: (a) sex and (b) chromosome inversions. Crossing over in *D. melanogaster* males is a very rare event. Likewise, pairing difficulties imposed by inversions effectively prevent crossing

over in chromosomes carrying them; infrequently, however, cross-overs in inversion heterozygotes of *D. melanogaster* have been reported (Kaufman and Gay, 1970, DIS, 45:81). In this communication I report a rare cross-over event that has occurred in the second chromosome of a male bearing the In(2L)Cy and In(2R)Cy inversions.

Three cinnabar-eyed mutant females were obtained among *D. melanogaster* larvae reared on basal medium containing 0.03% nitrosoguanidine. Mutant virgins were mated to Cy/B1 L<sup>2</sup> males with two objectives in view: (a) to confirm the second chromosome location of the cinnabar mutation and (b) to keep the treated chromosome intact. Two types of flies should normally be expected in the progeny - Curly (Cy) and Bristle Lobe (B1 L<sup>2</sup>). Actual results, however, showed that in addition to the above classes, Curly Bristle Lobe (Cy B1 L<sup>2</sup>) and wild type (+++) flies were also obtained (3.4% and 1.1% respectively; 172 flies were observed).

The recovery of these unusual types of flies can be explained on the basis of a cross-over between the Cy and B1 loci, presumably in the region of the chromosome lying between the two inversions. This would bring the Cy, B1 and L<sup>2</sup> markers on one chromosome and their respective (+) alleles on the other.

The recovery of cross-over products observed by me is unusual in two ways. Not only has a cross-over occurred in a male, a rare event in itself, but it has taken place in spite of the inversions that should have normally prevented it.

Mahowald, A.P. Institute for Cancer Research, Fox Chase, Philadelphia, Pennsylvania. Intracellular symbionts of *Drosophila*.

Many workers have demonstrated with the electron microscope the presence of Rickettsia or bacteria within cells of *Drosophila*. King (1970) has recently reviewed these findings and presented a detailed account of hitherto unpublished work from his labora-

tory. Since many laboratories are actively studying the nucleic acids of eggs of *Drosophila*, it is important to know whether there are bacterial contaminants within the cells. King points out the widespread occurrence of these organisms, which he terms "A-bodies", so that one gains the impression that they could be found in any stock examined with the electron microscope. But in extensive ultrastructural studies of oogenesis in two strains of *D. melanogaster* (Oregon R and Cochapontset, obtained from T.R.F. Wright), *D. hydei* (New Haven) and Zürich strains, obtained from S.J. Counce-Nicklas), *D. virilis* (Johns Hopkins), and *D. immigrans*, they have not been found. Furthermore, they have not been detected in my stocks of fs(1)N, dor 1(1)X2, 1(1)ff11 (obtained from S.J. Counce-Nicklas). On the other hand, they are present in high concentration within the oocytes of *D. willistoni*, Barbados III obtained from D.F. Poulson. In this species the distribution differs considerably from previous reports. Ullmann (1965) found  $\epsilon$ -granules (these appear to be the same as A-bodies) only in Stage 13 oocytes and later stages, and suggested that they may form from the oolemma. Furthermore, she only found them at the posterior tip of the embryo. In the strain of *D. willistoni* I have studied extensively, these structures are found in the oogonia and all cells derived from these cells, i.e., the nurse cells and oocytes. They have not been found in follicle cells. A brief survey of other adult and larval tissues also failed to detect any outside the germ tissue. During embryogenesis they are at first found throughout the ooplasm, but gradually they become concentrated in the posterior portion of the egg. Most of these A-bodies are included in the pole cells. A few A-bodies are also found in the blastoderm cells, even at the anterior end. Since apparently there are few, if any, A-bodies in somatic tissue, there must be some restriction on their growth outside the germ line. The role of these A-bodies or Rickettsia is unknown.

Reference: King, R.C. 1970, Ovarian Development in *Drosophila melanogaster*. Academic Press, New York. 227 pp.



Félix, R. and M.E. de la Rosa. Genetics and Radiobiology Program of the National Commission of Nuclear Energy, Mexico. Cytogenetic studies with sodium cyclamate in *D. melanogaster* females.

It is now almost 30 years since the first highly effective mutagens were detected. Since then, much interest has been concentrated on the agencies which produce mutations and chromosome breaks. By the application of chemical compounds, some of whose effects on the constituents of

the cell and nucleus are already known, it may be possible to obtain more information on the nature of the mechanisms involved in the production of genic and chromosomal alterations. Many agencies produce mutations and chromosome breaks. Their variety is such that it was wondered whether they do not achieve the same end effect by different means. Several mechanisms of mutagenesis were imagined, such as energy transfer to the chromosomes, chemical reactions with the genetic material, and interference with chromosome synthesis. Few of these would act equally well at all stages of the cell cycle. Thus, interference with chromosome synthesis should be restricted to interphase. Within the last two decades much interest has been concentrated on the variations in differential sensitivities exhibited by cells during gametogenesis in both plants and animals. These variations may be inherent in that they depend on processes or physiological states in the test object which, while sometimes recognized, cannot always be controlled, or they may be induced by different chemical and physical agents.

The reports of Sax and Sax (1968), D. Stone et al. (1969) and Legator et al. (1969) pointed out the cytogenetic damage induced by cyclamate and by its degradation product, cyclohexylamine, preceding the announcement of cyclamate's possible carcinogenic capabilities and the subsequent restrictions on dietary consumption.

Stone et al. (1969) demonstrated that cyclamate, in a minimum concentration of 200 microgram/ml can stimulate chromosome breakage in human cells in vitro. Whereas a high dosage (equivalent to 15 g/175 Kg) was required to obtain a demonstrable increase in chromosome breaks, Stone pointed out that there is some evidence of synergistic actions on chromosome damage between X-irradiation and radiomimetic chemicals (Merz et al., 1961); between the chemical agents and virus (Nichols et al., 1965), and between the chemical agents themselves (Moutschen-Dahem, 1962).

A considerable number of workers are now engaged in testing compounds for their mutagenic activity. Since in many cases the publication of negative data is not warranted, there is often a repetition of effort with similar negative results.

This study is concerned with the production of X chromosome loss and non-disjunction in *Drosophila* females by sodium cyclamate. Obtaining identifiable meiotic stages is possible in the case of the female, since the meiotic divisions do not occur until after the oocyte is laid. To obtain fairly uniform samples of a single stage it is necessary to limit the period of egg collection, so that not more than one oocyte is recovered from each ovariole. On the other hand, in the male, meiosis occurs long before the completion of sperm development and the insemination of the female, and at best one can distinguish meiotic stages from pre and post-meiotic ones, with little hope of subdividing the various stages of development of the spermatocyte.

Radiation studies to date have largely been restricted to older stages of oocytes in the vitellarium. King et al. (1956) have described the structure of the ovariole in the adult female, and have designated 14 developmental stages of the oocyte. They recognize three sensitive groups on the basis of recessive lethal mutations, X chromosome losses, and dominant lethal effects. However, much of the work of others has been concerned with but two of the stages they describe, stages 7 and 14 (Parker, 1963), which are, respectively, the oldest stages in the newly emerged females and the fully mature, chorionated oocyte found in females ready to begin egg laying (usually during the second day of adult life). Apparently strains differ in the rate of egg production, as well as in the number of stage 14 oocytes in each ovariole in 4-day old females (Williamson and Stubblefield, 1970). In the control group of our experiment an average of 30 eggs were obtained in 24 hours which correspond mainly to stage 14 oocytes.

Bridges (1913) identified non-disjunction by the recovery of exceptional females and males among the progeny, by their being matroclinous and patroclinous, respectively, in phenotype for sex linked characters. In the present work an improved method for detecting non-disjunction and chromosome X loss that gives particularly reliable evidence concerning the origin of each exceptional female and male makes use of a tester male stock with attached  $Y^{SX} \cdot Y^L$  chromosomes. This stock was derived from translocations between the X and Y chromo-



somes and has the markers (y) yellow and (B) Bar.

The females were taken from the cross of stocks having the  $sc^8Y$  chromosome:  $y^2 w^a/y^2 w^a$ ;  $e/e \times y^2 w^a/sc^8Y$ ;  $e/e$ . The existence of any secondary exceptions (from XXY mothers) among the  $y^2 w^a/y^2 w^a$  females was made unlikely by the  $sc^8Y$  chromosome which covers the effect of yellow in XXY females. The males have an attached  $Y^{SX} \cdot Y^L$  chromosome of the genotype:  $In(1)EN, Y^S B y \cdot Y^L$ . When virgin females are isolated, the marker ebony insures the virginity of such females, as the genotype of the females from the cross:  $y^2 w^a/y^2 w^a$ ;  $e/e \times In(1)EN, Y^S B y \cdot Y^L$ ;  $+/+$  is heterozygous for the ebony marker. The fertilization of an XX egg with an XY spermatozoon would produce meta-females with low viability which were excluded from the following analysis, whereas the fertilization of eggs of the same non-disjunctional chromosomal constitution with a non X chromosome bearing spermatozoon would produce matroclinous yellow, white apricot females of the same genotype as their mothers. These females are easily identified from their normal Bar eyed sisters. The no-X egg when fertilized with an SY bearing sperm would become an XY patroclinous male, which can be identified by the Bar, yellow eyes in its phenotype.

Virgin females of the genotype  $y^2 w^a/y^2 w^a$  were aged from 4 to 6 days and fertilized in cultures containing agar-cornmeal medium with sodium cyclamate. In each culture an aged female was mated with 3 attached  $Y^{SX} \cdot Y^L$  males, and eliminated after 24 hours. All the cultures were kept at  $25 \pm 1^\circ C$  throughout the experiment.

The  $F_1$  flies were scored for X-loss and non-disjunction from 13 to 15 days after the treatment with cyclamate of the P females. The female fruit fly has remarkable synthetic abilities, since during the period of maximum egg production it ingests a daily amount of yeast which approximately equals its body weight and manufactures from the raw material a quantity of eggs which approximately equals 1/3 its weight (King and Wilson, 1955).

Table 1. Progenies obtained from 28 cultures after feeding *Drosophila* adult females with media containing sodium cyclamate.

NaCy mg/ml	Control	0.05	0.10	0.20	0.40	0.80	1.60
$\delta$ (m.p.c.)	17.14	16.57	17.75	13.21	12.50	13.25	11.32
$\phi$ (m.p.c.)	12.46	9.79	13.46	10.82	8.29	9.11	7.82
$\delta/\phi$ (s.r.)	1.37	1.69	1.32	1.22	1.51	1.45	1.45
$\delta$ (total)	480	464	497	370	350	371	317
$\phi$ (total)	349	274	377	303	232	255	219

NaCy mg/ml, sodium cyclamate, milligram/milliliter; (m.p.c.) mean per culture; (s.r.), sex ratio.

As no exceptional progeny were obtained among 4,858 flies, a second experiment was started, maintaining during 6 days, 1 female previously aged during 4 days and mated to 3 males, from the stocks described above, in each culture. The agar-cornmeal medium contained no sucrose. After six days the P flies were eliminated. The feeding of the progenitors as well as the egg laying, embryonic and larval development of the progeny occurred in the medium with cyclamate. The results of this second experiment are contained in Table 2.

Table 2. Progenies obtained from adult and larval feeding during six days in media without sucrose and several concentrations of sodium cyclamate.

NaCy mg/ml	Control	10.00	40.00	100.00
$\delta$ (m.p.c.)	45.74	25.00	31.33	34.89
$\phi$ (m.p.c.)	25.16	11.39	12.00	17.33
$\delta/\phi$ (s.r.)	1.82	2.19	2.61	2.01
$\delta$ total	869	575	658	314
$\phi$ total	478	262	252	156
(n.c.)	19	23	21	9
(n.d.p.)	-	-	-	4
(c.l.p.)	-	-	2	9

NaCy mg/ml, sodium cyclamate, milligram/milliliter; (m.p.c.), mean per culture; (s.r.), sex ratio; (n.c.), number of cultures; (n.d.p.), non-junctional progeny (c.l.p.), chromosome loss progeny.

From Day and Grell (1966) the non-disjunction frequencies are calculated by the expression:

$$\% \text{ non-disjunction} = \frac{4 (\text{exc. } \phi\phi \times 100)}{\text{total} + \text{exc.}}$$

In multiplying the exceptional females by 4 and adding the exceptions to the denominator it is assumed that the number of exceptional males arising from non-disjunction is equivalent to the number of exceptional females and that one half of all XX and O oocytes are lost due to lethality.

The following expression was applied to calculate the frequencies of X chromosome loss among the

$$\text{progenies: } \% \text{ loss} = \frac{2 (\text{exc. } \delta\delta - \text{exc. } \phi\phi) \times 100}{\text{Total} + \text{exc.}}$$

The excess of exceptional males over exceptional females is considered to arise from loss of an X-chromosome during meiosis and not from non-disjunction, as the excess of exceptional male progeny is best explained as arising from spontaneous loss (Mavor, 1924; Patterson et al., 1932; Sturtevant and Beadle, 1936; Uchida, 1962). Oocytes lacking an X will lead to viable progeny only when fertilized by an X sperm; hence a correction factor of 2 is used to account for the nullo-X oocytes fertilized by a Y sperm (Day and Grell; 1966).

From table 2, exceptional progenies was restricted only to the concentrations of 40 mg/ml and 100 mg/ml. The computation of such data gives the following percentages of non-disjunction and chromosome loss:

NaCy mg/ml	40	100
XO	0.44	2.07
XX.Y	-	3.31

A comparison among the data from Stone et al. (1969) on induced chromosome breakage in human cells in vitro with a minimum concentration of 0.20 mg/ml of cyclamate and the minimum effective dose from this experiment (40 mg/ml) demonstrates a difference of sensitivity, equivalent to two orders of magnitude. Although the two experimental designs are hardly comparable, it seems evident that *Drosophila* oocytes are much less sensitive to cyclamates than human cells in vitro.

In a third experiment female flies aged for 3 days were fed during 24 hours with sodium cyclamate dissolved in distilled water at concentrations of 50 mg/ml, 100 mg/ml, and 160 mg/ml. (See Technical Note by R. Félix). Each treated female was mated afterwards with 3 males in individual cultures containing normal agar-cornmeal medium. After 6 days of egg-laying the flies were eliminated and the progenies were scored after 13 to 15 days (Tab. 3)

Table 3. Progenies obtained from the feeding of adults with concentrated solutions of sodium cyclamate.

NaCy mg/ml	50.00	100.00	160.00
♂ (m.p.c.)	30.75	36.25	35.31
♀ (m.p.c.)	21.78	21.15	22.95
♂/♀ (s.r.)	1.41	1.71	1.54
♂ (total)	861	725	671
♀ (total)	610	423	436
(n.c.)	28	20	20

NaCy mg/ml, sodium cyclamate, milligram/milliliter; (m.p.c.), mean per culture; (s.r.), sex ratio; (n.c.) number of cultures.

No exceptional progeny among 3,726 flies were obtained after the feeding of *Drosophila* with sodium cyclamate solution, in spite of the 160 mg/ml concentration, which is close to the saturation point.

Comparing the 100 mg/ml concentration in the food medium with the same concentration in solution, the size of the progenies are remarkably alike.

An aspect of the experiments, which called our attention, is the deviation of the sex ratio, indicating a lowered proportion of the X.XY females ( $y^{2wa}/\text{In}(1)\text{EN}$ ,  $Y^S$ ,  $By.Y^L$ ) as compared with the XO males ( $y^{2wa}$ ). The deviation that is also found in the control groups, is considerably

magnified by the prolonged treatment of adults with cyclamate, as shown in Table 2.

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Félix, R., J. Guzmán and A. de Garay Arellano. Genetics and Radiobiology Program. National Commission of Nuclear Energy. Mexico City, Mexico. Distribution of CO<sub>2</sub> sensitivity (sigma virus) in an urban population of *D. melanogaster* from Mexico City. II. Low dispersal, a factor which explains differences among locations.

Wright's (1955) model for optimum evolutionary opportunity is one of partially isolated sub-groups within which there is some opportunity for gene frequency shifts due to environmental differences between sub-populations or due to random processes (Crow, 1955). Chance gene frequency changes will be greater as populations become smaller, whereas fluctuations in selection coefficients have a similar effect

in larger populations. Kimura (1955) worked on the mathematically difficult problems of constructing the stochastic process of gene frequency change under various assumed conditions. The general conclusions of Kimura's work were anticipated by the earlier writings of Wright. The fluctuations in gene frequencies in sub-populations may be provided by random shift in small populations or by random changes in selective values in larger ones.

Wright (1951) considered a model in which the population has a homogeneous structure, but offspring travel only a certain distance from their parents. If the range of dispersal is restricted in such a way that the parents of a particular individual may be assumed to be drawn at random from a neighborhood of a certain size, the amount of local differentiation may be related to the effective size of the neighborhood.

In the study of Wallace (1966a) of the allelism, a model was applied, which predicts that for lethals collected at different times, the logarithm of the frequency of allelism due to inbreeding, should decline linearly with time. On a strictly empirical basis, Wallace (1966b) substituted the square root of the distance for time in the original model. The relationship was suggested by the dispersal of flies from a point of release: the logarithm of the numbers of recaptured flies decreases linearly with the square root of distance from the point of release.

Wallace analyzed the data of local collections of several species of *Drosophila* given by Dobzhansky and Wright (1943), by Burla et al. (1940) and by Timofeeff-Ressovsky (1941 a,b), which show that, despite its seemingly well developed powers of dispersal, (revealed by the migration of individual flies for several hundred meters in two days), more than one-quarter of the *D. pseudoobscura* captured at a given point have probably arisen within a radius of 25 meters from that point; one-eight of such collection of flies have probably arisen within a radius of ten meters. The data for *D. willistoni* reveal that this species is ever more sedentary; nearly one third of a collection of these flies have probably arisen within a radius of some ten meters from the point of capture. The data of Dobzhansky and Wright on the dispersal of *D. melanogaster*, suggest that the dispersal of this species resembles that of *D. funebris* and *D. willistoni* more closely than that of *D. pseudoobscura* (Wallace, 1966b).

When the linear relationship between the logarithm of the frequency and the square root of distance is plotted on a conventional arithmetical scale, the transformed curve clings to both the vertical and the horizontal axes. Wallace gives the following interpretation of this theoretical curve: immigrants form a small fraction of any population but, given that an individual is an immigrant, he is very nearly as likely to have come from any one distance as from another. The tall vertical portion of the curve, which hugs the Y axis means that the bulk of all individuals found at one spot has arisen initially at or very near that spot. Bateman (1947) refers to this feature of the dispersal pattern, pointing out its effect of subdividing what would otherwise appear to be a continuous population.

In this report, data are given, which suggest the low dispersal of *D. melanogaster* flies, when the percentage of sensitive flies recovered from neighboring collection sites is recorded.

Brun and Sigot (1955) pointed out the differences among stabilized and non-stabilized CO<sub>2</sub>-sensitive strains of *Drosophila*. In stabilized strains there is a maternal effect which assures the sensitivity to all the progeny, whereas stabilized males transmit the sigma virus in his sperm to a fraction of the offspring. Non-stabilized lines, on the other hand, typically produce resistant and sensitive progeny. The role of infectious virus sigma may be clarified by future investigations of the relationship between sigma and the *Drosophila* sensitivity to CO<sub>2</sub> (Seecof, 1962).

L'Héritier (1958), L'Héritier and Plus (1963) made an extensive research on the variations of the genotype of the virus, as well as on the genetic factors of the host which affects his hereditary transmission. Every quantitative aspect of the relationship between virus sigma and its host seems to be more or less genotype dependent.

The  $\text{CO}_2$ -sensitivity itself, is not a selective factor to be considered in population studies, as the concentration of the gas required to produce anoxia in *Drosophila*, is never found in natural conditions. However, the presence of sensitive, resistant, and "refractory" flies in the same population, points out that the presence of the hereditary virus is not a completely neutral character.

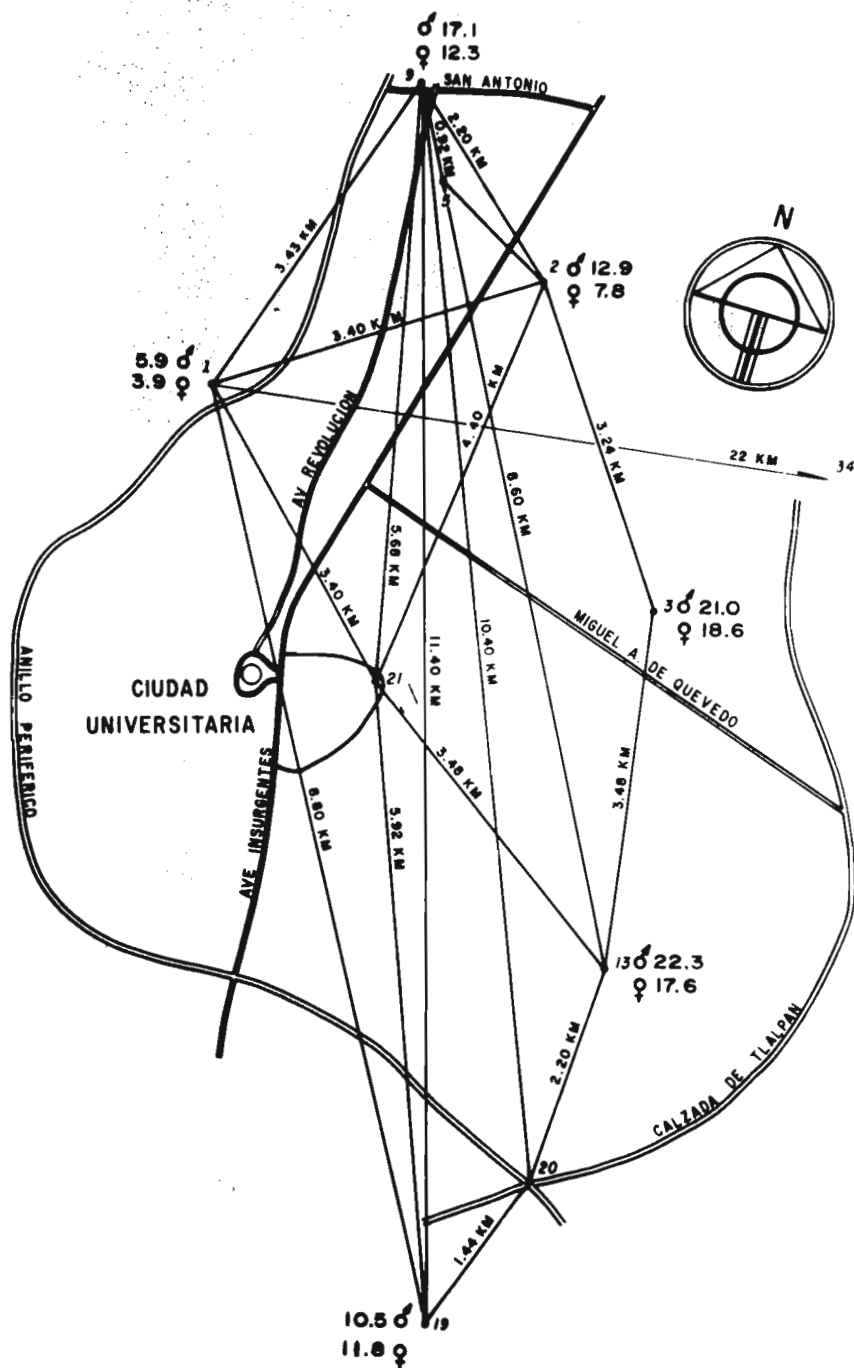


Fig. 1. Distribution of  $\text{CO}_2$  sensitivity of *D. melanogaster*. Percentages of sensitive males and females are indicated at six locations.

Seecof (1964) investigated the deleterious effects of the sigma virus infection of *D.*

melanogaster, demonstrating a direct relationship among deaths of the infected progeny and the logarithm of the number of infectious units inherited by the progeny. Seecof tested the effect of the variation of environmental conditions, looking for a selective advantage of the flies when they are infected. As a result, the infection proved to be a handicap under all the conditions tested. The presence of the virus in wild populations of flies, in absence of any known vector, is therefore, unexplained. However, the crowding of flies did not reduce the proportion of CO<sub>2</sub>-sensitive flies in the cultures. There is to be found, a compensating selective advantage for sigma-infected flies, in order to explain the proportion and persistence of such flies in wild populations.

In the present survey, the proportion of CO<sub>2</sub>-sensitive flies from six locations in Mexico City was tested by submitting them to the treatment with pure CO<sub>2</sub> at 8°C during 15 minutes. Collections were made from June 1969 to May 1970, in a southwest section of Mexico City. Flies were attracted to 1/4 liter containers of decaying cantaloupe. The spatial distribution of CO<sub>2</sub>-sensitivity, as well as the recorded averages of the monthly percentages of sensitive males and females from six locations are shown in Fig. 1. Only the data from the locations where the collections were significant, are included. The minimum and the maximum distances among locations, were respectively, 2.20 and 11.40 Km. The extreme difference of CO<sub>2</sub>-sensitivity was found among locations 1, and 3, approximately 4.70 Km. apart.

Monthly percentages of CO<sub>2</sub>-sensitivity in both sexes, from each location, are given in tables 1 and 2. The annual averages of the monthly percentages of sensitive males and females are shown in Table 3. These averages, are constantly larger for males than for females (ex-

Table 1. Percentage of CO<sub>2</sub>-sensitivity of males collected at six locations from Mexico City.

Loc.	June		July		August		September		October		November	
	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S
1	106/6	5.6 ±2.02	60/5	8.3 ±3.56	575/28	4.8 ±0.88	1426/139	9.7 ±0.78	261/33	12.6 ±2.05	60/1	1.8 ±1.33
2	215/32	14.8 ±2.39	198/30	15.1 ±2.54	213/28	13.1 ±2.31	43/5	11.6 ±4.88	18/4	22.2 ±9.79	-	-
3	117/9	7.6 ±2.44	339/32	9.4 ±1.69	130/25	19.2 ±3.45	149/16	10.7 ±2.53	150/0	0.0	-	-
9	-	-	75/11	14.6 ±4.07	-	-	61/3	4.9 ±2.76	26/1	3.8 ±3.74	-	-
13	60/23	38.3 ±6.27	-	-	93/14	15.0 ±3.70	18/5	27.7 ±10.54	-	-	-	-
19	-	-	70/10	14.2 ±4.17	52/4	7.6 ±2.01	158/11	6.9 ±2.01	101/6	5.9 ±2.34	46/1	2.1 ±2.11
	498/70	14.06 ±1.66	742/88	11.86 ±1.49	1063/99	0.3 ±0.99	1855/179	9.64 ±0.75	556/44	7.91 ±1.33	106/2	1.89 ±1.31

Loc.	December		January		February		March		April		May	
	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S
1	126/3	2.3 ±1.75	-	-	90/3	3.3 ±1.88	73/2	2.7 ±1.87	259/21	8.1 ±1.69	658/39	5.9 ±0.91
2	68/0	0.0	-	-	-	-	71/3	4.2 ±2.38	146/12	8.2 ±2.51	-	-
3	15/2	13.3 ±8.76	-	-	-	-	-	-	45/13	28.8 ±6.75	14/8	57.1 ±13.22
9	-	-	-	-	-	-	-	-	114/26	22.8 ±3.92	38/15	39.4 ±7.92
13	21/0	0.0	-	-	-	-	-	-	14/1	7.1 ±6.34	-	-
19	-	-	-	-	-	-	-	-	19/5	26.3 ±10.10	-	-
	230/5	2.17 ±1.77			90/3	3.33 ±1.88	144/5	3.47 ±1.52	597/78	13.07 ±1.55	710/62	8.73 ±1.78

N, number of collected flies; S, number of sensitive flies; %S, percentage of CO<sub>2</sub>-sensitive flies ± standard error.

Table 2. Percentage of CO<sub>2</sub>-sensitivity of females collect at six locations from Mexico City.

Loc.	June		July		August		September		October		November	
	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S
1	75/0	0.0	41/2	4.8 ±3.34	629/33	5.2 ±0.88	1247/51	4.0 ±0.55	255/20	7.8 ±0.60	41/2	4.8 ±3.33
2	216/19	8.7 ±1.92	177/21	11.8 ±2.42	222/34	15.3 ±2.42	31/3	9.6 ±5.29	14/1	7.1 ±6.86	-	-
3	58/4	6.9 ±3.32	337/36	10.6 ±1.63	39/5	12.8 ±5.35	152/18	11.8 ±2.61	9/1	11.1 ±10.47	-	-
9	-	-	58/5	8.6 ±3.68	-	-	54/2	3.7 ±2.56	-	-	-	-
13	53/29	54.7 ±6.83	-	-	65/11	16.9 ±4.64	17/0	0.0	-	-	-	-
19	-	-	51/2	3.9 ±4.81	48/2	4.1 ±2.66	107/8	7.4 ±2.5	137/11	8.0 ±2.31	18/0	0.0
	402/52	12.99 ±2.12	664/71	10.69 ±1.15	1003/85	8.47 ±0.71	1608/116	7.21 ±0.65	415/33	7.95 ±1.37	59/2	3.39 ±2.16

Loc.	December		January		February		March		April		May	
	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S
1	82/0	0.0	-	-	93/3	3.2 ±1.82	56/2	3.4 ±2.46	164/5	3.0 ±1.33	487/34	6.9 ±1.14
2	42/1	2.3 ±2.31	-	-	-	-	108/2	1.8 ±1.28	212/11	5.1 ±1.69	-	-
3	-	-	-	-	-	-	-	-	31/7	22.5 ±7.5	29/17	58.6 ±9.14
9	13/1	7.6	-	-	-	-	-	-	103/22	21.3 ±4.03	49/10	20.4 ±5.75
13	30/1	3.3 ±3.26	-	-	-	-	11/1	9.0 ±8.62	23/1	4.3 ±2.05	-	-
19	-	-	-	-	-	-	-	-	27/7	25.9 ±8.43	-	-
	167/3	1.80 ±1.58			93/3	3.23 ±1.82	175/5	2.86 ±1.60	560/53	9.46 ±1.46	565/61	10.80 ±1.90

N, number of collected flies; S, number of sensitive flies; %S, percentage of CO<sub>2</sub>-sensitivity ± standard error.

Table 3. Total number of male and female flies collected at six locations from Mexico City and percentage of CO<sub>2</sub> sensitivity ± standard error.

Loc.	Males		Females	
	Totals	%S	Totals	%S
1	3,694	5.92 ±0.38	3,170	3.92 ±0.33
2	972	12.99 ±1.08	1,022	7.89 ±0.84
3	959	21.09 ±1.32	655	18.62 ±1.52
9	314	17.14 ±2.13	277	12.32 ±1.97
13	206	22.36 ±2.89	199	17.64 ±2.70
19	446	10.51 ±1.45	388	11.82 ±1.64
	6,591	14.99 ±0.44	5,711	12.04 ±0.44

cluding location 19). Such a difference suggests that Y bearing spermatozoa are more effective in transporting the virus than X bearing spermatozoa, or that there exists a differential egg-adult survival depending on sex of infected flies among the progenies.

The differences of sensitivity among adjoining trapping sites may be explained by the relative isolation of the micropopulations, resulting from the low dispersal of *Drosophila*. The dispersal rate is perhaps further diminished in these densely inhabited urban areas, as compared with the field populations previously studied.

A *Drosophila* population which has been decimated during the winter and builds up again from a few new founder individuals will have a vastly different genetic structure than a population at the height of population density. There must be a "bottleneck" in the winter months in Mexico City, as it was impos-

sible to collect any flies during January.

The variations of the percentages of CO<sub>2</sub>-sensitivity in two localities (3 and 13), 2.48 Km. apart are shown in Fig. 2 and 3. The collections included the winter months, when the dwindling of the population took place. The pattern of variation of the proportion of sensitive flies differs considerably in the two trapping sites. Such differences may be due to a drift during the rebuilding of the populations, starting with a small population density, and to the low dispersal rate of urban *Drosophila* flies.

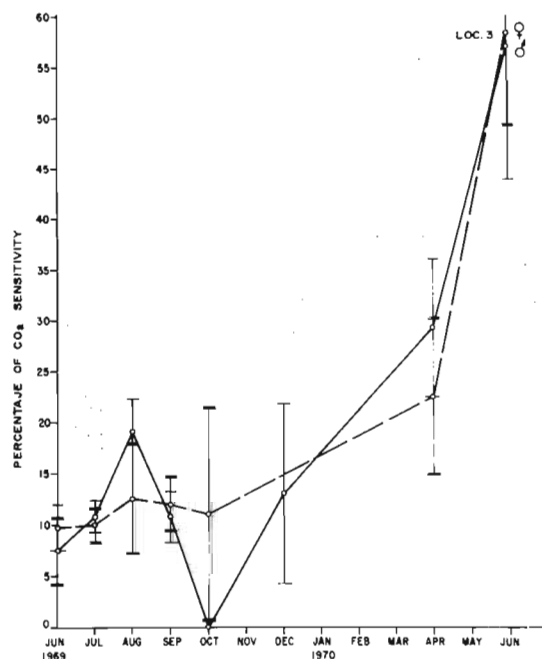


Fig. 2. Variation of CO<sub>2</sub>-sensitivity percentage in location 3.

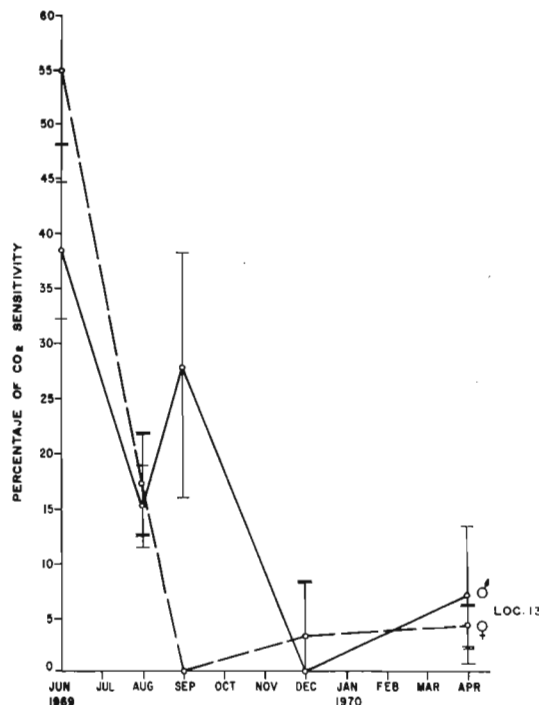


Fig. 3. Variation of CO<sub>2</sub>-sensitivity percentage in location 13.

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Félix R., J. Guzmán and A. de Garay Arellano. Genetics and Radiobiology Program. National Commission of Nuclear Energy. Mexico City, Mexico.  $\text{CO}_2$  sensitivity of *Drosophilid* flies from a location in the outskirts of Mexico City.

(L'Héritier and Teissier, 1945). In most respects the infection is typical of those caused by animal viruses (L'Héritier, 1958; Seecof, 1962), but there are several aspects of the infection which make it noteworthy.

$\text{CO}_2$ -sensitive strains show a maternal effect which assures sensitivity to all the progeny, whereas non-stabilized lines typically throw resistant as well as sensitive progeny (Goldstein, 1949). The non-stabilized hereditary transmission pattern is also displayed by flies that receive sigma virus initially by infection (L'Héritier, 1951).

From June 1969 to May 1970, a survey was made on the distribution of  $\text{CO}_2$ -sensitivity of *D. melanogaster* collected at six locations from the south-west of Mexico City. The bait used for trapping was fermented cantaloupe, and the proportion of sensitive flies was obtained by submitting all the collected samples to the treatment with pure  $\text{CO}_2$  at  $8^\circ$  during 15 minutes.

A first survey of *Drosophilid* species gathered in the same traps was done from October 1969, to March, 1970. Since the flies were trapped incidental to collecting samples of *D. melanogaster*, and since but one collecting technique was employed, the list of species is no doubt incomplete.

The distribution of the collected species; *immigrans*, *hydei*, *busckii*, *pseudoobscura* and *Drosophila* sp. (*repleta* group) shows a scarce dispersion into the trapping sites located in the urban area. It was possible to collect significant, although small, numbers of adults of the five species, only in location 1 (See: Félix, R. et al., 1971). The dominant population at this trapping site was a non-identified species of the *repleta* group. As location 1 is situated in a house in the outskirts of the city, the collected specimens are immigrants from the non-urban area surrounding this place, scarcely inhabited by man, and with an abundant arboreal vegetation (*Cupressus lindleyi* Krotzsch, *Casuarina equisetifolia* L., *Eucalyptus globulus* L. and *Schinus molle* L.)

Among natural populations of *Drosophilids* some individuals are readily found which show a physiological anomaly, sharply outlined and easy to recognize, when brought in contact with carbon dioxide. Sensitive flies will not recover after being anaesthetized with  $\text{CO}_2$  but, rather, will remain paralyzed and eventually die

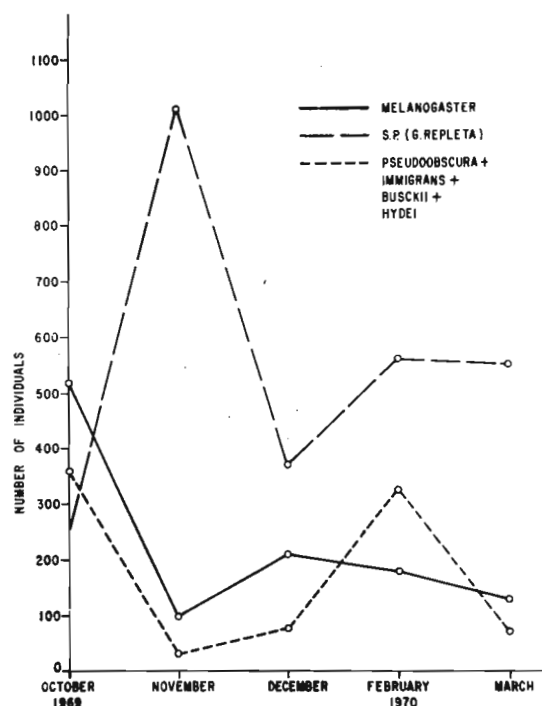


Table 1 shows the relative abundance of each species and its percentage of  $\text{CO}_2$ -sensitivity throughout the winter months. *Drosophilids*, other than *D. melanogaster* do not migrate into urban area, as only about 300 individuals belonging to the five species named above were collected at five trapping sites located in the densely inhabited urban area, at distances from 3.43 to 8.80 Km. apart from location 1 (Fig. 1).

Fig. 1. Relative abundance of *Drosophilid* species collected at location 1.

The data on the sensitivity related to sex, shows that the values for males are consistently larger than that for females (excluding *D. busckii*). The species which showed the largest proportion of  $\text{CO}_2$ -sensitivity is *D. immigrans*. No sensitive flies were found among 145 tested females of *D. pseudoobscura*, which constitutes an interesting feature of this species in a sub-urban area.

Table 2 shows the relative abundance, as compared to *D. melanogaster*, collected at the same traps, of the five species from the locations 1, 9 and 19.



Table 1. Relative abundance and percentage of CO<sub>2</sub> sensitivity of Drosophilid species collected at location 1. N, number collected; S, number of sensitive flies; %S, percentage of sensitive flies  $\pm$  standard error.

Species	October		November		December		February		March		Totals	Mean %S
	N/S	%S	N/S	%S	N/S	%S	N/S	%S	N/S	%S		
<i>D. immigrans</i>												
	110/19	17.3	18/0	0.0	19/0	0.0	2/0	0.0	2/0	0.0	110	17.3
		$\pm 3.60$										$\pm 3.60$
	80/7	8.7	3/0	0.0	12/0	0.0	1/0	0.0	2/0	0.0	80	8.7
		$\pm 3.15$										$\pm 3.15$
	190/27	13.7									190	13.7
		$\pm 2.48$										$\pm 2.48$
<i>D. hydei</i>												
	36/9	25.0	12/2	16.7	11/0	0.0	33/0	0.0	11/1	9.1	69	12.50
		$\pm 7.21$										$\pm 2.97$
	25/1	4.0	5/0	0.0	6/0	0.0	11/0	0.0	4/0	0.0	36	2.00
		$\pm 3.92$										$\pm 1.24$
	61/0	16.4					44/0	0.0			105	8.20
		$\pm 4.73$										$\pm 1.89$
<i>D. melanogaster</i>												
	261/33	12.6	60/1	1.8	126/3	2.3	90/3	3.3	73/2	2.7	610	4.54
		$\pm 2.05$		$\pm 1.33$		$\pm 1.65$		$\pm 1.88$		$\pm 1.87$		$\pm 1.40$
	255/20	7.8	41/2	4.8	82/0	0.0	93/3	3.2	56/2	3.5	527	3.86
		$\pm 0.60$		$\pm 3.33$				$\pm 1.82$		$\pm 2.46$		$\pm 1.79$
	516/53	10.3	101/3	3.0	208/3	1.4	183/6	3.3	129/4	3.1	1,137	4.21
		$\pm 1.57$		$\pm 1.67$		$\pm 0.31$		$\pm 1.21$		$\pm 1.53$		$\pm 0.71$
<i>Drosophila</i> sp. (repleta group)												
	165/20	12.1	849/3	0.4	202/0	0.0	410/17	4.1	427/5	1.2	2,053	3.56
		$\pm 2.59$		$\pm 0.21$				$\pm 0.98$		$\pm 0.50$		$\pm 0.26$
	88/9	10.2	161/1	0.6	174/0	0.0	152/0	0.0	125/0	0.0	700	2.16
		$\pm 3.23$		$\pm 0.61$								$\pm 0.54$
	253/29	11.4	1010/4	0.4	376/0	0.0	562/17	3.0	552/5	0.9	2,753	3.14
		$\pm 3.35$		$\pm 0.20$				$\pm 0.74$		$\pm 0.40$		$\pm 0.11$
<i>D. busckii</i>												
	50/1	2.0	11/0	0.0							61	1.00
		$\pm 2.07$										$\pm 1.27$
	30/2	6.7	12/0	0.0							42	3.35
		$\pm 2.05$										$\pm 2.75$
	80/3	3.7	23/0	0.0							103	1.85
		$\pm 2.11$										$\pm 1.32$
<i>D. pseudoobscura</i>												
	78/2	2.5			19/0	0.0	191/2	1.1	69/1	1.4	357	1.25
		$\pm 1.76$						$\pm 0.71$		$\pm 1.41$		$\pm 0.57$
	17/0	0.0			6/0	0.0	89/0	0.0	33/0	0.0	145	0.00
	95/2	2.1			25/0	0.0	280/2	0.7	102/1	1.0	502	0.95
		$\pm 1.46$						$\pm 0.48$		$\pm 0.93$		$\pm 0.42$

Table 2. Relative abundance of species collected at locations 1, 9 and 19. N, number collected; sp./mel., species/melanogaster.

Species	Loc.	October		November		December		February		March		Totals	sp./mel.
		N	sp./mel.	N	sp./mel.	N	sp./mel.	N	sp./mel.	N	sp./mel.		
Melanogaster	1	516	1.00	101	1.00	208	1.00	183	1.00	129	1.00	1,137	1.00
"	9	34	1.00	0	-	22	1.00	0	-	25	1.00	81	1.00
"	19	238	1.00	64	1.00	0	-	0	-	0	-	302	1.00
Sp. (repleta)	1	253	0.49	1,010	10.00	376	1.81	562	3.07	552	4.28	2,753	2.42
"	9	3	0.09	12	-	5	0.23	0	-	0	-	20	0.25
"	19	0	-	0	-	0	-	0	-	19	-	19	0.06
Pseudoobscura	1	95	0.18	0	-	25	0.12	280	1.53	102	0.79	502	0.44
"	9	0	-	0	-	3	0.14	0	-	22	0.88	25	0.31
"	19	0	-	0	-	0	-	0	-	0	-	0	-
Immigrans	1	190	0.37	21	0.21	31	0.15	3	0.02	5	0.04	250	0.21
"	9	34	1.00	13	-	0	-	0	-	2	0.08	49	0.60
"	19	4	0.02	0	-	0	-	0	-	0	-	4	0.01
Hydei	1	61	0.11	17	0.17	17	0.08	44	0.24	15	0.11	154	0.14
"	9	0	-	0	-	4	0.27	0	-	9	0.36	13	0.16
"	19	0	-	0	-	0	-	0	-	0	-	0	-
Busckii	1	80	15.5	23	-	0	-	0	-	0	-	103	0.09
"	9	44	1.29	0	-	0	-	0	-	0	-	44	0.54
"	19	0	-	0	-	0	-	0	-	0	-	0	-

Acknowledgments: The authors are most grateful to Dr. M.R. Wheeler, for the classification of the Drosophilid species collected during the present survey.

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Gupta, J.P. Banaras Hindu University,  
Varanasi, India. Key to Indian species of  
subgenus Scaptodrosophila.

During last few years taxonomists and  
geneticists in India have reported several  
new and unrecorded species of Drosophila,  
among which seven species belong to the  
subgenus Scaptodrosophila so far. A

taxonomic key is given here to distinguish them with an additional note on their distribution.

1. Mesonotum and scutellum unicolorous.....2  
Mesonotum and scutellum not unicolorous.....3
2. Tarsal segments of male fore legs with many long curved upright hairs  
.....latifshahi Gupta and Ray-Chaudhuri  
Tarsal segments of male fore legs with no such hairs.....4
3. Mesonotum and scutellum with silvery white striations arranged longitudinally  
.....silvalineata Gupta and Ray-Chaudhuri  
Mesonotum and scutellum with scattered silvery white spots arranged longitudinally  
.....chandraprabhiana Gupta and Ray-Chaudhuri
4. Posterior parameres forming a triangular flap-like structure  
.....paratriangulata Gupta and Ray-Chaudhuri  
Posterior parameres not forming a triangular flap-like structure.....5
5. Heel observable and produced into a large spur-like projection  
.....ebonata Parshad and Duggal  
Heel observable but not produced into a spur-like projection.....6

6. Acrostichal hairs in six rows. Or<sub>2</sub> less than half of vibrissa  
 .....bryani Malloch  
 Acrostichal hairs in eight rows. Or<sub>2</sub> not differentiated  
 .....bambuphila Gupta

Species	Source	Locality
D. chandraprabhiana	Bait	Chandraprabha (Chakia forest, Varanasi), Sirsi Dam (Mirzapur)
D. silvalineata	Bait	Chandraprabha (Chakia forest, Varanasi).
D. paratriangulata	Bait	Chandraprabha (Chakia forest, Varanasi); River Bank colony (Lucknow); Ayurvedic garden (B.H.U.).
D. latifshahi	Bait	Chandraprabha, Latifshah (Chakia forest, Varanasi); River bank colony (Lucknow).
D. ebonata	Bait	Srinagar, Pahalgam (Kashmir valley).
D. bryani	Bait and sweeping	Old Botanical garden (B.H.U.)
D. bambuphila	Bait and sweeping	Old Botanical garden (B.H.U.); Jatili near Padmapur (Berhampur).

Franklin, I.R. C.S.I.R.O. Division of Animal Genetics, Epping, N.S.W. Genetic variation at the Esterase-6 locus in *D. melanogaster*.

Wright (1963) in describing the Esterase-6 polymorphism in *D. melanogaster* reported two alleles, Est-6<sup>S</sup> and Est-6<sup>F</sup>. Subsequently Rodino and Martini (DIS 46:139) have reported a third allele, Est-6<sup>V</sup>. In a number of samples of *D. melanogaster*

from the Hunter Valley, N.S.W. four alleles have been observed, and a fifth seen rarely in other collections. Extension of the above notation would result in a cumbersome terminology, and I have followed Hubby and Lewontin (1966) in using the following designation -- Est-6<sup>1.0</sup>, Est-6<sup>1.1</sup>, Est-6<sup>1.15</sup>, and Est-6<sup>1.25</sup>. The first two are equivalent to Est-6<sup>S</sup> and Est-6<sup>F</sup>, and Est-6<sup>V</sup> probably corresponds to Est-6<sup>1.25</sup>. (The fifth allele alluded to above is slower than Est-6<sup>1.0</sup>, and has been represented Est-6<sup>0.95</sup>).

The frequencies of the four most common alleles are quite constant from site to site, and the genotypic frequencies are shown in Table 1.

Table 1. Genotypic frequencies at the Est-6 locus

	Genotype								Total
	1.0/1.0	1.0/1.1	1.0/1.15	1.0/1.25	1.1/1.1	1.1/1.25	1.25/1.25	Others	
Numbers	376	234	7	45	52	7	2	-	714
Frequency	.514	.328	.010	.063	.073	.010	.003	-	

The frequencies of the four alleles are 0.714, 0.242, 0.005, and 0.039.

Some additional data on the location of the Est-6 locus have been collected in test-crosses to 'rucuca'. Using a similar experiment to that described by Wright (1963), 185 flies showing recombination between hairy and thread were tested for their genotype at the Est-6 locus. In 81 cases the recombination had occurred between thread and Est-6. Wright observed 57 out of 149 tested. Pooling these data the location of the Est-6 locus is 3-35.9±0.5.

References: Wright, T.R.F. 1963 Genetics 48:787; Hubby, J.L. and Lewontin, R.C. Genetics 54:577.

Félix, R. and M.E. de la Rosa. Genetics and Radiobiology Program of the National Commission of Nuclear Energy, Mexico. Cytogenetic studies with cyclohexylamine in *D. melanogaster* females.

The artificial sweetener sodium cyclamate, increases chromosome breaks when added in relative high concentrations in human leukocytes in vitro (Stone et al., 1968), as well as in monolayer cultures derived from human skin and carcinoma of the larynx (Stone et al., 1969). The same

compound fails to induce chromosomal damage in *Haworthia variegata* Haw (Majumdar and Lane, 1970). This inability is probably due to the fact that chemical agents that break animal chromosomes may not induce chromosomal aberrations in plants, and different plant genera may react in different ways to the same agent as is found in mammals (Brodie, 1965).

Legator (1968), and Legator et al. (1969) demonstrated that cyclohexylamine, a breakdown product of cyclamate, also induces chromosome breaks in vitro in rodent cells in culture, as well as in vivo, in rat spermatogonia.

The Food Additives and Contaminants Committee provides evidence on the transformation of cyclamates to cyclohexylamine in persons ingesting cyclamates for 2 to 3 days (Food Additives and Contaminants Committee, 1967).

In another line of research a cyclamate-saccharin mixture was fed to male and female rats (FDRL strain, Wistar derived). The doses received in the food varied from 0 to 2,500 mg/Kg/day. Cyclohexylamine was assayed in the urine using the procedure of Derse and Daun (1966), omitting the oxidative steps. The research was carried on, in order to determine whether the conversion of cyclamate to cyclohexylamine takes place in the gastrointestinal tract or systematically as a biotransformation product. In the latter case, the reason for the differences on metabolic handling and the possibility for its genetical control would be of particular interest, considering the similar findings in man (Oser et al., 1968).

Kojima and Ichibagase (1966) and Leahy et al. (1967) also found that cyclamate is metabolized to cyclohexylamine in dogs and man. The most difficult areas in which to determine a cause-and-effect relationship in the human population are in the assessment of carcinogenicity, mutagenicity, or teratogenicity after exposure of the subject to a specific compound. Because of the long latent period between exposure and expression of effects as well as the high background rate of damage, it is difficult to detect effects of a given agent in the population even after years of exposure. Induction of chromosome damage is one of the methods used to evaluate potential carcinogenic, mutagenic, or teratogenic effects of the cyclamate metabolite.

The development of bladder neoplasms was reported in the Wistar strain of rats fed with cyclamate of saccharin (Price et al., 1970). There is no evidence that the use of cyclamate or saccharin has caused cancer in man, malformations in children, or any other abnormality in humans other than a rare skin hypersensitivity. However, in view of the requirements of the Delaney clause of the Food Additives Amendment, the removal of cyclamates from the classification of substances generally recognized as safe resulted in the prohibition of their use in general purpose food products.

The present report contains preliminary results on the sensitivity of oocytes of *D. melanogaster* using the adult feeding method. The process of oogenesis in adult females has already been described in detail (King et al., 1956), and the main features of germ-cell stage sensitivity to some mutagens are well known (Pelecanos, 1964). The chemical treatments administered by adult feeding require a longer period than the irradiation treatments, as the responsible mutagenic reaction is expectedly prolonged after the period of treatment. If newly emerged adult females are treated by adult feeding during 24 hours the stages affected are the stages immediately preceding stage 7, stage 7 itself, and stages 7-13, which have developed during such a period.

The sensitivity of stage 14 oocytes, reached at the 3rd day of adult life can be studied with more confidence, as there is no further development of oogenesis after the treatment.

The procedure adopted for the present study was to collect newly-emerged virgin females which were aged afterwards during 4 days and mating each of them with two males in vials containing regular agar-cornmeal food-medium, making further collection of less than 24 early eggs. The post-treated group includes the isolation of newly-emerged virgin females in regular medium, aging them during 4 days in food with cyclohexylamine, and further mating and oviposition in regular medium (Table 1). In the pre and post-treated group the embryonic, larval and first 4 days of adult life prior to mating took place in a medium with cyclohexylamine (Table 2). Afterwards, the flies were transferred to the regular agar-cornmeal medium employed during all the experiment, and mated in order to collect samples of stage-14

oocytes.

Cyclohexylamine (Merck) was added to the food medium and homogenized with a stirrer (Félix, 1970) at the temperature of  $40 \pm 2^\circ \text{C}$ . The examination of the progenies was done 13 days after oviposition.

In order to test the toxicity of cyclohexylamine, several concentrations were tested, feeding all the stages of development. The concentration of 8.60 mg/ml killed adults before 2 hours, while feeding with concentrations from 4.30 mg/ml to 6.88 mg/ml gave adult survival during 24 hours without development of the eggs layed during such a period. Cyclohexylamine at concentrations of 0.86 mg/ml did not noticeably affect the life-cycle of adult and larvae.

Table 1. Progenies obtained from adult feeding of females with cyclohexylamine.

cyc mg/ml	Control	0.08	0.86
♂ (m.p.c.)	10.40	11.17	10.54
♀ (m.p.c.)	6.40	7.00	7.42
♂/♀ (s.r.)	1.63	1.60	1.42
♂ (total)	260	201	274
♀ (total)	160	226	193
(n.c.)	25	18	26

cyc mg/ml, cyclohexylamine, milligram/milliter;  
(m.p.c.) mean per culture; (s.r.), sex ratio;  
(n.c.), number of cultures.

isolated, the marker  $sc^8$  in the Y chromosome, which contains the normal allele of y identifies XXY females which show gray phenotype instead of the yellow color showed by normal XX females. The marker ebony insures the virginity of such females, as the genotype of the females from the cross  $y^2 w^a/y^2 w^a; e/e \times In(1)EN, Y^S B y \cdot Y^L; +/+$ , is heterozygous for the ebony marker.

Table 2. Progenies obtained from larval and adult feeding of females with cyclohexylamine.

cyc mg/ml	Control	0.08	0.86
♂ (m.p.c.)	10.40	7.57	7.81
♀ (m.p.c.)	6.40	7.57	6.81
♂/♀ (s.r.)	1.63	1.33	1.15
♂ (total)	260	174	125
♀ (total)	160	131	109
(n.c.)	25	23	16

cyc mg/ml, cyclohexylamine, milligram/milliter;  
(m.p.c.), mean per culture; (s.r.), sex ratio;  
(n.c.), number of cultures.

$+/+$  males, and eliminated after the collection of stage 14 oocytes. All the cultures were kept at  $25 \pm 1^\circ \text{C}$  throughout the experiment.

$P \ y^2 w^a/y^2 w^a; e/e \text{ } \text{♀♀} \times In(1)EN, Y^S B y \cdot Y^L; +/+ \text{ } \text{♂♂}$

$F_1 \text{ regular: } y^2 w^a/In(1)EN, Y^S B y \cdot Y^L; e/+(B y/y^2) \text{ } \text{♀♀}$

$y^2 w^a; e/+$

$(y^2 w^a) \text{ } \text{♂♂}$

An improved method for detecting non-disjunction and chromosome X loss was applied, that gives particularly reliable evidence concerning the origin of the exceptional recovered progeny. Females were taken from one stock with  $sc^8 Y$  chromosome, to avoid the existence of any secondary exceptions from XXY mothers ( $y^2 w^a/sc^8 Y; e/e$ ). The tester stock which provides males has an attached  $Y^{SX} \cdot Y^L$  chromosome with the markers yellow and Bar ( $In(1)EN, Y^S B y \cdot Y^L / y^2 su-w^a bb/0$ ). When virgin females are

The fertilization of an XX egg with an XY spermatozoon would produce meta-females with low viability which were excluded from the following analysis, whereas the fertilization of eggs of the same non-disjunctional chromosomal constitution with a non X or Y chromosome bearing spermatozoon, would produce patroclinous yellow, white apricot females of the same genotype as their mothers. These females are easily identified from their normal Bar eyed sisters. The no X egg when fertilized with an XY bearing sperm would become an XY patroclinous male, which can be identified by the Bar eyes in its phenotype.

Virgin females of the genotype  $y^2 w^a/y^2 w^a; e/e$  were aged during 4 days. In each vial an aged female was mated with 2 attached  $Y^S X B y \cdot Y^L$ ;

exceptional:  $y^2 w^a/y^2 w^a;e/+$  ( $y^2 w^a$ ) ♀♀  
 $In(1)EN, y^S B y \cdot y^L;e/+$  ( $B y$ ) ♂♂

Data on the progenies obtained in the treated and control groups are included in Tables 1 and 2. No exceptional progenies were found either among 894 treated flies, from the adult feeding group, nor among 539 treated flies from the larval and adult feeding group.

Among 1,433 stage-14 oocytes treated with the maximum concentrations permissible for the female adult, no exceptions from non-disjunction and X-chromosome loss were recovered. However, the equivalence between the effects of the dosages employed in this experiment, and those from previous findings named above, is evidently difficult. Such factors as absorption, protein binding and excretion are to be considered to make pertinent comparisons with mammalian or human intake cyclamates and its transformation to cyclohexylamine.

It seems from this experiment that *Drosophila* shows a low sensitivity to the ingestion of cyclohexylamine when the genetic events mentioned before are recorded.

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Sreerama Reddy, G. and N.B. Krishnamurthy.  
 University of Mysore, Manasagangotri, Mysore-6, India. Preliminary survey of *Drosophilids* in Nilgiris and Kodaikanal ranges.

Maiden trips were made to Nilgiris and Kodaikanal ranges to explore the *Drosophilids*. Nilgiris is about 100 miles to the south of Mysore and Kodaikanal is about 300 miles also to the south of Mysore. Both these localities are hill stations

characterised by the combination of temperate and tropical forests. The sholas are ever-green with moist and humid atmosphere in most part of the year. Shifting cultivation forms the common biotic factor in both the localities. The annual rain fall ranges from 1524 to 2540 mm. The highest peak of Nilgiris is 2580 meters and that of Kodaikanal is 1900 meters.

Collections were made in the beginning of December 1970. In the Nilgiris, flies were trapped at 11 different altitudes ranging from 840 to 2580 meters, whereas in Kodaikanal the flies were trapped at six altitudes ranging from 1000 to 1900 meters. A total of 1470 flies were collected from both the localities, of which 1009 flies come from Nilgiris and 461 from Kodaikanal. The details of the collection record are depicted in tables 1 and 2. The various species collected from both the localities are *D. melanogaster*, *D. ananassae*, *D. melarkotliana*, *D. takahashi*, and *D. immigrans*. The species like *D. repleta*, *D. kikkawai*, *D. mysorensis*, *D. hoozani* like species and *D. busckii* are found in Nilgiris and absent in Kodaikanal range. In addition, two new species of *Drosophila* which will be described elsewhere were also found in the traps. In Kodaikanal range one individual belonging to the Genus *Leucophenga* was found in an orchard near Valegiri at an altitude of 1000 meters. It is quite remarkable to note that *D. immigrans* is wide spread in its distribution in almost all places scanned. This shows that *D. immigrans* thrives well in a moist and humid climate. Further the most interesting feature of our collection study is the lack of *Drosophila* flies

in the traps except one individual male of *Drosophila hoozani* like fly at a high altitude of 2580 meters in Doddabetta (Nilgiris).

Table 1. Distribution of different species of Drosophilids in Nilgiri Range

	Wild Domestic	Altitude in Metres	<i>D. melanogaster</i>	<i>D. ananassae</i>	<i>D. melarkotliana</i>	<i>D. repleta</i>	<i>D. takahashi</i>	<i>D. kikkawai</i>	<i>D. mysorensis</i>	<i>D. immigrans</i>	<i>D. busckii</i>	<i>D. species like hoozani</i>	Prob. new species (near truncata)A3	Prob. new species A2	Total
Thippakadu	Wild	840	-	-	5	-	35	-	-	-	-	-	-	-	40
Gudalur	Domestic	900	42	331	-	32	-	-	-	-	-	-	-	-	405
Gudalur	Wild	990	-	-	14	-	3	5	2	20	-	-	-	-	44
Merupalyam	Wild	1200	-	25	-	-	8	-	-	45	-	-	8	-	86
A virgin forest on the Ooty Road	Wild	1310	-	-	-	-	-	-	31	31	-	-	-	-	62
Coonur	Domestic	1800	25	-	-	-	5	9	-	15	-	-	-	10	64
Naduvattam	Domestic	1840	3	2	2	10	2	-	-	65	3	-	-	-	87
Naduvattam	Wild	1850	-	-	-	-	-	-	10	10	-	-	-	-	20
T.R. Bazar	Domestic	1970	5	-	-	90	-	-	-	10	-	-	-	-	105
Ooty city	Domestic	00	6	-	-	-	-	33	-	56	-	-	-	-	95
Doddabetta	Wild	580	-	-	-	-	-	-	-	-	-	1	-	-	01
			81	358	21	132	53	47	43	252	3	1	8	10	1009

Table 2. Distribution of Different Species of Drosophilids in Kodaikanal Range

	Wild or Domestic	Altitude in Metres	<i>D. melanogaster</i>	<i>D. ananassae</i>	<i>D. melarkotliana</i>	<i>D. takahashi</i>	<i>D. immigrans</i>	<i>Leucophenga</i> ( <i>Trichiasiphenga</i> ) <i>invicta</i> (Walker)	Total
Valegeri	Orchard	1000	-	-	4	2	65	-	72
Oothu	Domestic	1100	-	165	9	-	0	-	194
Oothu	Wild	1400	-	-	3	2	40	-	45
Perumalmalai	Domestic	1580	38	30	-	-	10	-	78
Kodaikanal	Wild	1750	-	-	9	5	18	-	32
Kodaikanal Town	Domestic	1900	-	-	-	-	40	-	40
			38	195	25	9	193	1	461

Acknowledgements: We are highly grateful to Dr. M.R. Rajasekarasetty, Professor and Head of the Department of Zoology, Manasagangotri, University of Mysore, Mysore for his help and encouragement. We are indebted to Dr. T. Okada, Professor of Biology, Tokyo Metropolitan University, Department of Biology, Setagaya-Ku, Tokyo, Japan for identifying the species. One of us (G. Sreerama Reddy) is thankful to the University of Mysore and University grants commission for the financial support.

Ringo, J.M. University of California, Davis, California. The effects of anesthetization upon survival and behavior of *D. grimshawi*.

To study some elements of behavior of *D. grimshawi* Oldenberg, it is necessary to mark and mutilate individuals, which in turn requires that the flies be lightly anesthetized. In order to find an anes-

thetization procedure which interferes the least with subsequent behavior, the effects of three agents ( $\text{CO}_2$ , cold, and ether) were evaluated.

To assess the effects of treatment upon survival, one control group ( $n=30$ ) was maintained and each agent was applied to five other groups of flies ( $N=20$  in each group) for different lengths of time (1/2, 1, 2, 4, and 8 mins.) Flies were chosen at random from a population of PK9 *D. grimshawi* adults, aged 1 to 22 days.  $\text{CO}_2$  was administered by suspending flies in a plastic tube over dry ice in a one lb. coffee can. The tube was fitted through a hole in a cardboard top, and the bottom of the tube was covered by a piece of bolting silk. This apparatus was modified from Seecof (1963). The temperature at the bottom of the tube was approximately  $4^\circ\text{C}$ . Anesthetization with cold was attained by placing flies in an aluminum cigar tube immersed in ice; the temperature was about  $0^\circ\text{C}$ . Ether was used in an ordinary small plastic etherizer. About two ml of ether was placed on the gauze at the bottom of the etherizer, and a few drops were added between treatment groups. The temperature was approximately  $20^\circ\text{C}$ . After treatment, each group was placed in a half pint bottle containing fresh food and maintained at  $20^\circ\text{C} \pm 1^\circ$ . Dead flies were removed and counted at 24, 48, 72, and 96 hours after treatment. The results are summarized in the following table and graphs:

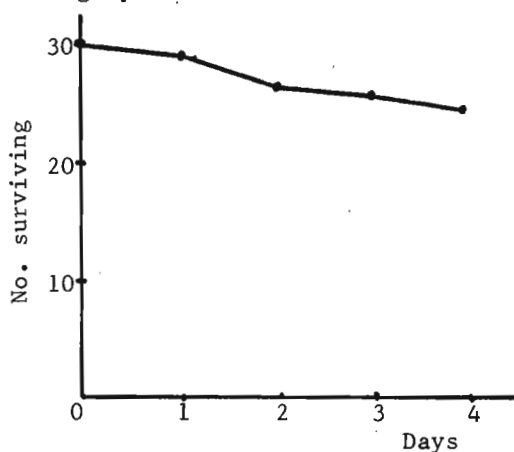


Figure 1.

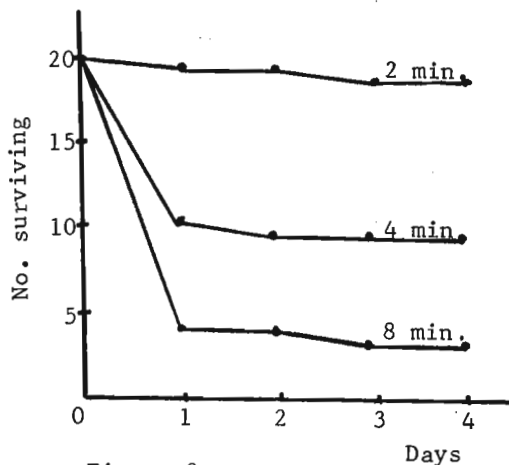


Figure 2.

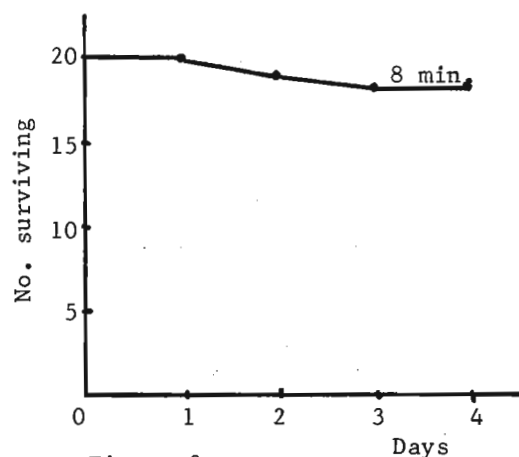


Figure 3.

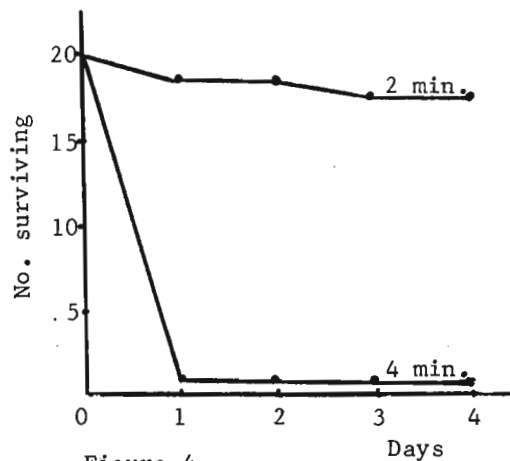


Figure 4.

Survival curves for six treated groups and control group. Fig. 1, controls; fig. 2,  $\text{CO}_2$  treatment for 2, 4, and 8 mins.; fig. 3, cold treatment for 8 min.; fig. 4, ether treatment for 2 and 4 mins.



Table 1

No. surviving	Min. with CO <sub>2</sub>						Min. with cold						Min. with ether						Controls
	1/2	1	2	4	8		1/2	1	2	4	8		1/2	1	2	4	8		
after 24 hours	19	19	19	10	4		20	18	19	20	20		19	20	18	1	0		29
" 48 "	19	19	19	9	4		19	14	19	18	19		18	19	18	1	0		26
" 72 "	18	18	18	9	3		17	14	18	18	18		18	18	17	1	0		25
" 96 "	18	18	18	9	3		17	14	18	18	18		18	18	17	1	0		24

We accept the hypothesis that the proportion of survivors among the controls and all flies treated for 30 sec. were equal ( $\chi^2=1.43$ ,  $df=2$ ,  $p>.20$ )

A second experiment sought to determine differences in behavior attributable to these three methods of anesthetization. The phenotype of greatest interest is jousting, a type of behavior found only in males of this species. Subjects were drawn at random from a population of adult PK9 males aged 19 to 25 days.  $N=30$  for each treatment group. Ss were anesthetized for 30 sec., their wings were marked with nail polish containing non-toxic dyes; they were isolated in individual half pint bottles containing fresh food and were maintained at  $20^\circ\text{C}\pm 1^\circ$ . Allowing at least two hours for recovery, Ss were observed in batches ( $N=10$ ) in plexiglass cells ( $2\times 5\times 9$  cm) with moist sponge at one end. Their interactions were observed for 20 minutes and recorded; the exact time spent jousting was recorded for each subject using an Esterline Angus 10-channel event recorder. The observations were repeated four more times for each S.

There were marked behavioral differences between treatments. Aggression and courting were very much reduced in cold-treated Ss, and somewhat reduced in CO<sub>2</sub>-treated Ss relative to etherized Ss. The quantitative results for jousting show a similar pattern:

Table 2

Treatment	Total of all scores	No. of Ss
CO <sub>2</sub>	834.3	28
cold	932.9	25
ether	1842.1	30

The data can be analyzed in two ways. One can simply record whether or not a subject jousted during a given observation period, or one can consider the relative amount of jousting for each test period. An ordinary analysis of variance is impossible, since the scores have a J-shaped distribution. Out of 415 observations (7 Ss died) or scores,

271 were zero. Using  $271/415 = .653$  as the expected proportion of zero scores among treatments and testing  $H_0: \theta_1 = \theta_2 = \theta_3$  against the alternative that the proportions are not equal, we reject  $H_0$  ( $\chi^2=10.37$ ,  $df=2$ , and  $p<.01$ ). The large number of zero scores in all groups of Ss indicates that a simple dichotomous measure has as much biological significance as the amount of time spent jousting. The simplest non-parametric test using the scores is the Friedman two-way analysis of variance by ranks (Siegel 1956). The Friedman test requires equal sample sizes, but 7 Ss died during the experiment and could not be replaced so we averaged the scores for each batch. We reject the hypothesis that treatments do not differ in their effects ( $\chi^2=6.50$ ,  $df=2$ ,  $p<.05$ ).

The results of these experiments show that light etherization is a better method of anesthetization for behavioral studies in *D. grimshawi* than the use of either CO<sub>2</sub> or low temperature.

References: Seecof, R.L. 1963 DIS 37:145; Siegel, S. 1956 Nonparametric Statistics for the Behavioral Sciences, McGraw-Hill, Inc., New York.

Hunt, D.M. University College London, England. A haemolymph protein anomaly associated with the lethal-giant-larvae mutant in *Drosophila melanogaster*.

Faulhaber (1959) demonstrated a reduction in the haemolymph protein content of larvae homozygous for the *lgl* mutant. However, the paper electrophoresis technique employed by Faulhaber allowed the clear separation of only two protein fractions. With the introduction of acrylamide gel as a supporting medium for electrophoresis, it is now possible to

identify a large number of protein fractions in larval haemolymph. A re-examination of the situation in *lgl* therefore would seem appropriate.

Two alleles *lgl* and *lgl<sup>B</sup>* have been studied. Both mutants were maintained as balanced lethals over the SM5 chromosome. Haemolymph samples from non-lethal larvae were collected at about 5 days post-hatching when the larvae leave the food medium prior to pupation. Development in lethal larvae is delayed and haemolymph samples were taken therefore from at least 6 days post-hatching. The technique of acrylamide gel disc electrophoresis was used. The procedure follows Davis (1964) except that the spacer and sample gels were omitted and 50  $\mu$ l of sample applied directly to the top of each gel. Gels were stained in 0.5% amido black in 7% acetic acid.

No differences could be detected in the haemolymph proteins from *lgl/lgl* and *lgl/SM5* third instar larvae, and *lgl<sup>B</sup>/SM5* larvae also gave a normal protein pattern. However, the electrophoretic separation of haemolymph samples from *lgl<sup>B</sup>/lgl<sup>B</sup>* larvae revealed clear differences in protein content; fraction 7 was entirely missing and the amount of stainable material in fractions 12 and 15 considerably elevated (Fig. 1).

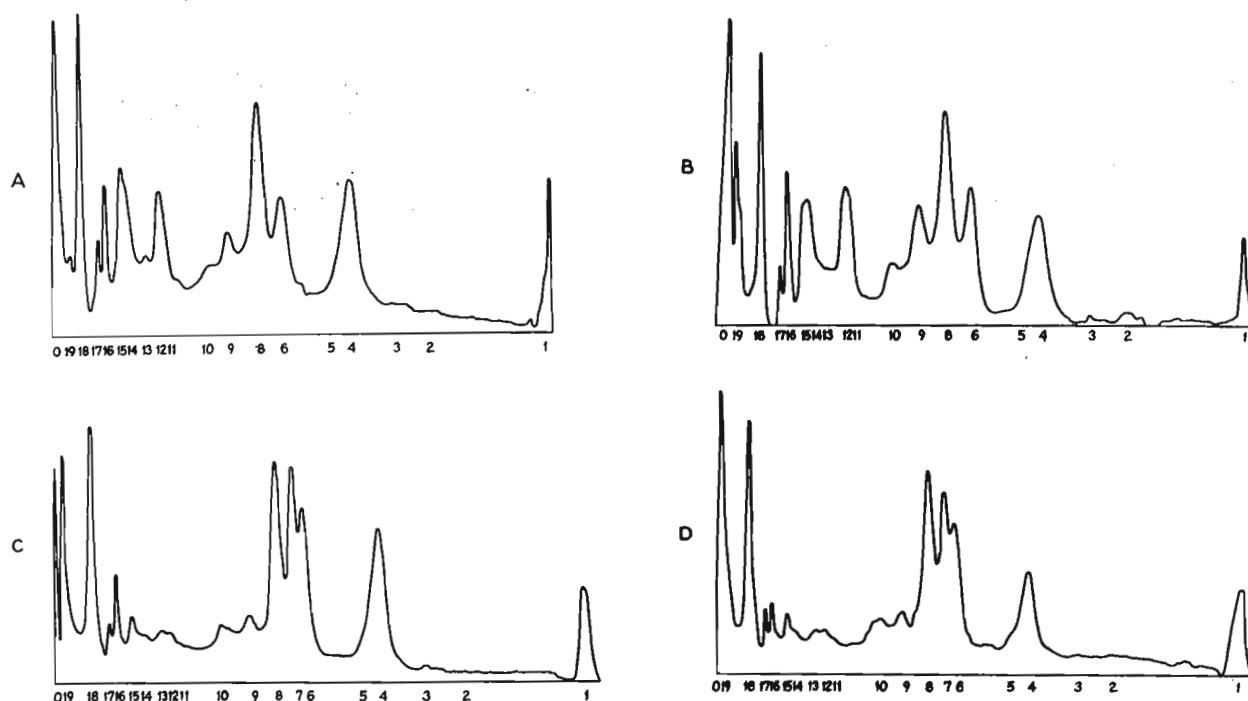


FIGURE 1. Densitometer tracings of electropherograms obtained from haemolymph samples of third instar *lgl<sup>B</sup>/lgl<sup>B</sup>* larvae (A,B) and *lgl<sup>B</sup>/SM5* larvae (C,D). The protein fractions are numbered from the running front to the origin (0).

Fractions 12 and 15 show considerable quantitative variation between strains so the inheritance of the elevated quantities in *lgl<sup>B</sup>* homozygotes was not examined further. The possibility that the absence of fraction 7 in lethal larvae depends on another gene locus on the *lgl<sup>B</sup>* second chromosome was tested by outcrossing the *lgl<sup>B</sup>/SM5* strain to the Edinburgh wild type. Non-SM5 *F<sub>1</sub>* progeny was mated to expose the *lgl<sup>B</sup>* chromosome to recombination and haemolymph samples were taken from the resulting third instar larvae. In a total of 104 *lgl<sup>B</sup>/lgl<sup>B</sup>* larvae examined, no recombinants were obtained.

References: Davis, B.J., 1964, *Ann. N.Y. Acad. Sci.* 121: 404; Faulhaber, I., 1959, *Z. Induktive Abstammungs-Vererbungslehre* 90: 299.

Nirmala Sajjan, S., and N.B. Krishnamurthy,  
University of Mysore, Mysore-6, India.  
Karyotype of *Drosophila nasuta*.

*D. nasuta* subgroup of the immigrans group created by Wilson et al (1969) includes 8 morphologically similar species. Males of this subgroup have silvery markings on the frons, in all but one species.

However, whitish to silvery sheen over the entire frons is present only in *D. albomicans*, *D. kohkoa*, *D. kepulauana* and *D. nasuta* Lamb (1914). The species described here is characterized by silvery sheen over the entire frons but differs cytologically from that of *D. albomicans*, *D. kohkoa* and *D. kepulauana*. Though Ray Chaudhuri and Jha (1969) have given an account on the cytology of *D. nasuta*, it is not known to which species proper it belongs under *nasuta* subgroup. The karyotype of *D. nasuta sensu strictu* is yet unknown. Hence the karyotype is reported here.

The flies collected from Soundatti (Mysore state) are big and yellowish in color with silvery frons. There is brown longitudinal streak on pleura reaching back to the wing base in both males and females. Other morphological characters are similar with that of *D. nasuta* reported by Okada (1964). The metaphase karyotype of the larval neuroblast cells (Fig. 1) consists of a pair of rods which represents X chromosome in females, one of which is replaced by V-shaped Y chromosome in males, a pair of V's (chromosome 2), a pair of double length rods (chromosome 3) and a pair of dots (chromosome 4). No additional heterochromatin is found in the dot.

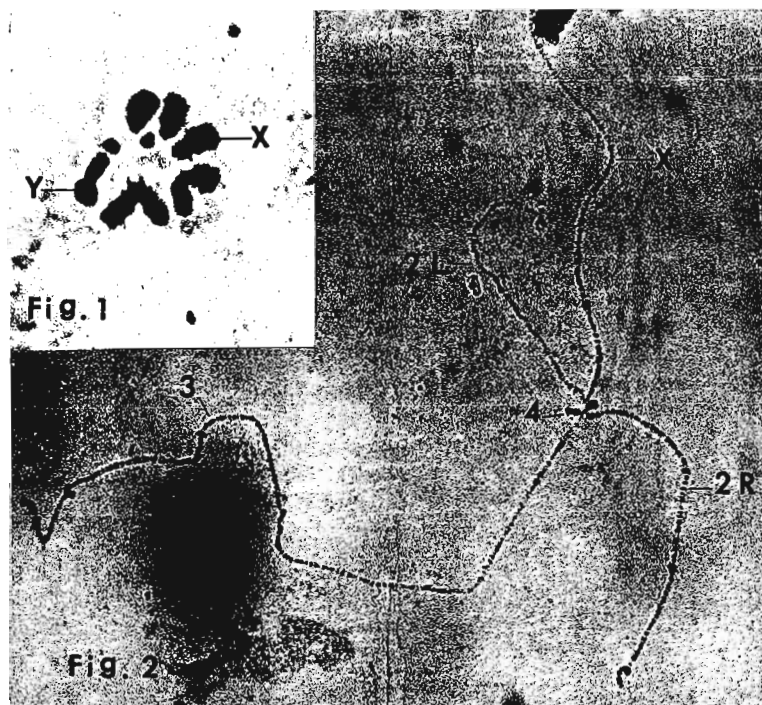


Fig. 1. The Metaphase Karyotype of male larval neuroblast cell.

Fig. 2. Salivary gland chromosomes

The salivary gland chromosomes show four long arms and one small arm as shown in the fig. 2. Centric heterochromatin is practically absent except for a little between 2L and 2R. Like other species of the subgroup, here also a loop, which is not an inversion is frequently observed in the basal region of 2R. The longest arm represents the double length rod of the metaphase karyotype and the arm next to third chromosome in length is the X chromosome and the remaining two long arms are the left and right arms of the metacentric second chromosome. The small arm represents the dot chromosome of the somatic metaphase.

The species under study is allied to *D. albomicans*, *D. kepulauana* and *D. kohkoa* in having entire frons with silvery sheen. This is also true with the original species, *D. nasuta* according to published notes for which cytology is not known. Based on the cytological analysis, the species here described differs from *D. albomicans* in that *D. albomicans*

has  $2n=6$ , whereas here we have  $2n=8$ . It also departs from *D. kepulauanana* in having V-shaped Y chromosome and basic type of dots while in *D. kepulauanana* Y is rod shaped and the dot chromosomes with added heterochromatin are slightly thicker and longer. The other member of the same series with entire silvery frons -- *D. kohkoa*, is characterized by the pinched constriction in the third chromosome which is always accompanied by the dot. This species also has a small amount of added heterochromatin to the dot which gives it a comma-shaped appearance (Wilson et al, 1969). This has not been observed in the present species. The karyotype described by Ray Chaudhuri and Jha (1969) consists of 6 pairs of chromosomes in metaphase configuration and 6 arms (5 long and one short arm) in salivary gland nuclei. Our findings are different from this.

Recounting the similarities and differences that are exhibited by the members of the *nasuta* subgroup, the species herein described must be either *D. nasuta* sensu strictu or a new species of the *nasuta* subgroup for which confirmation is needed. Further this species is highly polymorphic in having duplications and deficiencies and a multitude of inversions which will be presented elsewhere.

Acknowledgements: The authors are very grateful to Dr. M.R. Rajasekarasetty, Professor and Head of the Department of Zoology, University of Mysore, Manasagangotri, Mysore for his advice and encouragement. We are thankful to Mr. Ramakrishna Raju for preparing Photomicrographs. This work is supported by the department of Atomic Energy, Government of India.

References: Ray Chaudhuri, S.P. and A.P. Jha. 1969, *The Nucleus*, Vol. XII(1): 9-13; Wilson, F.D., M.R. Wheeler, Margaret Harget and Michael Kambyssellis. 1969, Cytogenetic relations in the *D. nasuta* subgroup of the *immigrans* group of species.

Sanjeeva Rao, M. and S. U. Devi. Osmania University, Hyderabad-7, AP., India. Induction of mutations in *D. melanogaster* with radioisotopes -  $^{90}\text{Sr}$  and  $^{131}\text{I}$ .

Even though much work was done on the induction of mutations in *Drosophila* by ionizing radiations and chemicals, the possible mutagenic effects of radioisotopes have received little attention. Blumel (1950) reported that phosphorus-32 induces muta-

tions in *Drosophila* while Rubin (1950) observed mutagenicity in microorganisms.  $\text{Sr}^{90}$  and  $\text{I}^{131}$  are more powerful radioisotopes than phosphorus-32 and to assess their genetic damage in *Drosophila* the following experiments were carried out.

Two concentrations of each isotope were tried. The isotope was mixed in food medium. Flies were allowed to lay eggs on this medium and the offspring were allowed to grow on the medium containing the isotope. The treated males were crossed individually with 3 virgin females of  $y\ sc^{S1}\ In-49\ sc^8; bw; st$  for three days only to assess the genetic damage in spermatozoa alone. The  $F_1$  females were mated individually with  $y\ sc^{S1}\ In-49\ sc^8$  males while the males were mated with  $bw; st$  females to score for sex linked recessive lethals and translocations, respective in the  $F_2$  generation. The results are presented in Table 1.

Table 1

Treatment	Sex linked recessive lethals				Translocations			
	T	l	%	Chi-square value	T	l	%	Chi-square value
1. Control	505	1	0.2	-	712	-	-	-
2. $\text{Sr}^{90}$ 0.2 $\mu\text{cc}$								
in 100cc of food	329	8	2.12	9.3	439	3	0.68	4.94
3. $\text{Sr}^{90}$ 1.0 $\mu\text{cc}$								
in 100cc of food	268	5	1.86	6.33	247	3	1.21	8.74
4. $\text{I}^{131}$ 1.00 $\mu\text{cc}$								
in 100cc of food	436	8	1.83	6.64	-	-	-	-
5. $\text{I}^{131}$ 2.00 $\mu\text{cc}$								
in 100cc of food	363	5	1.40	4.28	347	2	0.6	4.2

T = Total number of X chromosomes or  $F_1$  sons scored; l = Lethals recorded;

t = translocations recorded

These preliminary studies indicate that  $^{90}\text{Sr}$  and  $^{131}\text{I}$  cause mutations in *D. melanogaster* similar to phosphorus - 32.

Kastritsis, C.D.<sup>1</sup> and J. Grossfield<sup>2</sup>. <sup>1</sup>University of Texas Southwestern Medical School, Dallas, Texas. <sup>2</sup>Purdue University, Lafayette, Indiana. Balbiani rings in *D. auraria*.

Since different strains of *D. auraria* differ with respect to their ability to mate in darkness, and since this trait is at least under partial genetic control (Grossfield, 1970), an investigation was undertaken to explore possible cytological correlations.

Different strains were found to differ by a number of inversions and, in addition, two strains were found to exhibit two Balbiani rings (Fig. 1) in one of the chromosomes of the

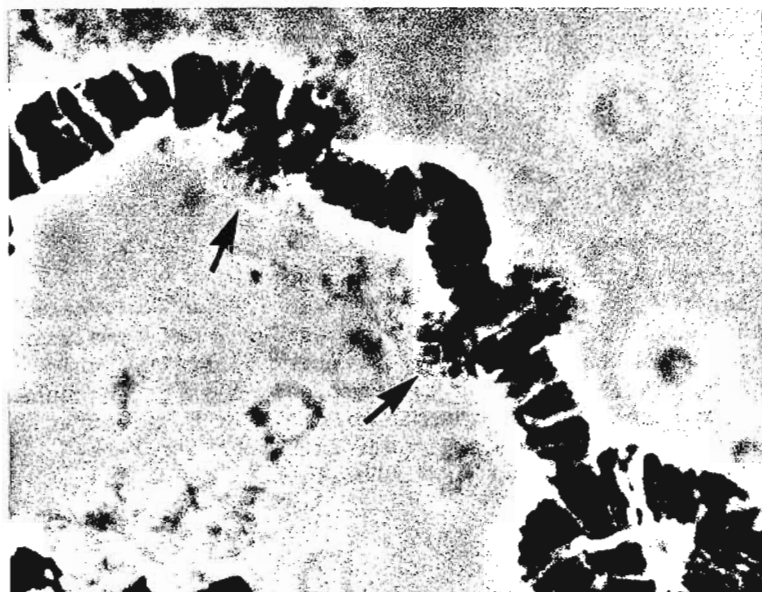


Fig. 1. Phase contrast photomicrograph of *D. auraria* polytene chromosome. Arrows point at two Balbiani rings.

salivary glands cells. Due to the fact that Balbiani rings have not been described in *Drosophila* before, we feel that our observation warrants this note. Our preliminary data indicate that these may be stage-specific structures. Research is now under way to further investigate the implications of the phenomenon.

Reference: Grossfield, J., 1970. Genetics 65:s27.

Würgler, F.E., R. Büchi and P. Maier. Swiss Federal Institute of Technology, Zürich, Switzerland. Relative viability of different types of *Drosophila melanogaster* males without a free Y chromosome.

If  $R(1)2,y B / B^S Y y^+$  males are X-rayed and mated to nonirradiated females partial as well as complete loss of sex chromosomes is indicated by non-Bar males. The reports of Graf and Würgler (DIS-46, 73-74, 1971) and Würgler and Kälin (DIS-46, 79-80, 1971) show that the rates of chromosome

losses recorded depend on the type of females used in the test crosses. The data obtained for X-irradiation of ring-X males with 2000 R in nitrogen are summarized in the following table ("Oster" = Inscy;dp bw;st pP and "XY" =  $y^2 su(w^a)w^a KS.KL y^+$  (Parker 110-8)):

females	spontaneous rate of loss	X-ray experiment	corrected for spontaneous loss	relative rate
Oster	0.54 %	2.33 %	1.8 %	1
y sn	0.71	4.37	3.7	2.1
XY	2.2	8.7	6.7	3.7

In the three tests males of different genotype are indicative for a sex chromosome loss. The difference found between stocks might therefore simply reflect the relative viability of the non-Bar males. To test this possibility the relative viability of y sn /0 and XY/0 males compared to Oster/0 males was determined. For this purpose hybrid females y sn /Oster and XY/Oster were mated to XY/0 males and the progeny classified according to the genotype. This is a more rigorous test than the one described by Graf and Würgler because the different types of males to be compared develop under identical conditions within the same vial. The data obtained are given in the following table. For comparison, results from some other comparable test crosses are also included: (All hybrid females are heterozygous for dp bw st pP.)

females (X*/Oster)	total progeny	X*/0 males	Oster/0 males	male ratio
y sn /Oster	6117	1945	1627	1.20
XY / Oster	5871	1790	1374	1.30
+/Oster ; t/vg	6561	2194	1522	1.44
Oregon-R/Oster	5175	1680	1136	1.47
Hikone-R/Oster	10043	3090	1836	1.68
Berlin wild/Oster	3370	1081	628	1.72

For the y sn stock we find a relative viability of the y sn/0 males compared to Oster /0 males of 1.2. The corresponding ratio for the induced chromosome losses in the X-ray experiments is 2.1. A similar result is found for the XY stock : 1.30 versus 3.7. This analysis shows that, although the various types of males show a slightly different viability, this difference is not sufficient to explain the variation of the chromosome loss rates encountered in the X-ray experiments. It is postulated that after insemination of the nonirradiated egg some factors which are under the control of the maternal genome influence the X-ray lesions induced in mature sperms. Work supported by Schweizerischer Nationalfonds zur Förderung der wissenschaftlichen Forschung.

Ondřej M. Prague, Czechoslovakia. The induction of large chromosomal fragments by ethylnitrosourea and radiation.

Ethylnitrosourea is one of the most effective mutagens in producing recessive lethals. It is of interest to know the relative effectiveness of ethylnitrosourea

(ENH) in producing large chromosomal aberrations. The relationship of large chromosomal fragments to lethals in the X chromosome after ENH treatment was compared with that after X-radiation. ENH was applied by injections; each fly received, on the average, 0.2 µl of solution. X-radiation was applied by means of Siemens apparatus with those parameters: 22 mA, 200 kV, OK - 17.5, filter 0.5 mm Cu, dose - rate 394.7 R/min., overall dose 1500 r. The concentration of ENH was near the upper limit of applicable concentrations. The radiation dose was of medium magnitude, as regards the induction of sterility. The results show, that while ENH induces extremely high frequency of recessive lethals it is very weak chromosome breaker. The ratio of the frequencies of large fragments to recessive lethals after ENH treatment was in our experimental conditions of about two orders of magnitude lower than that after irradiation.

The frequencies of large fragments and recessive lethals in the X chromosome induced by ENH and x-rays.

Treatment	Fragments		Rec. lethals		Fragments/ lethals
	No. of F <sub>1</sub> females	% of fragments	No. of X chromosomes	% of rec. lethals	
ENH 10 mM	4,770	0.02	550	41.5	0.0005
X-rays 1500 R	18,577	0.27	715	5.6	0.048
Control	19,570	0.005			

Jacob, M. and S.P. Ray-Chaudhuri, Banaras Hindu University, Varanasi, India. Protective effect of glutathione (reduced) against X-ray induced sex-linked recessive lethals in *D. melanogaster*.

The thiol tripeptide, glutathione, was shown to be protective against radiation-induced lethality in mice (Chapman et al. 1950). This chemical yielded significant protection against chromosomal aberrations also, in *Tradescantia* (Mikaelsen 1952) and grasshopper (Chaudhuri 1965). But Mittler (1964) reported the failure

of glutathione as well as a few other well-known chemicals to protect *Drosophila* chromosomes against X-ray induced sex-linked lethals, dominant lethals, translocations and deletions.

A number of experiments were conducted to test the protective effect of reduced glutathione against sex-linked recessive lethals using Oster's stock. One-day old males ( $X^{c2}yB$ )

Frequency of sex-linked recessive lethals after 2000 r and pretreatment with glutathione(reduced)=GSH

Experiment No.	Treatment	No. of sex linked lethals	No. of chromosomes tested	% of lethals	$\chi^2$	Probability	Degree of protection
1	GSH + 2000r	41	770	5.3	4.1	$0.05 > P > 0.025$	35.3
	Saline+2000r	50	604	8.2			
2	GSH + 2000r	48	923	5.2	2.6	$0.25 > P > 0.10$	24.6
	Saline+2000r	76	1096	6.9			
3	GSH + 2000r	51	932	5.4	7.1	$0.01 > P > 0.005$	36.6
	Saline+2000r	147	1722	8.5			
4	GSH + 2000r	134	2729	4.9	5.3	$0.02 > P > 0.01$	28.9
	Saline+2000r	72	1037	6.9			
Total	GSH + 2000r	274	5354	5.1	24.9	$P > 0.005$	33.7
	Saline+2000r	345	4456	7.7			

were pretreated with the chemical dissolved in saline to a concentration of 10 mM and each fly received 0.6  $\mu$ l of the solution. The controls were treated with saline and both the lots received a dose of 2000 r X-ray after 30 minutes of the treatment. Crosses were made with these males and the virgins of the Oster  $\phi$  stock. Nearly twenty pair-matings were made from the  $F_1$  progeny of each treated male and the  $F_2$  offspring were examined for the absence of Bar males, indicating the presence of a lethal induced in the paternal X-chromosome.

The consistent results of the four repeated experiments are presented in the table. Significant protection was observed in all the experiments, only one being slightly below the border line. The pooled data of all the experiments show a highly significant protection,  $P(\chi^2) 0.005$ , the degree of protection being 33.7.

References: Chapman, W.H., C.R. Sipe, D.C. Eltzholtz, E.P. Cronkite and F.W. Chambers 1950, Radiology 55: 865; Chaudhuri, J.P., 1965, Ph.D. Thesis, Banaras Hindu University; Mikaelsen, K., 1952, Science 116 (3007): 172-174; Mittler, S., 1964, Int. J. Rad. Biol. 6 (5): 405-413.

Lee, T.J. Chungang University, Seoul, Korea. Frequency of races in males of *D. auraria* in natural populations.

*D. auraria* is a polymorphic species in Korea populations. This species was divided into three races, A, B and C, mainly by forms of genitalia. In natural populations of Korea, the distribution

area of races A and C is much wider than race B. In general, race A is the most domestic of the three, abundant around areas of human habitation, while race B inhabits rather cool, and mountainous or relatively northern regions. The habitat of race C is, in general, wider in environmental and higher altitude than that of race A. However, it is sometimes found that the two or three races live sympathetically.

Among the flies collected in twelve localities, a few flies of *D. auraria* showed a

hybrid character in male phallic organs. The shapes in the phallic organs found in these males seems to be the same as those found in experimental hybrids obtained in the laboratory. However, no natural AB hybrid has yet been detected.

Races		A	B	C	AC	BC	Total
Is. Quelpart	No.	11		58	2		71
	%	15.4		81.6	2.8		
Mt. Chiri	No.	41		35	6		82
	%	50.0		42.6	7.3		
Mt. Kaya	No.	33	13	41	4	2	93
	%	35.4	13.9	44.	4.3	2.2	
Muju	No.	75		13	2		90
	%	83.3		14.4	2.2		
Mt. Palkong	No.	15		48	7		70
	%	21.4		68.5	10.0		
Mt. Kyeryong	No.	58		25	5		88
	%	65.9		28.4	5.6		
Kongju	No.	75		7	3		85
	%	88.2		8.2	3.5		
Mt. Sokli	No.	75	20	26	3	2	126
	%	59.5	15.9	20.6	2.3	1.6	
Daekwanryong	No.	19	30	53	4	2	108
	%	17.5	27.7	49.0	3.7	1.8	
Kwangneung	No.	40	32	22	3	1	98
	%	40.8	32.6	22.2	3.1	1.0	
Mt. Soyo	No.	17		22	7		46
	%	36.9		47.8	15.		
Mt. Sulak	No.	20	39	25	2	2	88
	%	22.7	44.3 c	28.4	2.7	2.7	

The sexual isolating mechanisms among the three races were analyzed. The mean coefficient of interracial sexual isolation in the three classes are arranged in descending order,  $A \leftrightarrow B$  (0.845) >  $B \leftrightarrow C$  (0.600) >  $A \leftrightarrow C$  (0.426) at 25°C, and  $A \leftrightarrow B$  (0.788) >  $B \leftrightarrow C$  (0.417) >  $A \leftrightarrow C$  (0.165) at 19°C.

In crosses between A and B, a highly significant sexual isolation was demonstrated at both temperatures of 19°C and 25°C.

It is conjectured that races A and B, in natural populations, have been completely precluded genetically from one another by many isolating mechanisms.

References: Kurokawa, H. 1960. Japan J. Genet., 35:161-166; Kurokawa, H. 1967. Annot. Zool. Japan., 40:154-160; Lee, T.J. 1970. Chungang Univ. Theses Collection, 15: 239-258.

#### PERSONAL AND LABORATORY NEWS

D. Sperlich is now at Tübingen, Germany, from Vienna, Austria. New address: Institut für biologie, Lehrstuhl für Genetik, Aus der Morgenstelle 1, D-7400 Tübingen, Germany.

After September 1, 1971 the address of Th. Dobzhansky, F.J. Ayala and Mrs. Olga Pavlovsky will be: Department of Genetics, University of California, Davis, California 95616.

J.K. Choo is now at the Physiological Department, National Institute of Genetics, Misima, Japan (from Chungang University, Seoul, Korea).

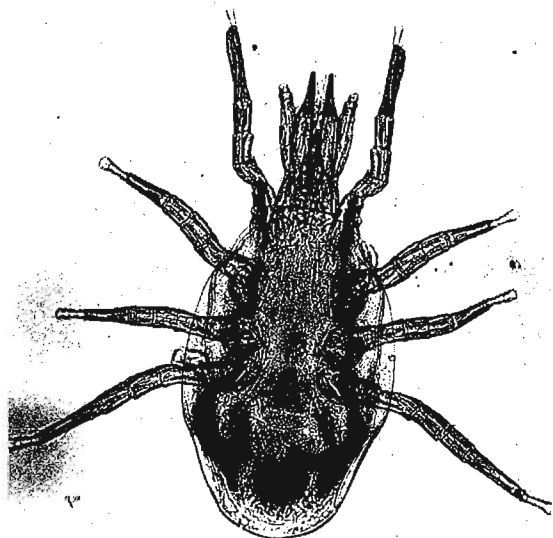
A. Shearn is now Assistant Professor in the Department of Biology, Johns Hopkins University, Baltimore, Maryland 21218 (from Yale University, New Haven, Conn.).

P.Dennis Smith is now Assistant Professor in the Department of Biology, Emory University, Atlanta, Georgia 30322 (from Storrs, Conn.).



Félix, R., J. Guzmán, M.E. de la Rosa and O. Olvera. Control of mites in Drosophila cultures.

Mites are introduced into laboratories when collections of wild flies are being made, or when Drosophila cultures received from other laboratories are not carefully examined before new cultures are started from such flies.



Histiotostoma sp.

Histiotostoma sp. is the most serious pest found in Drosophila cultures, as the control of infestation with this predatory mite has proved to be more difficult than the well known contaminations with either molds or bacteria.

Histiotostoma sp. has a hypopus stage which attaches itself to Drosophila, as well as to other insects. It differs from other less dangerous mites mainly by its absence of long hairs. Besides, the predatory mite has a squatting body build in contrast with the long and thinner non-parasitic mites. As female adults produce both male and female progeny by parthenogenesis, a laboratory might become infected from a single introduced mite in the hypopus stage.

The newly eclosed smallest nymphs thrive on the culture medium. After a week or so, they metamorphose into the migratory (hypopus) state. These hypopi, that develop in large numbers in old, infested cultures, are extremely active. They crawl up out of the culture medium and, as they penetrate tiny crevices, they may infest other cultures, unless very tightly stoppered. The migratory nymphs attach to any insect they may come into contact with, sucking the mouth parts into the insect. After ten days of attachment, they leave the host and grow to the adult stage on the surface of the medium, where they reproduce.

A heavy infestation which extended to stock cultures occurred last year at this laboratory.

The mites were introduced with samples of wild flies regularly collected at several trapping sites in Mexico City. All the infested cultures were submitted to the treatment detailed below, in order to eliminate all the mites from the cultures. Contaminated bottles and instruments were heated in a furnace before washing. The instruments used to manipulate flies, as well as the surface of the microscope, the outer surface of bottles, and upper surface of tables were continuously washed with a solution of benzyl-benzoate (20%) in 96° ethanol.

To start new cultures, flies were examined under the microscope, to use only adults that were apparently free from mites. As it is difficult to avoid contamination of the new medium, the adults were allowed to lay eggs on it, only during a period of 24 hours. When a few of the small hypopus nymphs from contaminated flies, or from other cultures were found in the new medium, it was necessary to cover its surface with a solution of benzyl-benzoate (20%) in 96° ethanol. This treatment kills the nymphs, without producing any noticeable effect in Drosophila larvae, which develop to the adult stage without hinderance from depredatory mites. Newly emerged flies were transferred each day to new cultures, to avoid the attachment of mites which survived after the treatment.

In heavily infested cultures all of the flies died, and there was found to be a crowding of mite nymphs among Drosophila larvae. In such a case, it was necessary to apply another treatment, thoroughly washing the larvae by immersion in a solution of benzyl-benzoate (20%) in ethanol. After 2-4 minutes in the benzyl-benzoate solution larvae were washed with Ringer solution and transferred to fresh vials. Following the above steps, the pest was effectively controlled after three weeks.

Grossfield, J. and J. Smith. Purdue University, Lafayette, Indiana. Video taping *Drosophila* behavior.

general applicability for the analysis of *Drosophila* behavior. A TV camera with its lens removed (A in Fig. 1) is mounted vertically on a trinocular dissecting scope. This allows the microscope adjustment to focus the camera. A 10X eyepiece is located in the phototube supporting the camera. A lower power would give a wider field of view. The problems of glare from wings and thorax and heavy shadows can be compensated for by diffusing incident light, placing a set of polarizers in the light path, balancing light with aluminum foil reflectors, and using a deep pile underlay (velvet) on the bottom of the lucite observation chamber. A  $1/4$  wave plate

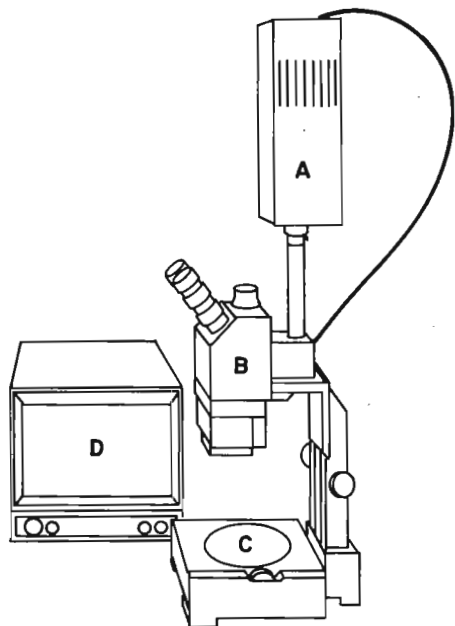


Figure 1. A. TV Camera; B. Dissecting scope with trinocular head; C. Observation Chamber, flat surface; D. TV monitor.

can also be used to cut glare. Any remaining glare can be compensated for by turning down the automatic gain control on both the camera and the monitor. A light coat of vaseline on the inside vertical surfaces of the observation chamber is reasonably effective in keeping the flies from assuming poorly photogenic positions on the corners or walls of the chamber.

For work in the dark a flashlight with a red filter (650 nm cut off, no UV transmittance) is a sufficient light source since the vidicon tube in the TV camera is

sensitive to infrared light. IR Image Converter equipment can be used to work with wavelengths further towards that region of the spectrum (RCA laboratories, David Sarnoff Research Center, Princeton, N.J.).

A videotape recorder can be interposed in the system. This yields the capability to stop action at any point in a behavioral sequence and measure distances (angles of body parts, etc.) on the face of the TV monitor during playback. The videotape records can, of course, be stored to form a library of behavioral activities. In the long run this method is less expensive than using and processing 16mm film.

If you have sufficient funds to think about color TV, we'd be glad to hear where you got them.

Zalokar, M. Centre de Génétique Moléculaire, CNRS 91, Gif sur Yvette (France). Fixation of *Drosophila* eggs without pricking.

Because of the impermeability of the vitelline membrane, the usual cytological fixatives can not penetrate the *Drosophila* egg and the egg has to be pricked to facilitate their entry. Only

Carnoy fixative can be used directly and then only if its content of chloroform is higher than in recommended formulas, but this fixation shrinks eggs very badly.

Lipid solvents can penetrate the vitelline membrane, and if they contain a fixative in solution, they can carry it across the membrane. Any fixative which is soluble both in the solvent and in water will diffuse into the ooplasm and partition itself between its aqueous phase and the solvent according to the phase rule. If we want to fix an egg with 50% acetic acid, we should shake the solvent with the acid of this concentration. The solvent will take up the acid at the proper concentration so that the acid entering an egg submerged in the solution will reach 50%.

If we use a solvent which does not disrupt the egg lipids too drastically, we can achieve

fixation which is equivalent to fixation by the corresponding aqueous fixative. It was found that heptane or octane did not injure the cytoplasm unduly, while penetrating well through the vitelline membrane. An egg remains alive if submerged in these solvents for 10 minutes or more. The eggs become fixed in heptane loaded with acetic acid, picric acid, acrolein or glutaraldehyde, in less than one minute and can remain in the fixative for several minutes before beginning to shrink.

In order to facilitate the penetration of post-fixatives, colorants or dehydrating liquids, the vitelline membrane should be removed after initial fixation. To do this, the egg is transferred into the aqueous phase of the fixative and the membrane torn away with sharp needles. Surface tension helps to remove the membrane and the egg falls into the liquid. This operation can be performed best in 30% acetic acid, but after some practice, one can do it also in other fixatives.

Fixation in heptane containing acrolein or glutaraldehyde is quite adequate for electron microscopy. Cell inclusions and organelles are well fixed, the ergastoplasm has its normal appearance and mitochondria have well preserved cristae. The following procedure is used:

1. Dechorionate eggs.
2. Fix in heptane which has been shaken with a 10% solution of acrolein or 25% solution of glutaraldehyde, for 1 to 2 minutes.
3. Remove the vitelline membrane in a buffered glutaraldehyde solution (conventional electron microscopy fixative).
4. Fix in the same solution for 1 hour.
5. Wash with buffered physiological solution.
6. Post fix with osmic acid 2 to 24 hours.
7. Further processing for embedding like any other tissue.

This fixation may be useful also in other cases where lipophilic membranes prevent the penetration of the usual fixatives, e.g. to fix *Drosophila* larvae and adults.

Félix, R. National Commission of Nuclear Energy, Mexico City, Mexico. A system for feeding liquids to adult flies.

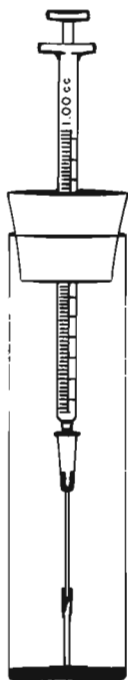
The following method may be used as an alternative to injection of solutions to *Drosophila* flies, especially when several treatments with liquids should be tested in adults at separate time intervals. This

system is particularly effective as the solution is administered during a period of time that may be lengthened to several days. It proved effectual for feeding cyclamates and cyclohexylamine to *Drosophila melanogaster*.

The liquid is gradually injected by means of a thin hypodermic syringe that goes through a hole of a rubber plug occluding the 2.8 x 9.0 cm vial, into a double layer of filter paper. The piece of polyethylene tubing (Intra-med, Clay Adams, Inc.) adapted to the needle of the syringe, touches the filter paper, assuring a continuous delivery of the solution, when the embolus is pushed in.

The quality of filter paper cut to fit the bottom of the vial is important because it must be sufficiently absorbent to remain moist, without retaining an excessive amount of solution, which would drown the flies. Whatman 3 filter paper was used for such a purpose.

The syringe may be removed without the removal of the rubber plug, thus avoiding the escape, as well as the squashing of the flies, that occurs if the plug is removed and replaced. The amount of solution contained in the syringe (1.00 cc B.D. Yale turbeculin, Becton Dickinson) is enough to feed flies during several days. An additional pasteur pipette made at the laboratory with thin glass tube may be adapted through another hole, assuring the proper aeration of the vial, if the system is to be used during a period of several days without the removal of the flies.



Erk, F.C., H.V. Samis, M.B. Baird and H.R. Massie. Masonic Medical Research Laboratory, Utica, New York. A method for the establishment and maintenance of an aging colony of *Drosophila*.

The routine maintenance of an aging colony is highly desirable for studies involving senescence in *Drosophila*. A major difficulty in maintaining such a colony is the lack of coincidence between the generation time of *D. melanogaster* and the normal work week. In this note the details are given

of a method for the initiation and maintenance of such a colony, as developed in this laboratory. The method will be described in terms of codes we arbitrarily use. Details of the environmental conditions for the maintenance of the colony are given elsewhere<sup>1</sup>.

The original stock of *D. melanogaster* (Oregon-R) was obtained from the Division of Biological Sciences, State University of New York at Stony Brook. These flies, designated  $P_0$  (parental), were allowed to lay eggs over a single 24-hour period; these eggs were collected on Thursday (Th). Cultures resulting from these eggs were labelled  $A_1P$  (A group, parental). Eggs were again collected on Friday (F), and these cultures were designated  $A_0A$  (A group, Aging Colony). During the following week, eggs from  $P_0$  were again collected on Th, but labelled  $B_1P$ . Eggs from  $P_0$  collected on F were labelled  $B_0A$  (B group, Aging Colony). By Monday (M) of the third week,  $A_1P$  flies had eclosed and were transferred to fresh medium.  $A_0A$  flies eclosed on Tuesday (T) and entered the aging colony.  $P_0$  eggs were collected on Th and labelled  $C_1P$ , while eggs collected on Friday were labelled  $C_0A$  (C group, Aging Colony).  $P_0$  flies were then discarded.

During week 4,  $B_1P$  flies were collected on M and placed on fresh medium, and  $B_0A$  flies entered the aging colony on T. Eggs from  $A_1P$  flies were collected on Th and designated  $A_2P$ , while  $A_1P$  eggs collected on F were designated  $A_1A$ . The  $A_1P$  flies were then mixed with  $A_0A$  flies which were already in the aging colony; these groups of flies had the same parents. This process is repeated during week 5 and subsequent weeks. During week 5, for example,  $C_1P$  flies were placed on fresh medium on M,  $C_0A$  flies were placed in the aging colony on T. Eggs from  $B_1P$  were collected on Th and designated  $B_2P$ , while eggs collected on F were labelled  $B_1A$ .

It becomes apparent (with pencil, paper, and time) that an aging colony of *Drosophila* may be maintained conveniently through standardized manipulation of these 3 groups of flies, all of which were derived from common parental stock. The continued input to the aging colony (e.g.,  $A_1P$  and  $A_0A$ ) may be adjusted to experimental needs. Our usual input is approximately 10,000 flies (mixed males and females) per week, but is being expanded to 30,000 flies per week. This method produces large numbers of flies which are one week apart in age. In addition, the parental age of all flies in the aging colony is held constant at 11-12 days.

Reference: 1. Samis, H.V., Jr., Erk, F.C. and Baird, M.B. 1971. *Exper. Gerontol.* 6: 9-18.

Merriam, J.R., University of California, Los Angeles, California. A low cost disposable bottle for *Drosophila* culture.

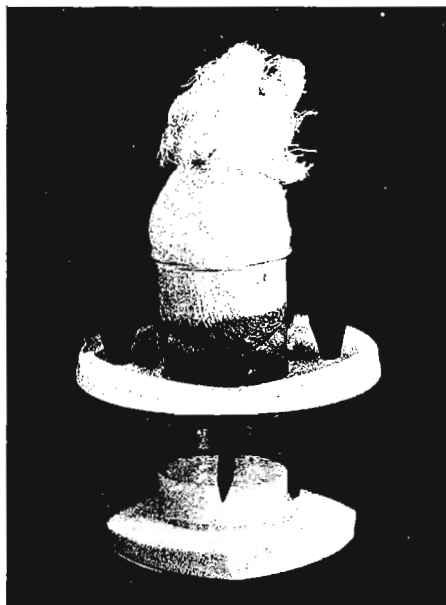
Propak-California Corp., 211 N. Willow Ave. City of Industry, Ca. 91746 (phone 213-968-6447) sells 1/2 pint cylindrical milk bottles 5.3 cm in diameter, made of translucent plastic, which we use for all

our cultures. The bottles are transparent enough so that the food condition and the pupae on the bottle walls can be easily seen. Although we originally bought them because of their low cost (currently \$1.70 per 100 bottles) other advantages over the usual glass bottles are also important: they weigh much less and take up less space. A shipping carton with 400 bottles weighs just 15 3/4 lbs. This enables us to use low cost baskets "homemade" from welded wire fencing. We plug the bottles with dispo plugs 28 x 35 mm obtained from Scientific Products. Snap lids did not work. We find it economical to clean and reuse the bottles, although other labs might want to use them for large one-shot projects or classes, or as a low cost reserve. Schools that want to set up fly keeping on a small scale or for a limited time will find these bottles especially attractive. If anyone would like to see a bottle we will be glad to send an empty on request within the U.S.

Hedrick, P.W. University of Kansas, Lawrence, Kansas. A culture which allows sand pupation.

Many species of *Drosophila* prefer to pupate in sand rather than on paper or the sides of a culture bottle. Several types of cultures have been used for these *Drosophila* but they usually entail trans-

ferring the larvae to a second container. The device I am using now for study of niche separation permits larvae to have the option of pupating either in sand or on the side of a chimney. Furthermore, this container permits downward migration of larvae into the sand simulating the behavior of *Drosophila* larvae in fallen fruit.



The container used is a clear plastic, eight ounce refrigerator container made by Deka Plastics, Inc. (see photograph). In the bottom of the container is placed approximately one eighth cup of sterilized white sand. In order to prevent dehydration of the media, the sand is saturated with water. The media is poured into a 3 inch high, 1-1/2 inch diameter pyrex chimney which has been placed on aluminum foil. After the media has set, the foil is peeled away and the chimney slipped through a hole in the container lid made earlier by a hot pyrex chimney. Larvae or adult flies are placed in the chimney and it is stoppered by a cotton-cheesecloth plug.

One must use an aspirator in order to remove flies from the container. To remove those inside the container, an aspirator tube is placed inside a plug. Flies which emerge outside the chimney, that is pupate in the sand, are aspirated through several holes which have been drilled in the container lid. These are stoppered by golf tees when not in use. Even with saturated sand, I have encountered some shrinkage of the media. This is dependent on the media used, the humidity, and the amount of larvae working.

When adult flies are used, I suggest that they be allowed to lay on the media for 24-48 hours before placing the chimney in the container. I am indebted to J.S.F. Barker for suggesting many of the ideas in the design of this container.

Tomkins, J.K. and T. Billington. Monash University, Clayton, Vic., Australia. Analysis of *D. melanogaster* RNA by acrylamide gel electrophoresis of single fly homogenates.

The acrylamide gel electrophoresis procedure of Loening (1967) as modified by Becker et al. (1971) has been adapted for the analysis of RNA from homogenates of single individuals of *D. melanogaster* developmental stages. Third instar, pupa or adult individuals have been used successfully in this study. The method

permits the resolution of ribosomal and transfer RNA as distinct sharp bands.

An individual of the developmental stage to be studied is briefly washed in distilled water before transfer to a small all-glass homogenizer. Homogenization is carried out at room temperature in 0.1 ml Loening's electrophoresis buffer containing 1% (w/v) sodium dodecyl sulphate (SDS) and 10% (w/v) sucrose. The homogenate is then stood at room temperature for 1 1/2 hours.

Acrylamide gels are prepared by the method of Loening. The gels are pre-run at 5 mA/gel for 1 hour at 4° in electrophoresis buffer containing 0.1% (w/v) SDS. This pre-run is in the direction the sample electrophoresis is to take place. The buffer is then renewed and a further pre-run, in the opposite direction, of 1 hour is carried out.

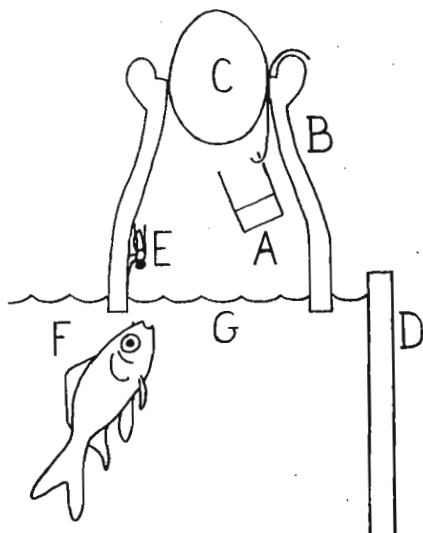
The homogenate is applied to the gels and electrophoresis at 5 mA/gel in fresh 0.1% SDS buffer is carried out at 4°. After electrophoresis the gels are removed from the tubes, fixed, stained and destained according to the method of Solymosy et al. (1970).

References: Loening, U.E., 1967, *Biochem. J.* 102: 251-257; Becker, H., C.P. Stanners, and J.E. Kudlow, 1971, *J. Cell. Physiol.* 77: 43-50; Solymosy, F., G. Lazar and G. Bagi, 1970, *Anal. Biochem.* 38: 40-45.

Wong, P.T. and W.E. Trout III. City of Hope National Medical Center, Duarte, California. The psychic fly fish feeder: A reply to Ogden Nash.

"The Lord in His wisdom made the fly  
And then forgot to tell us why."  
The Fly, by Ogden Nash

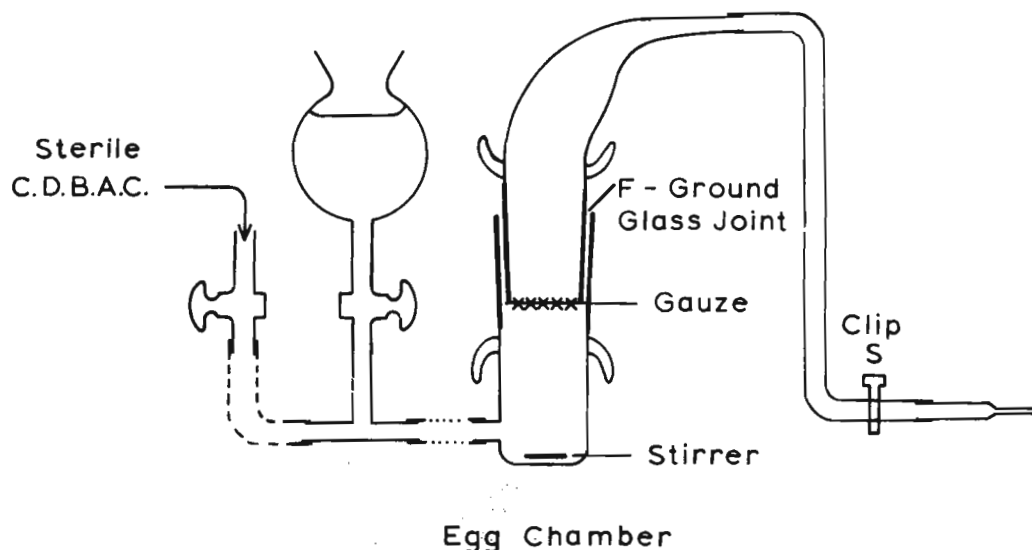
A culture of the neurological mutant  $Hk^1$  (who falls over when he sees something moving), is placed in container (A), in bottomless bottle (B), plugged with cotton (C), which is suspended at the surface of the water in aquarium (D). When a fly emerges (E) and sees fish (F) moving below him, he loses control of himself and falls into the water (G) thereby feeding fish (F).



Sparrow, J.C. University of Sussex, Brighton, England. Eggwashing apparatus.

This apparatus replaces the reversing pump described by Sang (1956) in our *Drosophila* egg sterilization procedure for axenic culture. The eggs are col-

lected by the method of Sang (1956), dechorionated with hypochlorite and suspended in 1.0% saline. This stops the dechorionated eggs from aggregating. Any larvae are removed by hand using a binocular microscope. The eggs are then placed in the apparatus through the thistle funnel and enter the egg chamber. They are washed in this chamber by a sterile 0.1% C.D.B.A.C. (cetyltrimethylbenzylammonium chloride) solution which runs continuously through the apparatus. The flow rate is controlled by the screw clip (S) at a rate of 90 drops/minute



from the standard pasteur pipette. The eggs are kept in suspension and from sticking to the gauze by means of a small magnetic stirrer, revolving at about 74 revs./minute. The figures for flow rate and stirring are those which give optimum egg-hatching rates with minimum culture infection. The eggs undergo 15 minutes of this treatment and then the egg chamber is removed from the apparatus. The egg chamber must be washed through with sterile water before plating so as to remove any C.D.B.A.C. Prolonged exposure to this detergent was found to kill eggs. Some strains appear more sensitive than others to the length of time spent in C.D.B.A.C. The egg chamber is then dismantled at (F) in a sterile hood and the eggs plated on agar using the procedure of Sang (1956).

Reference: Sang, J.H. (1956). J. Exp. Biol. 33:45-72.

Lewis, E.B. and L. Craymer. California Institute of Technology, Pasadena, California. Quinacrine fluorescence of *Drosophila* chromosomes.

We describe below a modification we have developed for *Drosophila* of the quinacrine-fluorescent staining methods developed by Caspersson and his colleagues (Expt. Cell Res. 58: 141-151, 1969) for plant and, later, human chromosomes. Our findings are in general

agreement with those of Vosa (Chromosoma 31: 446-451, 1970) who reports selective staining with this dye of the long arm of the Y chromosome and of the fourth chromosome in *Drosophila melanogaster*. In addition we have observed a bright fluorescing band in region 8lF (3R) of the salivary gland chromosomes. Also in larval ganglion metaphases a weakly fluorescent spot is visible in the basal heterochromatic region of X, 2L, 2R and 3L, while 3R has two such spots. The Y has at least three strongly fluorescing spots in Y-long and at least one in Y-short (Y<sup>closed</sup> has five spots visible); the fourth chromosomes appear at metaphase as two very bright fluorescent bodies. Adult muscle or brain tissues show in resting nuclei a large, usually single, fluorescent body, which may represent a chromocentral fusion of the fluorescent spots found in the basal part of each chromosome arm. The XY male has a somewhat brighter body, evidently due to fusion of the Y chromosome as well. It may prove possible therefore to "sex" somatic resting nuclei of adult tissues.

The resting nuclei of imaginal disc cells also have a single large fluorescing body in XX or XO tissues but tend to have two such bodies, of approximately equal fluorescent intensity, in XY or XXY resting nuclei. We interpret this to mean that in these rapidly dividing tissues the Y often does not fuse with the chromocenter. We have also extended this observation by studying males carrying an extra Y-long arm attached to X in addition to a normal Y. Such males often show three fluorescing bodies in the resting nuclei of their imaginal wing discs instead of two, suggesting that extra Y's do not tend to fuse with one another in imaginal disc tissue.

A. Procedure for staining *Drosophila* salivary gland chromosomes or resting nuclei of many larval or adult (except brain) tissues.

1. Dissect larvae in 45% acetic acid. Place tissue in a small drop of 45% acetic acid on a siliconed coverslip. Lower a slide which has been "subbed"\* over the drop; after it touches the drop, invert the slide; blot to remove excess mounting fluid; tap the coverslip sharply over the tissue area with a blunt instrument to disperse cells; cover with absorbent tissue and squash with strong pressure.

2. Immerse slide in liquid nitrogen until bubbling stops (or freeze on a block of dry ice).

3. Pry off coverslip with a razor blade. Dip slide in 95% alcohol for about a minute and then into absolute alcohol for a minute. Remove and dry by waving in the air.

4. Flood area over tissue with a few drops of an 0.5% to 2% solution of quinacrine hydrochloride in 45% acetic acid and stain for one or two minutes. (Batches of Gurr's "atebrin" or Sigma's quinacrine have proved satisfactory.)

5. Drain off staining solution and quickly dip slide into a jar of 95% ethanol followed by one or two transfers through absolute ethanol. The total time in the alcohols should be 20 seconds or less at 25°C to avoid excessive destaining. Insufficient rinsing may result in excessive background fluorescence in the final preparation.

6. Remove slide from the absolute ethanol and quickly dry by waving the slide vigorously in air. Heating at this stage tends to destain the preparation.

7. To mount, place a drop of sucrose solution (0.5 to 1.0 molar in double distilled water) in the center of a coverslip. Invert the slide over this coverslip until it touches

the drop. Quickly reinvert and blot firmly to remove excess mounting fluid. Seal with clear nail polish.

**B. Procedure for staining adult brain tissue.**

1. The same procedure as that described for the larvae is used except that the percentage of acetic acid in both the dissecting fluid and the quinacrine staining solution is reduced from 45% to 10%.

**C. Procedure for staining larval ganglia for metaphase chromosome studies (modified from DIS 34: 118-119).**

1. Dissect larvae in a solution of 1.0% Na Citrate in distilled water. Place the dorsal ganglia in a drop of this solution for 10 minutes on a slide. Warm the slide on a hot plate at 40°C for one minute (this hastens separation of sister chromatids). Pass the ganglia into a pre-fixative composed of equal parts of 45% acetic acid and 95% ethanol and leave for 30 seconds. Then remove tissue and place in a drop of 45% acetic acid on a siliconed coverslip. Continue with procedure described in part A, par. 1 above.

\*3 gm gelatin. 600 ml distilled water. Heat to dissolve gelatin. Cool. Add chrom. alum -  $\text{KCr}(\text{SO}_4) \cdot 12 \text{H}_2\text{O}$  - 300 mg. Dip slides, drain and allow to dry in dust-free container.

#### TEACHING NOTE

Potter, J.H. University of Maryland, College Park, Maryland. A demonstration of compensation for an inherited biochemical defect in *D. melanogaster*.

A simple demonstration of compensation for an inherited biochemical defect can be carried out by beginning students using *D. melanogaster*. In essence, students supply kynurenine to larvae of vermilion mutants which cannot convert tryptophan to kynurenine, one of the steps

in the synthesis of ommochrome pigments. Since students frequently do not distinguish vermilion from wild type flies, they use the white-eyed, double mutant, vermilion brown. Vermilion brown larvae fed kynurenine develop brown eyes. To emphasize the specificity of the block, students also feed kynurenine to the double mutant, cinnabar brown. Cinnabar brown mutants develop white eyes whether or not they receive kynurenine.

**Experimental procedure:** Students set up two cultures each of vermilion brown and cinnabar brown mutants in 80 x 25 mm. shell vials containing 5 ml of Carolina Instant *Drosophila* Medium. As soon as larvae appear the parents are removed and the medium in one vial of each genotype is injected with 0.2 ml of a kynurenine-antibiotic solution. The medium in the other two vials is injected with 0.2 ml of plain antibiotic solution. The injections are made with a 2 1/2 ml syringe without a needle inserted in a hole made in the medium with an applicator stick. Injections are repeated every two days until pupae appear. The adults are scored in the usual way. The kynurenine treated, vermilion brown, flies are mated after scoring and their progeny scored for eye color to demonstrate that the genotype has not been changed by the kynurenine treatment.

The kynurenine antibiotic solution is similar to that used by Parsons and Green (1959) for culturing eye discs: 0.05% streptomycin, 0.033% penicillin and 1.00% D.L. kynurenine can be obtained from Sigma Chemical Co., St. Louis, Missouri, at \$14.00/gram.

References: Parsons, P.A. and M.M. Green, 1959, Proc. Nat. Acad. Sci., Wash. 45: 993.

#### MATERIALS REQUESTED OR AVAILABLE

H.R. Feijen, University of Malawi, Genetics Section, P.O. Box 5200, Limbe, Malawi, would be grateful to obtain reprints on speciation in *Drosophila* and reprints on systematics of *Drosophila*.



R.C. King, Dept. of Biology, Northwestern University, Evanston, Illinois, is undertaking the editing of a Handbook of Genetics for the Van Nostrand-Reinhold Company, and intends to include a cytogenetic map of *Drosophila melanogaster*. In his monograph on Ovarian Development in *Drosophila melanogaster*, he published a fold-out which included the cytological localization of about 71 genes or gene clusters. This used the data from Lindsley and Grell which by now are four years old. He would therefore appreciate hearing from *Drosophila* workers who have new data (published or unpublished) that would enable him to update the comparative maps. Also any suggestions as to other material that should be covered will be greatly appreciated.

Discontinuance of *Drosophila* Stock Service: After 31 years, the *Drosophila* service at Cold Spring Harbor has been discontinued. Copies of the "*Drosophila* Guide" will continue to be available from: Office of Information, Carnegie Institution of Washington, 1530 P Street, Northwest, Washington, D.C. 20005. Flies can be purchased from several sources, including the following: Carolina Biological Supply Company, Burlington, North Carolina, 27215 and Gladstone, Oregon 97027; Ward's Natural Science Establishment, Inc., Post Office Box 1712 Rochester, New York 14603.

The Seton Hall University stock list has been discontinued.

Announcing publication of "Ovarian Development in *Drosophila melanogaster*," by R.C. King, Department of Biological Sciences, Northwestern University, Evanston, Illinois, June 1970, 227 pp, \$16.50, Academic Press, New York, N.Y.

The Egyptian Society of Genetics is putting out a new Journal entitled "Egyptian Journal of Genetics and Cytology". Submission of publications from the different fields of genetics and cytology is encouraged. It is hoped that this new Journal will be of great help to researchers in genetics all over the world. The Journal will be in two numbers and about 300 pp. a year, starting Vol. 1 No. 1, January 1972. The subscription rates are \$10 for institutions and \$5 for individuals. Subscriptions are to be ordered through the Editorial Office (Prof. A.O. Tantawy, Dept. of Genetics, Faculty of Agriculture, Alexandria University, Alexandria, Egypt). All checks should be made payable to the Egyptian Society of Genetics.

The Genetics and Biology of *Drosophila*. Announcement of a proposed three volume work, intended to cover comprehensively the genetics and biology of *Drosophila*.

The field of *Drosophila* research suffers from the absence of any modern comprehensive account of the biology and genetics of the organism. No complete account of *Drosophila* genetics has been published since 1925, although individual topics have often been well reviewed in recent years. In addition the current accounts of *Drosophila* biology (Demerec) and evolution (Patterson and Stone) are, although irreplaceable, either out of date or fail to cover the many fields of research which have opened up dramatically in the last 20 years. The size and breadth of the current *Drosophila* literature is a severe handicap to students starting research in *Drosophila*, to workers switching to *Drosophila* from other fields and to *Drosophila* workers themselves with interests in aspects of *Drosophila* research other than their immediate research interest.

The current project, to publish in three volumes an account of the genetics and biology of *Drosophila*, is an attempt to fulfill the requirements of the research biologist. Each volume will consist of a number of chapters by invited authors. The standing brief to each author is that he makes an attempt to cover comprehensively his particular topic with emphasis on its historical context and in particular the techniques used and its place in modern *Drosophila* research. Although emphasis will obviously be on *melanogaster*, especially in the volume on genetics, information from other species, and where necessary from other insects, will also be included. It would, for example, be quite illogical to consider certain aspects of the developmental biology of *Drosophila* without drawing on the literature concerned with similar problems in other species of flies and other insect groups.

The three volumes, which will be published consecutively at yearly intervals, will be as follows: Vol. I. Genetics; Vol. II Biology and Development; Vol. III Evolution, Taxonomy and Ecology. Volume I is being edited by Michael Ashburner and Ed Novitski and is planned for publication in 1973.

ARGENTINA

Buenos Aires: Comisión Nacional de Energía Atómica, Departamento de Radiobiología, División Genética. Libertador 8250. Tel 70-7711 Ext 124

Kirschbaum, W.F. B.Sc.Agr. Research Associate Salivary cytology  
 Mazar Barnett, B. (Mrs.) Ph.D. Radiation genetics & chemical induction of mutations  
 Muñoz, E.R. M.D. Radiation genetics (on leave during 1971 at the University of Leiden Netherlands)  
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Almazán, M. Spain, Madrid  
Alvarez, M. C. Spain, Madrid  
Anderson, M. Berkeley, California  
Anderson, S. Austin, Texas  
Anderson, W.W. New Haven, Connecticut  
Andjelković, M. Yugoslavia, Belgrade  
Andrés, R. de Spain, Madrid  
Aotani, S. Japan, Osaka  
Aotsuka, T. Japan, Tokyo  
Araújo, H.S. Brazil, Pôrto Alegre  
Arcos-Teran, L. Germany, Tübingen  
Arens, M.F. France, Lyon  
Arias, T.B. Colombia, Bogotá  
Armstrong, M. Canada, Vancouver  
Arnheim, N. Jr. Stony Brook, New York  
Ash, W.J. Canton, New York  
Ashburner, M. England, Cambridge  
Atherton, J. England, Brighton  
Averhoff, W.W. Austin, Texas  
Ayala, F.J. Davis, California  
Ayers, A.J. San Marcos, Texas  
Babb, E.E. Lafayette, Indiana  
Bächli, G. Switzerland, Zürich  
Back, H. Korea, Chungang  
Badr, E. U.A.R., Alexandria  
Bahn, E. Denmark, Copenhagen  
Baillie, D.L. Storrs, Connecticut  
Baird, M.B. Utica, New York  
Balkin, K.E. Pullman, Washington  
Baker, B. Seattle, Washington  
Baker, J. England, London  
Baker, W.K. Chicago, Illinois  
Band, H. East Lansing, Michigan  
Banerjee, M. India, Calcutta  
Bannerjee, R. India, Calcutta  
Barigozzi, C. Italy, Milan  
Barker, J.S.F. Australia, Sydney  
Barnes, B.W. England, Birmingham  
Barr, H.J. Madison, Wisconsin  
Barrett, J.A. England, Cambridge  
Basden, E.B. Scotland, Edinburgh  
Battaglia, B. Italy, Padova  
Bauer, G. Germany, Düsseldorf  
Baxa, H. Austria, Vienna  
Beatty, M.E. Morgantown, West Virginia  
Beavers, G. Austin, Texas  
Becerra, E.L. Colombia, Bogotá  
Beck, H. Switzerland, Geneva  
Beck, M.L. Fayetteville, Arkansas  
Becker, G.L. Germany, München  
Becker, H. Germany, München  
Beerefenger, D. Netherlands, Leiden  
Beermann, W. Germany, Tübingen  
Beeson, V. Notre Dame, Indiana  
Belitz, H.J. Germany, Berlin  
Bell, A.E. Lafayette, Indiana  
Bender, H.A. Notre Dame, Indiana  
Benedik, J. Czechoslovakia, Brno  
Benner, D.B. Johnson City, Tennessee  
Bennett, J. DeKalb, Illinois  
Bennett, K. DeKalb, Illinois  
Benozatti, M.L. Brazil, São Paulo  
Bentley, M. Wooster, Ohio  
Benzer, S. Pasadena, California  
Berger, E. Chicago, Illinois  
Berkes, J. Syracuse, New York  
Bernard, J. France, Gif-sur-Yvette  
Bhalla, D. India, Chandigarh  
Biemont, C. France, Lyon  
Biggers, J.M. Carbondale, Illinois  
Bigonnet, C. France, Lyon  
Bijlsma, R. Netherlands, Haren  
Bingham, D. San Bernardino, California  
Bischoff, W.L. Chapel Hill, North Carolina  
Bishop, A.M. Lexington, Kentucky  
Black, M. Urbana, Illinois  
Bleyman, M.A. Chapel Hill, North Carolina  
Bloom, P. Iowa City, Iowa  
Blumenfeld, M. Austin, Texas  
Boam, T.B. England, Sheffield  
Bock, I.R. Austin, Texas  
Bocquet, C. France, Gif-sur-Yvette  
Bodenstein, D. Charlottesville, Virginia  
Boerema, A.C. Netherlands, Haren  
Boesiger, B. New York, New York  
Boesiger, E. New York, New York  
Boettcher, B. Australia, Adelaide  
Boo, H.A. de Netherlands, Leiden  
Booker, S.J. Raleigh, North Carolina  
Boot-Wassenaar, M.C. Netherlands, Leiden  
Borai, F.M. U.A.R., Alexandria  
Borner, P. Switzerland, Zürich  
Bortolon, A.H. Brazil, Pôrto Alegre  
Bos, M. Netherlands, Haren  
Boshes, R. New Haven, Connecticut  
Bosiger, E. France, Gif-sur-Yvette  
Bouletreau, M. France, Lyon  
Bouletreau-Merle, J. France, Lyon  
Bowman, H. England, Heslington  
Bownes, M. England, Brighton  
Boyd, J.B. Davis, California  
Boyer, J.F. Iowa City, Iowa  
Bras, F. France, Gif-sur-Yvette  
Bray, R. Australia, Bundoora

Bregliano, J.-C. France, Clermont-Ferrand  
Bremner, T.A. Washington, D.C.  
Brenman, M. Utica, New York  
Bretschneider, F. Netherlands, Urecht  
Breugel, F.M.A. van Netherlands Leiden  
Brigman, J. Buffalo, New York  
Brink, N.G. Australia, Adelaide  
Brito da Cunha, A. Brazil, São Paulo  
Broderick, D. Carvallis, Oregon  
Brodie, A.E. Madison, Wisconsin  
Brosseau, G.E. Iowa City, Iowa  
Browder, L. Canada, Calgary  
Brown, J. Raleigh, North Carolina  
Brown, S.W. Chicago, Illinois  
Brown, W.P. Marietta, Ohio  
Brugger, C. Zürich, Switzerland  
Brun, G. France, Gif-sur-Yvette  
Brüschweiler, W. Switzerland, Zürich  
Bryant, S. Riverside, California  
Buchanan, J.S. Cold Spring Harbor, New York  
Büchi, R.S. Switzerland, Zürich  
Buchner, E. Germany, Tübingen  
Bultmann, H. Seattle, Washington  
Burckhardt, H. Switzerland, Zürich  
Burdette, W.J. Houston, Texas  
Bürki, K. Switzerland, Zürich  
Burla, H. Switzerland, Zürich  
Burnet, B. England, Sheffield  
Burnham, M.B. Ann Arbor, Michigan  
Burton, S. Lafayette, Indiana  
Buruga, J.H. Uganda, Kampala  
Busby, N.J. Riverside, California  
Bussereau, F. France, Gif-sur-Yvette  
Bzdega, E. Scotland, Aberdeen  
Cama, J. Spain, Barcelona  
Camfield, R.C. Canada, Vancouver  
Campanario, E. Spain, Madrid  
Canabal, F.L. Brazil, Porto Alegre  
Capps, A.S. Austin, Texas  
Cardellino, R.A. Raleigh, North Carolina  
Carfagna, M. Italy, Naples  
Carlson, E.A. Stony Brook, New York  
Carpenter, A. Seattle, Washington  
Carpenter, J.M. Lexington, Kentucky  
Carpenter, N. Ann Arbor, Michigan  
Carson, H.L. Honolulu, Hawaii  
Carver, J.E. Jr. Houston, Texas  
Castillo, B.J. Duarte, California  
Castro, L.E. Colombia, Bogotá  
Catchside, D.E.A. Australia, Adelaide  
Caten, C.E. England, Birmingham  
Cavataio, P. San Bernardino, California  
Čejnová, H. Czechoslovakia, Prague  
Cetl, I. Czechoslovakia, Brno  
Chace, A. Chapel Hill, North Carolina  
Chan, L.L. New Haven, Connecticut  
Chang, Y.-C.K. Austin, Texas  
Charlesworth, B. Chicago, Illinois  
Chassagnard, M.T. France, Gif-sur-Yvette  
Chatterjee, B. India, Calcutta  
Chattopadhyay, S. India, Calcutta  
Chawdhury, K. India, Calcutta  
Chen, P.S. Switzerland, Zürich  
Chen, T. Canada, Edmonton  
Chew, G.K. Australia, Bundoora  
Chikushi, H. Japan, Fukuoka  
Chirombo, H. Malawi, Timbe  
Choi, Y. Austria, Vienna  
Choo, J.K. Korea, Seoul  
Chornoma, R. Upton, New York  
Chovnick, A. Storrs, Connecticut  
Christensen, P.T. Pasadena, California  
Christopher, W. Chapel Hill, North Carolina  
Christopoulou, A. Greece, Patras  
Chun, S.B. Korea, Kwangju  
Chung, C.U. Korea, Kwangju  
Chung, J.K. Korea, Seoul  
Chung, O.K. Korea, Seoul  
Chung, Y.-J. Korea, Seoul  
Clark, A.M. Australia, Adelaide  
Clayton, F.E. Fayetteville, Arkansas  
Clise, R. Cleveland, Ohio  
Cobbs, G. Riverside, California  
Cohen, E. New Haven, Connecticut  
Cohet, Y. France, Lyon  
Colaianne, J. Lafayette, Indiana  
Colin, C. Duarte, California  
Collett, J. England, Brighton  
Conner, B. Atlanta, Georgia  
Connolly, K.J. England, Sheffield  
Conscience-Egli M. Switzerland, Zürich  
Contamine, D. France, Gif-sur-Yvette  
Cook, R.M. England, Sheffield  
Cooper, K.W. Riverside, California  
Cooper, M. Australia, Adelaide  
Cordeiro, A.R. Brazil, Porto Alegre  
Cordova, F. Italy, Naples  
Corwin, H.O. Pittsburgh, Pennsylvania  
Counce, S.J. Durham, North Carolina  
Coyne, J.A. Williamsburg, Virginia  
Craddock, E. Honolulu, Hawaii  
Crépin-de Jong, T.A. Netherlands, Leiden  
Croft, J.H. England, Birmingham  
Crossley, S.A. Australia, Clayton  
Crow, J.F. Madison, Wisconsin  
Crowder, E. Tuscaloosa, Alabama  
Crumpacker, D.W. Boulder, Colorado  
Cuello, J. Spain, Barcelona  
Cummings, M.R. Chicago, Illinois  
Cuthbertson, D. Raleigh, North Carolina  
Cybul, G. Macomb, Illinois  
Dailie, J. France, Lyon  
D'Amora, D. Italy, Naples  
Danieli, G.A. Italy, Padova  
Dapples, C.C. Billings, Montana  
Darst, R.P. Chapel Hill, North Carolina  
Das, A.K. India, Calcutta  
Datta, R.K. India, Calcutta  
David, I. Dallas, Texas  
David, J. France, Lyon  
Davidson, T. Atlanta, Georgia  
Davis, B.K. San Diego, California

- Davis, D. Dallas, Texas  
Davis, D.G. Tuscaloosa, Alabama  
Day, John W. Ames, Iowa  
De, A. India, Calcutta  
Debouzie, D. France, Lyon  
De Domenicis, M.A. Italy, Naples  
Dee, J. England, Leicester  
De Garay, A.L. Mexico, Mexico City  
De Jongh, L.F. Madison, Wisconsin  
Deland, M.C. Storrs, Connecticut  
De la Rosa, E. Mexico, Mexico City  
Delden, W. van Netherlands, Haren  
De Marco, A. Italy, Rome  
DeMarinis, F. Cleveland, Ohio  
De Mitri, S. Italy, Naples  
Denell, R.E. San Diego, California  
De Niro, M.J. Pasadena, California  
De Reggi-Mourgues, C. France, Lyon  
DeRemer, J. Rochester, New York  
DeStefano, J. Philadelphia, Pennsylvania  
Deutsch, V. France, Gif-sur-Yvette  
Devaux, J. France, Gif-sur-Yvette  
Dewhurst, S. Duarte, California  
Diamantopoulou, E. Greece, Athens  
Diatla, F. France, Gif-sur-Yvette  
Dickerman, R.C. Cleveland, Ohio  
Dickson, E. Austin, Texas  
Dickson, L.R. Pasadena, California  
Diehl, E. Brazil, Pôrto Alegre  
Dijken, F.R. van Netherlands, Haren  
Dimsdale, C.H. Canada, Edmonton  
Divelbiss, J. Le Mars, Iowa  
Doane, W.W. New Haven, Connecticut  
Dobzhansky, Th. Davis, California  
Doira, H. Japan, Fukuoka  
Dolfini, S. Italy, Milan  
Donady, J.J. Duarte, California  
Donini, S. Davis, California  
Douma, O. Netherlands, Leiden  
Downing, B. Galesburg, Illinois  
Doyle, P. England, Cambridge  
Driskell, W.J. Pasadena, California  
Druger, M. Syracuse, New York  
Duberstein, R. New Haven, Connecticut  
Duck, P.D. Storrs, Connecticut  
Dudick, M. Charlottesville, Virginia  
Duffy, C. Los Angeles, California  
Dunn, B.K. Madison, Wisconsin  
Du Pui, M.L.L. Netherlands, Haren  
Dutta Gupta, A.K. India, Calcutta  
Dyer, K.F. Australia, Clayton  
Early, D.M. Lafayette, Indiana  
Eastwood, L. England, Sheffield  
Eaves, L.J. England, Birmingham  
Ebitani, N. Japan, Tokyo  
Edney, E.B. Riverside, California  
Eggert, R. Davis, California  
Ehrenfeld, J.G. New York, New York  
Ehrlich, E. Eugene, Oregon  
Ehrman, L. Purchase, New York  
Eiche, A. Sweden, Stockholm  
Eichenberger, E. Pasadena, California  
Elens, A. Belgium, Namur  
El-Kouni, M.H. Canada, Edmonton  
Ellison, J.R. Madison, Wisconsin  
El-Masry, A. Scotland, Aberdeen  
Elmer, W.A. Atlanta, Georgia  
Emara, M.K. U.A.R., Alexandria  
Emmens-Pieters, J. Netherlands, Haren  
Engel, C.M. Brazil, Pôrto Alegre  
Engelking, A.B. San Marcos, Texas  
Englert, D.C. Carbondale, Illinois  
English, D.S. Flagstaff, Arizona  
Eppenberger, H.M. Switzerland, Zürich  
Epstein, D. Garden City, New York  
Erickson, J. Bellingham, Washington  
Erk, F.C. Stony Brook, New York  
Eschiletti, J.A.F. Brazil, Pôrto Alegre  
Evans, Wm. Bellingham, Washington  
Ezell, S.D. Jr. Madison, Wisconsin  
Ezinga, K. Netherlands, Haren  
Fabergé, A.C. Austin, Texas  
Fahmy, A.M. U.A.R., Alexandria  
Fahy, T. Notre Dame, Indiana  
Fain, P. Boulder, Colorado  
Falk, D.R. Canada, Edmonton  
Falke, E. Charlottesville, Virginia  
Falkiner, J. Australia, Canberra  
Favalora, V. Garden City, New York  
Feijen, H.R. Timbe, Malawi  
Feijen-van Soest, J.J. Timbe, Malawi  
Félix, R. Mexico, Mexico City  
Fernandes, N. Brazil, São Paulo  
Ferreira, A. Brazil, Pôrto Alegre  
Feverbach-Mravlag, H. Austria, Vienna  
Figueroa, J.A. Chicago, Illinois  
Finlay, D.E. Australia, Sydney  
Fishburn, J. Iowa City, Iowa  
Fiorio, P. Duarte, California  
Fitz-Earle, M. Canada, Vancouver  
Fleming, C. Cleveland, Ohio  
Fleuriet, A. France, Clermont-Ferrand  
Fong, W.-F. Notre Dame, Indiana  
Fontaine, T.E. Minneapolis, Minnesota  
Fontdevila, A. Spain, Barcelona  
Forbes, C. Philadelphia, Pennsylvania  
Ford, A. Washington, D.C.  
Forero, I. Colombia, Bogotá  
Forrest, H.S. Austin, Texas  
Fouillet, P. France, Lyon  
Fourche, J. France, Lyon  
Fox, A.S. Madison, Wisconsin  
Fox, D.J. Switzerland, Zürich  
Francis, H. Australia, Adelaide  
Frankel, A. Iowa City, Iowa  
Frankham, R. Chicago, Illinois  
Franklin, I.R. Australia, Sydney  
Frei, H. Switzerland, Geneva  
Frey, J.J. Lafayette, Indiana  
Friedman, L.D. St. Louis, Missouri  
Friedman, T.B. Ann Arbor, Michigan  
Frijters, D.A.M. Netherlands, Urecht

- Fritz, M. Sweden, Stockholm  
 Fritz-Niggli, H. Switzerland, Zürich  
 Frutos, R. Spain, Barcelona  
 Fuchs, M.S. Notre Dame, Indiana  
 Fujii, H.M. Japan, Fukuoka  
 Fujii, S. Japan, Kobe  
 Fujimoto, M. Japan, Tokyo  
 Fukatami, A. Japan, Sakado-Machi  
 Fulch, D.G. San Diego, California  
 Fullilove, S.L. Austin, Texas  
 Gabay, S.J. Urbana, Illinois  
 Gale, J.S. England, Birmingham  
 Galia, M.S. Brazil, Pôrto Alegre  
 Gall, J.G. New Haven, Connecticut  
 Galmuzzi, G. Italy, Naples  
 Ganguly, R. India, Calcutta  
 Garcia-Bellido, A. Spain, Madrid  
 Garcia, P. Spain, Barcelona  
 Garen, A. New Haven, Connecticut  
 Garrido, P. Spain, Madrid  
 Gavin, J. Canada, Calgary  
 Gay, H. Ann Arbor, Michigan  
 Gay, P. France, Gif-sur-Yvette  
 Gaylord, C.A. Washington, D.C.  
 Greer, B.W. Galesburg, Illinois  
 Gehring, W. New Haven, Connecticut  
 Gelbart, W.M. Pasadena, California  
 Geltosky, J.E. Pasadena, California  
 Gerdy, J.R. Carbondale, Illinois  
 Gerresheim, F. Germany, München  
 Gersh, E.S. Pasadena, California  
 Gethmann, R.C. San Diego, California  
 Gibson, J.B. England, Cambridge  
 Gill, R.W. Riverside, California  
 Glass, B. Stony Brook, New York  
 Glätzer, K.H. Germany, Düsseldorf  
 Gloor, H. Switzerland, Geneva  
 Gnes, A. Italy, Padova  
 Gochberg, C. Madison, Wisconsin  
 Godbole, N.N. India, Poona  
 Gold, E.E. Urbana, Illinois  
 Gold, J.R. Davis, California  
 Gonzales, F.W. Upton, New York  
 González, C. Spain, Barcelona  
 González, R. Spain, Barcelona  
 Gorodenski, S.A. Raleigh, North Carolina  
 Gorospe, M. J. Spain, Madrid  
 Gossi, S.J. Pullman, Washington  
 Gottlieb, F.J. Pittsburgh, Pennsylvania  
 Götz, K.G. Germany, Tübingen  
 Gouveia, G. England, Oxford  
 Grabicki, E. New Haven, Connecticut  
 Grace, D. Netherlands, Leiden  
 Graf, U. Switzerland, Zürich  
 Graham, R.S. Chapel Hill, North Carolina  
 Granobles, L.A. Colombia, Bogotá  
 Grant, B.S. Williamsburg, Virginia  
 Gray, K.F. Williamsburg, Virginia  
 Graziano, M.H. Charlottesville, Virginia  
 Green, M.M. Davis, California  
 Greer, G. Australia, Bundoora  
 Greisen, K.S. Pasadena, California  
 Grell, E.H. Oak Ridge, Tennessee  
 Grell, R.F. Oak Ridge, Tennessee  
 Grimes, W.P. Dallas, Texas  
 Groh, G. Germany, Berlin  
 Groot-van Stralen, C. Th. de Netherlands, Leiden  
 Grossfield, J. Lafayette, Indiana  
 Grüneberg, H. England, London  
 Gruwez, G. Belgium, Heverlee  
 Guedes, M.A. Brazil, Pôrto Alegre  
 Guest, W.C. Fayetteville, Arkansas  
 Haapala, O. Finland, Turku  
 Hablas, A.A. U.A.R., Alexandria  
 Hackman, W. Finland, Helsinki  
 Hadorn, E. Switzerland, Zürich  
 Haendle, J. Germany München  
 Hale, G. New York, New York  
 Halfer, C. Milan, Italy  
 Hall, J. Seattle Washington  
 Hall, L. Canada, Vancouver  
 Halm, J. Houston, Texas  
 Hama, H. Japan, Chiba  
 Hamdalla, H. U.A.R. Assuit  
 Hammerschmidt, H. Germany, München  
 Hammond, K. Australia, Sydney  
 Hanks, G.D. Gary, Indiana  
 Hannah-Alava, A. Finland, Turku  
 Hanratty, W.P. Pittsburgh, Pennsylvania  
 Hanson, T.E. Pasadena, California  
 Hanstein, B. Duarte, California  
 Hara, Miss Japan, Fukuoka  
 Hardwick, J.S. Chapel Hill, North Carolina  
 Hardy, D.E. Honolulu, Hawaii  
 Hardy, R.W. San Diego, California  
 Harrison, B. England, Sheffield  
 Harrison, B.J. England, Norwich  
 Harrod, M.J.E. Dallas, Texas  
 Hartley, P. Durham, North Carolina  
 Hartmann-Goldstein, I.J. England, Sheffield  
 Harvey, K. Ithaca, New York  
 Hashem, Y.D. U.A.R., Alexandria  
 Hashim, M. U.A.R. Assuit  
 Hauri, H.-P. Switzerland, Zürich  
 Havermans, E.M.J. Netherlands, Utrecht  
 Hawk, J.A. Lafayette, Indiana  
 Hawkins, E. Stony Brook, New York  
 Hay, D.A. England, Birmingham  
 Hayashi, K. Japan, Osaka  
 Hazevoet, I. Netherlands, Leiden  
 Hebert, P. England, Cambridge  
 Heisenberg, M. Germany, Tübingen  
 Hengstenberg, B. Germany, Tübingen  
 Henstenberg, R. Germany, Tübingen  
 Hennig, I. Germany, Tübingen  
 Hennig, W. Germany, Tübingen  
 Hensen, A.E. Netherlands, Leiden  
 Hess, O. Germany, Düsseldorf  
 Hewitt, N.E. Washington, D.C.  
 Hihara, F. Japan, Tokyo  
 Hihara, Y.K. Japan, Tokyo  
 Hill, C.L. Madison, Wisconsin

Hillman, R. Philadelphia, Pennsylvania  
Hinton, C.W. Wooster, Ohio  
Hippard, J. Los Angeles, California  
Hiraizumi, Y. Austin, Texas  
Hirose, Y. Japan, Kobe  
Hiroyoshi, T. Japan, Osaka  
Hodges, C. Raleigh, North Carolina  
Hodgetts, R.B. Canada, Edmonton  
Hoenigsberg, H.F.B. Colombia, Bogotá  
Hoff, D. Madison, Wisconsin  
Hofman, J.D.D. Netherlands, Haren  
Holden, J. Canada, Vancouver  
Hollander, W.F. Ames, Iowa  
Hollie, E. Austin, Texas  
Hollingsworth, M.J. England, London  
Hollingworth, M. Australia, Brisbane  
Hollis, R.J. Williamsburg, Virginia  
Holzworth, K.W. Pittsburgh, Pennsylvania  
Honda, Y. Japan, Nagasaki  
Hooper, G.B. Poughkeepsie, New York  
Hoste, C. Belgium, Heverlee  
Hotta, Y. Pasadena, California  
Hoyland, M. England, Sheffield  
Hoyt, J.P. Ann Arbor, Michigan  
Huang, S.-L. Austin, Texas  
Huang, S.-M. Austin, Texas  
Hubby, J.L. Chicago, Illinois  
Hubert, L.M. Brazil, Pôrto Alegre  
Hudson, A.S. Chapel Hill, North Carolina  
Hudson, G. England, Cambridge  
Hunt, D.M. England, London  
Hürlimann, R. Switzerland, Zürich  
Huth, A.C. Germany, Tübingen  
Ibrahim, H. U.A.R., Assuit  
Ibrahim, S. U.A.R., Alexandria  
Ikeda, H. Japan, Tokyo  
Ikeda, K. Duarte, California  
Illmensee, K. Germany, München  
Imaizumi, Y. Japan, Chiba  
Inove, A. Japan, Nagasaki  
Ito, S. Japan, Tokyo  
Ivic, G. Yugoslavia, Belgrade  
Iyama, S. Japan, Misima  
Jack, K. Garden City, New York  
Jacobs, M.E. Goshen, Indiana  
Jacobs, P.A. San Diego, California  
Jacobson, A.G. Austin, Texas  
James, J.W. Australia, Sydney  
Janning, W. Germany, Münster  
Jefferson, M. Boulder, Colorado  
Jeliasvčić, B. Ugoslavia, Belgrade  
Jen, K.-d. DeKalb, Illinois  
Jha, A.P. India, Bhagalpur  
Jinks, J.L. England, Birmingham  
Johnsen, R.C. Garden City, New York  
Johnson, F.M. Raleigh, North Carolina  
Johnson, G. Philadelphia, Pennsylvania  
Johnson, J.H. Chicago, Illinois  
Johnson, L. Madison, Wisconsin  
Johnson, T. Northridge, California  
Johnson, W.E. Honolulu, Hawaii  
Johnston, C.J. Pasadena, California  
Johnston, F. Storrs, Connecticut  
Jones, D.A. England, Birmingham  
Jones, G.H. England, Birmingham  
Jones, J.S. Chicago, Illinois  
Jones-Mortimer, M.C. England, Birmingham  
Jonker, F.H. Netherlands, Haren  
Jousset, F.X. France, St. Christol les Alès  
Judd, B.H. Austin, Texas  
Julien, J. Chicago, Illinois  
Jungen, H. Switzerland, Zürich  
Kachur, F. Gary, Indiana  
Kaji, S. Japan, Kobe  
Kalicki, H. Garden City, New York  
Kalisch, W.-E. Davis, California  
Kalisz, A. Brazil, Pôrto Alegre  
Kamra, O.P. Canada, Halifax  
Kanehisa, T. Japan, Kobe  
Kaneshiro, K.Y. Honolulu, Hawaii  
Kang, M.-J. Korea, Seoul  
Kang, S.-H. Notre Dame, Indiana  
Kang, (Song), S.-J. Korea, Seoul  
Kang, Y.S. Korea, Seoul  
Kankel, D.R. Pasadena, California  
Kaplan, M.L. New York, New York  
Kaplan, W.E. Duarte, California  
Karlik, A. Austria, Vienna  
Kastritsis, C.D. Dallas, Texas  
Katt, S. Atlanta, Georgia  
Kaufman, T.C. Canada, Vancouver  
Kaufmann, B.P. Ann Arbor, Michigan  
Kawabe, M. Japan, Kobe  
Kawaharada, M. Japan, Nagasaki  
Kawanishi, M. Japan, Misima  
Kearney, M. England, Sheffield  
Kearsey, M.J. England, Birmingham  
Keith, M.J. Houston, Texas  
Kekić, V. Yugoslavia, Belgrade  
Kellen, C.M. Iowa City, Iowa  
Keller, E.C. Morgantown, West Virginia  
Keller, H.E. Morgantown, West Virginia  
Kelly, J.F. Canton, New York  
Kelly, M. Australia, Brisbane  
Kercher, M.D. Lexington, Kentucky  
Kernaghan, R.P. Stony Brook, New York  
Kersten, H.J.M.G. Netherlands, Utrecht  
Kessenich, M. Madison, Wisconsin  
Kessler, S. Madison, Wisconsin  
Khishin, A. U.A.R., Assuit  
Kieft, P. Netherlands, Leiden  
Kikkawa, H. Japan, Osaka  
Kim, K.-J. Korea, Seoul  
Kim, K.W. Korea, Kwangju  
Kimura, M. Japan, Misima  
King, R.C. Evanston, Illinois  
Kiriasis, W.C. Upton, New York  
Kirschbaum, W.F. Argentina, Buenos Aires  
Kitagawa, O. Japan, Tokyo  
Kitazume, Y. Japan, Kobe  
Kitos, R. Syracuse, New York  
Kleisch, U. Germany, Berlin

- Klug, M. Germany, Freiburg  
 Knaap, A.G.A.C. Netherlands, Leiden  
 Kobel, H.R. Switzerland, Geneva  
 Koepfer, R. New York, New York  
 Köhler, B. Germany, Tübingen  
 Kojima, K.-I. Austin, Texas  
 Kolodzieg, G. Germany, Freiburg  
 Komori, M. Japan, Nagasaki  
 Konopka, R.J. Pasadena, California  
 Kooistra, J. Netherlands, Haren  
 Koref-Santibanez S. New York, New York  
 Korge, G. Germany, München  
 Korinek, E. Canada, Vancouver  
 Kosuda, K. Japan, Tokyo  
 Kothari, R.M. India, Poona  
 Kovarik, A. Austin, Texas  
 Kramer, P.G. Netherlands, Leiden  
 Kratz, F.L. Brazil, Pôrto Alegre  
 Krause, E. South Orange, New Jersey  
 Kreisman, D.J. Pullman, Washington  
 Kress, H. Germany, München  
 Kreteski, D. Atlanta, Georgia  
 Krimbas, C. Greece, Athens  
 Krivshenko, J.D. Rochester, New York  
 Krunić, M. Yugoslavia, Belgrade  
 Kubli, E. Switzerland, Zürich  
 Kucherlapati, R.S. Urbana, Illinois  
 Kuchino, C. Japan, Misima  
 Kunze-Mühl, E. Austria, Vienna  
 Kuroda, Y. Japan, Misima  
 Kurokawa, H. Japan, Tokyo  
 Label, E. France, Gif-sur-Yvette  
 Lachaise, D. France, Gif-sur-Yvette  
 Lai, Y.-H. Pasadena, California  
 Laird, C. Seattle, Washington  
 Lakovaara, S. Finland, Helsinki  
 Lamb, M.J. England, London  
 Landner, L. Sweden, Stockholm  
 Lang, E.L. Carban, Illinois  
 Langley, C. Madison, Wisconsin  
 Langlois, B. France, Gif-sur-Yvette  
 LaPushin, R. Houston, Texas  
 Lasa, L. Spain, Madrid  
 Lasseter, A. Austin, Texas  
 Latter, B.D.H. Australia, Sydney  
 Laughnan, J.R. Urbana, Illinois  
 Laurent, J. France, Gif-sur-Yvette  
 Lawrence, M.J. England, Birmingham  
 Lebherz, H. Switzerland, Zürich  
 Lechien, J. Belgium, Namur  
 Lee, B.W. Korea, Chungang  
 Lee, C.C. Korea, Seoul  
 Lee, C.S. Korea, Chungang  
 Lee, J.-Y. Pasadena, California  
 Lee, K.-S. Korea, Seoul  
 Lee, L.Y. Lafayette, Indiana  
 Lee, T.J. Korea, Chungang  
 Lees, G. Duarte, California  
 Lefevre, G. Northridge, California  
 Leffel, L.E. Madison, Wisconsin  
 Legay, J.M. France, Lyon  
 Leightwood, J. Australia, Canberra  
 Lemeunier, F. France, Gif-sur-Yvette  
 Leonard, J. Purchase, New York  
 Leto, G. DeKalb, Illinois  
 Levine, H. Storrs, Connecticut  
 Levins, R. Chicago, Illinois  
 Lewandowski, Y. Buffalo, New York  
 Lewellyn, S. Australia, Canberra  
 Lewis, E.B. Pasadena, California  
 Lewis, J. Pasadena, California  
 Lewontin, R.C. Chicago, Illinois  
 Lezzi, M. Switzerland, Zürich  
 L'Helias, C. France, Gif-sur-Yvette  
 L'Héritier, Ph. France, Clermont-Ferrand  
 Libion, M. Belgium, Namur  
 Lidington, K. Lafayette, Indiana  
 Lieb, E. Germany, München  
 Lifschytz, E. San Diego, California  
 Limbird, D. Washington, D.C.  
 Lindsley, D.L. San Diego, California  
 Linney, R. England, Birmingham  
 Lints, C. Belgium, Heverlee  
 Lints, F.A. Belgium, Heverlee  
 Lipps, K. Davis, California  
 Liszczynskyj, J. Utica, New York  
 Llewellyn, M.A. Bellingham, Washington  
 Lloyd, B. Australia, Adelaide  
 LoCascio, H. Buffalo, New York  
 Lokki, J. Finland, Helsinki  
 Lommerse, M.A.H. Netherlands, Leiden  
 Long, G. Chapel Hill, North Carolina  
 Long, T.C. Chapel Hill, North Carolina  
 Loos, M.J. Netherlands, Leiden  
 Louis, M. Grance, Gif-sur-Yvette  
 Loukas, M. Greece, Athens  
 Lowy, P.H. Pasadena, California  
 Lu, C.C. Houston, Texas  
 Lu, M.-H. Austin, Texas  
 Lucas, K. New Haven, Connecticut  
 Lucchesi, J.C. Chapel Hill, North Carolina  
 Luce, W.M. Urbana, Illinois  
 Ludwig, M.R. Brazil, Pôrto Alegre  
 Lüers, H. Germany, Germany  
 Lumme, J. Finland, Helsinki  
 Luna, E. San Marcos, Texas  
 Lundelius, J. Austin, Texas  
 Lüning, K.G. Sweden, Stockholm  
 Lütolf, H.-U. Switzerland, Zürich  
 Lyttle, T.W. Madison, Wisconsin  
 Macaulay, S. Canada, Vancouver  
 MacBean, I.T. Australia, Bundoor  
 Machado, D.M. Brazil, Pôrto Alegre  
 Machida, I. Japan, Chiba  
 Machová, H. Czechoslovakia, Brno  
 Macht, V.C. Notre Dame, Indiana  
 MacPhail, S. Corvallis, Oregon  
 Madhaven, K. Switzerland, Zürich  
 Maeda, M. Japan, Misima  
 Maeda, Y. Japan, Kobe  
 Magalhães, L.E. de Brazil, São Paulo  
 Magnusson, J. Sweden, Stockholm

- Maier, P. Switzerland, Zürich  
Maier, V. Switzerland, Zürich  
Mainx, F. Austria, Vienna  
Maitra, S.N. India, Calcutta  
Majoral, J. Spain, Barcelona  
Mange, A. Seattle, Washington  
Manna, P.K. India, Calcutta  
Marien, D. New York, New York  
Marinković, D. Yugoslavia  
Markowitz, E.H. Madison, Wisconsin  
Maroni, G. Madison, Wisconsin  
Marques, E.K. Brazil, Pôrto Algere  
Marree, C.M. Netherlands, Utrecht  
Marsh, D. Tuscaloosa, Alabama  
Martensen, D. Galesburg, Illinois  
Marstokk, A. Norway, Oslo  
Martin, A.O. Cleveland, Ohio  
Martínez, J.M. Spain, Barcelona  
Martinez, M.N. Brazil, Pôrto Alegre  
Martinsen, D. Davis, California  
Maruyama, T. Japan, Misima  
Marynick, S.P. Dallas, Texas  
Masry, A.M. U.A.R., Alexandria  
Massie, H.R. Utica, New York  
Masterson, J.E. Ames, Iowa  
Masuda, H. Japan, Misima  
Mather, W.B. Australia, Brisbane  
Matheson, A.C. Australia, Bundoora  
Matsubara, T. Japan, Nagasaki  
Mayoh, H. Canada, Vancouver  
Mazar Barnett, B. Argentina, Buenos Aires  
McAdams, L.W. Raleigh, North Carolina  
McCabe, K. Australia, Brisbane  
McCaman, R. Duarte, California  
McCarron, M. Storrs, Connecticut  
McCormack, M.K. South Orange, New Jersey  
McCrady, E. Greensboro, North Carolina  
McCaman, M.W. Duarte, California  
McDonald, R.P. England, London  
McDonnell, J. Garden City, New York  
McFarland, J.L. Riverside, California  
McKenzie, G. Scotland, Aberdeen  
McKenzie, J.A. Australia, Bundoora  
McLean, J.L. Chapel Hill, North Carolina  
McMahon, M. Utica, New York  
McMurtrey, M. Houston, Texas  
McNeil, H.M. Dallas, Texas  
McNeil, R.M. Raleigh, North Carolina  
McQue, R. Flagstaff, Arizona  
Meeles, E. Netherlands, Haren  
Meer, B. Germany, Tübingen  
Megaheid, M.A. U.A.R., Assuit  
Megna, F. Italy, Naples  
Melon, I. Italy, Naples  
Meltzer, P.S. Pasadena, California  
Memhauser, I. New York, New York  
Ménsua, J.L. Spain, Barcelona  
Mercader, J. Mexico, Mexico City  
Mercio, A.L. Brazil, Pôrto Alegre  
Merrell, D.J. Minneapolis, Minnesota  
Merriam, J.R. Los Angeles, California  
Merritt, R. Rochester, New York  
Messerschmid, V. Germany, München  
Metcalf, J.A. England, Heslington  
Mettler, L.E. Raleigh, North Carolina  
Meyer, G.F. Germany, Tübingen  
Meyer, H.U. Madison, Wisconsin  
Micheli, A. Italy, Rome  
Miklos, G.L. San Diego, California  
Mikušová, J. Netherlands, Utrecht  
Milkman, R. Iowa City, Iowa  
Miller, D.D. Lincoln, Nebraska  
Miller, D.H. Australia, Sydney  
Miller, S. Seattle, Washington  
Millington-Ward, A.M. Netherlands, Leiden  
Milton, M.K. Australia, Sydney  
Minato, K. Japan, Misima  
Mindek, Géza Switzerland, Zürich  
Mishra, M. India, Bhagalpur  
Mitchell, H.K. Pasadena, California  
Mitchell, J.P. Tuscaloosa, Alabama  
Mittra, N. India, Calcutta  
Mittler, S. DeKalb, Illinois  
Mizobuchi, K. Japan, Chiba  
Mizuguchi, Y. Brazil, São Paulo  
Moffitt, S. England, Oxford  
Mohler, J. Iowa City, Iowa  
Mohler, J.D. Iowa City, Iowa  
Mohr, M.A. Houston, Texas  
Moisand, R.E. Buffalo, New York  
Mollet, P. Switzerland, Zürich  
Monclús, M. Spain, Barcelona  
Monjeló, L.A. dos S. Brazil, Pôrto Alegre  
Montalenti, G. Italy, Rome  
Montana, D. Syracuse, New York  
Montelius, I. Sweden, Stockholm  
Montgomery, S.M. Honolulu, Hawaii  
Morales, N.B. Brazil, Pôrto Alegre  
Moran, C. Corvallis, Oregon  
Morata, G. Spain, Madrid  
Moree, R. Pullman, Washington  
Moreno, N.I. Colombia, Bogotá  
Mori, S. Japan, Nagasaki  
Moriwaki, D. Japan, Misima  
Morrow, D. Canada, Calgary  
Mortensen, M. Denmark, Copenhagen  
Morton, M.D. Charlottesville, Virginia  
Moskwinski, T. Notre Dame, Indiana  
Mosna, G. Italy, Milan  
Mossige, J. Norway, Oslo  
Moth, J.J. Australia, Sydney  
Moura, V.L.P. de Brazil, Pôrto Alegre  
Mourad, A.M. U.A.R., Alexandria  
Mueller, D. Iowa City, Iowa  
Mukai, T. Raleigh, North Carolina  
Mukherjee, A.S. India, Calcutta  
Mukherjee, T.K. India, Calcutta  
Mukherjee, U. Austria, Vienna  
Mulley, J.C. Australia, Sydney  
Munoz, E. Netherlands, Leiden  
Muñoz, E.R. Argentina, Buenos Aires  
Murakami, A. Japan, Misima

Murata, M. Japan, Chiba  
Murch, A. Australia, Adelaide  
Murnik, M.R. Illinois, Macomb  
Muzyka, G. Galesburg, Illinois  
Myszewski, M.E. Des Moines, Iowa  
Nagib, F.M. U.A.R., Alexandria  
Naguib, F.N.M. Canada, Edmonton  
Nahas, P. Notre Dame, Indiana  
Nakai, S. Japan, Osaka  
Nakai, S. Japan, Chiba  
Nakashima-Tanaka, E. Japan, Sakai  
Narise, S. Japan, Sakado  
Narise, T. Japan, Sadado-Machi  
Nash, D. Canada, Edmonton  
Nash, W.G. Washington, D.C.  
Natori, S. New Haven, Connecticut  
Nawa, S. Japan, Misima  
Nayudu, P.L. Australia, Clayton  
Neeley, J.C. Portland, Oregon  
Neto, C.C. Brazil, Pôrto Alegre  
Nettleton, R.W. Lincoln, Nebraska  
Nevin, E. Gary, Indiana  
Newton, J.A. Chapel Hill, North Carolina  
Nicoletti, B. Italy, Rome  
Miet, J.P. van der Netherlands, Leiden  
Nigon, V. France, Lyon  
Nill, A. Austin, Texas  
Nilsson, B. Sweden, Stockholm  
Nishiura, J.T. Seattle, Washington  
Nix, C.F. Oak Ridge, Tennessee  
Noble, W.R. Raleigh, North Carolina  
Nogués, R. Spain, Barcelona  
Norby, S. Denmark, Copenhagen  
Norton, S. Charlottesville, Virginia  
Nöthel, H. Germany, Berlin  
Nöthiger, R. Switzerland, Zürich  
Novitski, E. Eugene, Oregon  
O'Donald, P. England, Cambridge  
Oftedal, P. Norway, Oslo  
Ogaki, M. Japan, Sakai  
Ogita, Z. Japan, Osaka  
Oguma, Y. Japan, Tokyo  
Ohanessian, A. France, Gif-sur-Yvette  
Ohba, S. Japan, Tokyo  
Ohki, K. Japan, Nagasaki  
Ohnishi, O. Madison, Wisconsin  
Ohta, T. Japan, Misima  
Oishi, K. Japan, Misima  
Ojeda, A.A. Colombia, Bogotá  
Okada, T. Japan, Tokyo  
Oksala, T.A. Finland, Turku  
Okubo, K. Japan, Nagasaki  
Olive, M. Houston, Texas  
Oliver, D. Iowa City, Iowa  
Oliver, D. Austin, Texas  
Olivieri, G. Italy, Rome  
Olthoff, H.M. Netherlands, Haren  
Olvera, O. Mexico, Mexico City  
Ondřej, M. Czechoslovakia, Prague  
Ortiz, E. Spain, Madrid  
Oshima, C. Japan, Misima  
Outwater, T.W. Chapel Hill, North Carolina  
Ouweneel, W.J. Netherlands, Utrecht  
Paez, D. Colombia, Bogotá  
Paik, S.K. Korea, Seoul  
Paik, Y.K. Honolulu, Hawaii  
Paika, I.J. Lincoln, Nebraska  
Pak, W.L. Lafayette, Indiana  
Pakonen, C.Z. Pullman, Washington  
Pal, T. India, Calcutta  
Pankow, W. Switzerland, Zürich  
Parisi, G. Italy, Naples  
Park, E.H. Korea, Seoul  
Park, M.S. Korea, Kwangju  
Parker, D.R. Riverside, California  
Parker, M. Houston, Texas  
Parry, D. Seattle, Washington  
Parshad, R. India, Chandigarh  
Parsons, P.A. Australia, Bundoora  
Parzen, S.D. Madison, Wisconsin  
Pasteur, G.D. Dallas, Texas  
Pasteur, N. Dallas, Texas  
Patricolo, M.R. Italy, Naples  
Patrouli, H. Greece, Athens  
Patty, R.A. Lincoln, Nebraska  
Pavlich, L. Flagstaff, Arizona  
Pavlovsky, O. Davis, California  
Paxman, G.J. England, Lancaster  
Paz, C. Argentina, Buenos Aires  
Peacock, W.J. Australia, Canberra  
Peeding, N. Durham, North Carolina  
Peers, E. New York, New York  
Pelecanos, M. Greece, Patras  
Perdrix-Gillot, S. France, Lyon  
Perekovic, V. Canada, Edmonton  
Pereyra, E. Argentina, Buenos Aires  
Perkins, J.M. England, Birmingham  
Perreault, W.J. Ann Arbor, Michigan  
Peterson, K. Northridge, California  
Petković, D. Yugoslavia, Belgrade  
Pex, A.M. Netherlands, Leiden  
Picard, G. France, Clermont-Ferrand  
Piekieniak, M. Utica, New York  
Pierre, A.M. France, Gif-sur-Yvette  
Pinsker, W. Austria, Vienna  
Pipkin, S.B. Washington, D.C.  
Phelps, C.G. Chapel Hill, North Carolina  
Phillips, G. Austin, Texas  
Phillips, J. Austin, Texas  
Plagens, U. Madison, Wisconsin  
Plaut, W.S. Madison, Wisconsin  
Plus, N. France, St. Christol les Alès  
Poirier, M. Durham, North Carolina  
Polan, M.L. New Haven, Connecticut  
Pomato, N.J. Notre Dame, Indiana  
Poodry, C. Canada, Vancouver  
Pospisil, Z. Eugene, Oregon  
Postlethwait, J.H. Eugene, Oregon  
Porter, H.N. Austin, Texas  
Portin, P. Finland, Turku  
Posch, N.A. Pasadena, California  
Poulson, D.F. New Haven, Connecticut



- Powell, J.R. New York, New York  
Powers, L.M. Washington, D.C.  
Prakash, S. Rochester, New York  
Prendergast, L. San Diego, California  
Prevosti, A. Spain, Barcelona  
Prins, F.W. Netherlands, Haren  
Printz, P. France, Gif-sur-Yvette  
Procunior, D. Canada, Calgary  
Pronk, P. Netherlands, Haren  
Prout, T. Riverside, California  
Prudhommeau, C. Netherlands, Leiden  
Pulliam, R. Chicago, Illinois  
Puro, J. Finland, Turku  
Pyati, J. Boulder, Colorado  
Querubin, M.A. Brazil, São Paulo  
Räber, E. Switzerland, Zürich  
Rahman, R. India, Calcutta  
Rahman, S.M.Z. India, Bhagalpur  
Raibley, D.W. Carbondale, Illinois  
Raichaudhuri, A. India, Calcutta  
Raikow, R. Honolulu, Hawaii  
Rajaraman, R. Canada, Halifax  
Ramel, C. Sweden, Stockholm  
Ramila, D. Brazil, Pôrto Alegre  
Rankin, S. Bellingham, Washington  
Rathie, K.A. Australia, Sydney  
Ratty, F.J. San Diego, California  
Rawls, J.M. Chapel Hill, North Carolina  
Ray, C. Atlanta, Georgia  
Razzini, A. Italy, Milan  
Reguly, M.L. Brazil, Pôrto Alegre  
Relichová, J. Czechoslovakia, Brno  
Relton, J.M. England, Sheffield  
Remondini, D.J. Spokane, Washington  
Rendel, J.M. Australia, Sydney  
Renka, M.M. Austin, Texas  
Resch, K.M. Austin, Texas  
Reveley, M.A. Austin, Texas  
Rha, C.H. Korea, Kwangju  
Rhodes, K. Atlanta, Georgia  
Ribó, G. Spain, Barcelona  
Rice, T. New Haven, Connecticut  
Richard-Molard, C. France, Gif-sur-Yvette  
Richardson, M. Austin, Texas  
Richardson, R.H. Austin, Texas  
Richmond, R.C. Bloomington, Indiana  
Rickoll, W. Durham, North Carolina  
Rijnsburger, T. Netherlands, Leiden  
Rinehart, R.R. San Diego, California  
Ringo, J.M. Davis, California  
Ripoll, P. Spain, Madrid  
Ristow, H. New Haven, Connecticut  
Rivera, M.L. Spain, Barcelona  
Rizki, R.M. Ann Arbor, Michigan  
Rizki, T.M. Ann Arbor, Michigan  
Roach, S. Galesburg, Illinois  
Robbins, L.G. Austin, Texas  
Roberts, C.F. England, Leicester  
Roberts, D.B. England, Oxford  
Roberts, M.A. Upton, New York  
Roberts, P. Corvallis, Oregon  
Roberts, R. Chicago, Illinois  
Robertson, F.W. Scotland, Aberdeen  
Robertson, G.C. Raleigh, North Carolina  
Robertson, K. Houston, Texas  
Rocha, D. San Marcos, Texas  
Rodell, C.F. Minneapolis, Minnesota  
Rodinò, E.B. Italy, Padova  
Rokop, S. San Diego, California  
Romans, P. Canada, Edmonton  
Rose, R.W. Glenside, Pennsylvania  
Rosenbluth, R. Canada, Vancouver  
Rosenfeld, A. Seattle, Washington  
Rotter, D. DeKalb, Illinois  
Rowe, L. Riverside, California  
Roy, D. Atlanta, Georgia  
Ruch, P. Switzerland, Zürich  
Ruderer-Doschek, E. Austria, Vienna  
Rumball, W. Australia, Sydney  
Runger, E. Switzerland, Geneva  
Russell, M.R. Canada, Edmonton  
Rutherford, P. Scotland, Aberdeen  
Saeki, T. Japan, Chiba  
Saitta, F. Austin, Texas  
Sakaguchi, B. Japan, Fukuoka  
Sakai, K.I. Japan, Misima  
Sakoyama, Y. Japan, Osaka  
Sakri, B. Lafayette, Indiana  
Salas, E. Spain, Madrid  
Saleh, F.M. U.A.R., Assuit  
Sallam, T.M. U.A.R., Assuit  
Salverson, H.M. Madison, Wisconsin  
Samis, H.V. Utica, New York  
Sanches, F. Netherlands, Leiden  
Sandler, L.M. Seattle, Washington  
Sang, J.H. England, Brighton  
Sankaranarayanan, K. Netherlands, Leiden  
Santamaría, P. Spain, Madrid  
Santos, E.P. dos Brazil, São Paulo  
Sanyal, C. India, Calcutta  
Sato, J.E. Honolulu, Hawaii  
Saul, S. Chicago, Illinois  
Saura, A. Finland, Helsinki  
Savolainen, S. Finland, Turku  
Savontaus, M.-L. Finland, Turku  
Sayers, E.R. Tuscaloosa, Alabama  
Sayles, C. Canada, Calgary  
Schaerer, H.-R. Switzerland, Zürich  
Schäfer, U. Germany, Düsseldorf  
Schaffer, H.E. Raleigh, North Carolina  
Schalet, A. Netherlands, Leiden  
Scharloo, W. Netherlands, Utrecht  
Scheid, W. Germany, Münster  
Scheidt, G.C. Pasadena, California  
Schewe, M. Canada, Vancouver  
Schneider, I. Washington, D.C.  
Schouten, S.C.M. Netherlands, Utrecht  
Schümperli, R. Switzerland, Zürich  
Schüpbach, P. Switzerland, Zürich  
Schultz, E.G. Brazil, Pôrto Alegre  
Schweizer, P. Switzerland, Zürich  
Schwochau, M. Germany, Düsseldorf

- Sciandra, R. DeKalb, Illinois  
 Scowcroft, W.R. Australia, Canberra  
 Seecof, R.L. Duarte, California  
 Seki, T. Japan, Osaka  
 Semeonoff, R. England, Leicester  
 Sene, F.M. Brazil, São Paulo  
 Serizawa, S. Japan, Misima  
 Settegast, M. Houston, Texas  
 Sevela, A. Czechoslovakia, Brno  
 Sewell, D. England, Sheffield  
 Seybold, W.D. Pasadena, California  
 Seyffert, W. Germany, Tübingen  
 Sfier, G. Utica, New York  
 Shafer, G.T. Philadelphia, Pennsylvania  
 Shafer, S.J. Philadelphia, Pennsylvania  
 Shannon, M.P. Austin, Texas  
 Sharma, G.P. India, Chandigarh  
 Shearn, A. New Haven, Connecticut  
 Sheehy, A.J. Australia, Clayton  
 Sheldon, B.L. Australia, Sydney  
 Shellenbarger, D. Iowa City, Iowa  
 Shen, M.W. Austin, Texas  
 Sherald, A. Charlottesville, Virginia  
 Sherif, T.H. U.A.R., Assuit  
 Sherwin, R.N. Chicago, Illinois  
 Shideler, D.M. Lafayette, Indiana  
 Shield, G. England, Brighton  
 Shiomi, T. Japan, Nagasaki  
 Shmookler, R. England, Brighton  
 Shoeb, Y. U.A.R. Alexandria  
 Shorrocks, B. England, Leeds  
 Sick, K. Denmark, Copenhagen  
 Siddiqi, O. Pasadena, California  
 Sieber, F. Switzerland, Zürich  
 Sillans, D. France, Lyon  
 Silva, I.F. da Brazil, Pôrto Alegre  
 Silver, E. Philadelphia, Pennsylvania  
 Simmons, M.J. Madison, Wisconsin  
 Simpson, A. Gary, Indiana  
 Singeisen, C. Switzerland, Zürich  
 Singh, A. India, Chandigarh  
 Singh, V.K. India, Bhagalpur  
 Sistonon, P. Finland, Helsinki  
 Sladká, D. Czechoslovakia, Brno  
 Slati, H.M. East Lansing, Michigan  
 Slepkeis, N.J. Madison, Wisconsin  
 Sloane, C.A. Chicago, Illinois  
 Smit, A.M. Pasadena, California  
 Smith, D.A. England, Birmingham  
 Smith, J.M. England, Brighton  
 Smith, L.M. Portland, Oregon  
 Smith, L.M. Pittsburgh, Pennsylvania  
 Smith, P. Atlanta, Georgia  
 Smith, P.D. Pittsburgh, Pennsylvania  
 Smouse, P. Austin, Texas  
 Smurtleff, E. Corvallis, Oregon  
 Snook, M. Australia, Canberra  
 Synder, A.G. Williamsburg, Virginia  
 Synder, R.D. Atlanta, Georgia  
 Sobels, F.H. Netherlands, Leiden  
 Sokoloff, A. San Bernardino, California  
 Somero, M.G. San Diego, California  
 Sondhi, G. Edison, New York  
 Sondhi, K.C. Edison, New York  
 Song, C.Y. Korea, Chungang  
 Sorsa, M. Finland, Helsinki  
 Sorsa, V. Finland, Helsinki  
 Souza, H.M.L. de Brazil, São Paulo  
 Sparrow, J. England, Brighton  
 Spassky, B. Davis, California  
 Spear, B. New Haven, Connecticut  
 Sperlich, D. Germany, Tübingen  
 Spiess, E.B. Chicago, Illinois  
 Spieth, H.T. Davis, California  
 Spillmann-Faller, E. Switzerland, Zürich  
 Spofford, J.B. Chicago, Illinois  
 Sprague, G. New Haven, Connecticut  
 Sprechman, L. Austin, Texas  
 Springer, R. Austria, Vienna  
 Spuhler, P. Boulder, Colorado  
 Stafford, D.W. Chapel Hill, North Carolina  
 Stahl, G. Sweden, Stockholm  
 Stankevych, A.J. Chicago, Illinois  
 Stattard, R. Atlanta, Georgia  
 Steen, L. van der Netherlands, Leiden  
 Steffensen, D.M. Urbana, Ill.  
 Steiner, E. Switzerland, Zürich  
 Steiner, W.W. Honolulu, Hawaii  
 Stennek, A. Sweden, Stockholm  
 Stern, C. Berkeley, California  
 Stewart, B. Los Angeles, California  
 Stocker, A.J. Dallas, Texas  
 Stoddard, A. Pittsburgh, Pennsylvania  
 Stoliar, M.A. Argentina, Buenos Aires  
 Stoltz, J. Austin, Texas  
 Střebická, M. Czechoslovakia, Brno  
 Strickberger, M.W. St. Louis, Missouri  
 Stroman, P. Denmark, Copenhagen  
 Strub, S. Switzerland, Zürich  
 Stunell, M. Houston, Texas  
 Subbarao, S.K. Urbana, Illinois  
 Sullivan, D.T. Syracuse, New York  
 Sullivan, M.C. Syracuse, New York  
 Sunbom, R. Northridge, California  
 Sung, K.C. Honolulu, Hawaii  
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 Surver, W.M. Notre Dame, Indiana  
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 Suzuki, D.T. Canada, Vancouver  
 Suzuki, H. Japan, Chiba  
 Swartz, P. Austin, Texas  
 Szymanski, D. Gary, Indiana  
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 Thieker, K. Iowa City, Iowa  
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 Tobari, Y.N. Japan, Tokyo  
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