

The activity in the carcass and head of adults very likely corresponds to what is found in the dispersed fat tissue; thus, this tissue probably accounts for most of the ADH found in imagos.

Figure 2 shows the developmental pattern of accumulation of active enzyme in some of these organs. Of the organs examined, only the Malpighian tubules persist from larval to adult stages; it is interesting to notice that they do not display the general rise in activity during the third instar but they do so in the adult.

References: Ursprung et al. 1970; W.Roux Archiv. 164:201-208; Maroni 1978, Biochem. Genet. 16:509-523; Maroni et al. 1982, Genetics 101:431-446.

Martinez-Sebastian, M.J. & R.deFrutos.
University of Valencia, Spain. Chromosomal polymorphism in Drosophila subobscura populations submitted to selection for a quantitative character.

Table 1. Frequencies of chromosomal arrangements (%) in the laboratory population.

Capture	3/79 n=118	6/79 n=156	11/80 n=109	12/81 n=145
A _{st}	38.36	45.65	15.95	16.1
A ₁	4.11	6.52	5.08	4.3
A ₂	57.53	47.83	77.97	78.5
A ₁₊₂	-	-	-	1.1
J _{st}	27.97	24.34	16.51	16.0
J ₁	72.03	75.66	83.49	84.0
U _{st}	4.31	0.66	-	-
U ₁₊₂	43.97	47.37	45.37	41.3
E _{st}	34.75	46.05	47.66	40.5
E ₁₊₂	10.17	11.19	8.41	2.1
E ₁₊₂₊₉	18.64	9.21	16.82	16.1
E ₁₊₂₊₉₊₁₂	29.66	27.63	26.17	41.3
E ₁₊₂₊₉₊₃	5.09	3.95	0.93	-
E ₈	1.69	1.97	-	-
O _{st}	20.34	21.79	33.94	22.1
O ₃₊₄	27.12	24.36	45.87	72.4
O ₃₊₄₊₇	32.20	25.64	14.68	2.75
O ₃₊₄₊₈	5.09	19.87	1.83	-
O ₃₊₄₊₂	1.69	4.49	1.83	2.75
O ₃₊₄₊₂₂	0.85	0.64	-	-
O ₃₊₄₊₁	11.02	-	1.83	-
O ₇	1.60	3.21	-	-

The evolution of chromosomal polymorphism in several abdominal bristle selection lines was analyzed.

A laboratory population (R) was established with individuals from nature, and was developed without selection throughout the entire duration of the experiment. Four selection lines, two high (P₁ and P₂) and two low (N₁ and N₂), and two control lines (C₁ and C₂) were taken from the laboratory population and run during 24 generations.

The sum of the bristles on the 4th and 5th abdominal sternites was the criterion of selection and the intensity of selection used was 20%.

The first time the chromosomal polymorphism of the natural population was analyzed, and later, periodic analyses of the selection lines, control lines and laboratory population were carried out.

The results of the analyses of the laboratory population are given in Table 1. As it can be seen, the chromosomal arrangements present in the initial population at a low frequency tend to be eliminated. A χ^2 homogeneity test comparing the different analyses shows no significant differences in chromosome U ($\chi^2=1.29$; d.f.=3; P=0.73). However, chromosomes J ($\chi^2=7.91$; d.f.=3; P=0.05), A ($\chi^2=27.59$; d.f.=3; P<0.001), E ($\chi^2=48.99$; d.f.=9; P<0.001) and O ($\chi^2=133.49$; d.f.=9; P<0.001) show clear differences. In A chromosome the A₂ arrangement tends to be selected, in E chromosome the E_{st} and E₁₊₂₊₉₊₁₂ arrangements increase and in O chromosome the O₃₊₄ arrangement increases strongly.

Selection lines and control lines were analyzed several times during the experiment, but in this paper we give only the results of the last analysis, which was done in the 24th generation of selection. The results (see Table 2) show a general tendency to homozygosis, stronger in selection lines than in control lines.

In chromosome A, all lines except C₁ are practically homozygotic. In the two high selection lines the A₂ arrangement is fixed, but in low selection lines the A₂ is fixed in N₁ and

Table 2. Frequencies of chromosomal arrangements (%) in selection lines.

	R n=157	P1 n=128	P2 n=89	N1 n=116	N2 n=145	C1 n=126	C2 n=149
A _{st}	45.65	1.3	-	-	98.75	52.1	-
A ₁	7.52	-	-	-	-	-	-
A ₂	47.83	98.7	100.0	100.0	1.25	47.9	100.0
J _{st}	24.34	0.8	-	-	33.79	9.5	-
J ₁	75.66	99.2	100.0	100.0	66.21	90.5	100.0
U _{st}	0.66	-	-	-	-	-	-
U ₁₊₂	47.37	100.0	100.0	-	72.73	60.0	48.30
U ₁₊₂₊₈	51.97	-	-	100.0	27.27	40.0	51.70
E _{st}	46.05	16.4	51.69	-	43.75	46.0	23.81
E ₁₊₂	11.19	-	1.12	73.3	24.31	2.4	12.93
E ₁₊₂₊₉	9.21	-	-	-	-	-	-
E ₁₊₂₊₉₊₁₂	27.63	83.6	47.19	26.7	31.94	51.6	63.27
E ₁₊₂₊₉₊₃	3.95	-	-	-	-	-	-
E ₈	1.97	-	-	-	-	-	-
O _{st}	21.79	-	56.82	-	-	58.1	32.21
O ₃₊₄	24.36	-	43.18	100.0	100.0	17.7	67.11
O ₃₊₄₊₇	25.64	100.0	-	-	-	24.2	-
O ₃₊₄₊₈	19.87	-	-	-	-	-	-
O ₃₊₄₊₂	4.49	-	-	-	-	-	0.67
O ₃₊₄₊₂₂	0.64	-	-	-	-	-	-
o ₇	3.21	-	-	-	-	-	-

In some chromosomes the same arrangement was fixed in the two high selection lines or in the two low selection lines, as happens with the A₂ arrangement in the two high selection lines and with the O₃₊₄ arrangement in the two low selection lines. These two arrangements tend to be increased in the laboratory population. Also the J₁ and U₁₊₂ arrangements were fixed in high selection lines. In the rest of the chromosomes the behaviour of high and low selection lines is similar.

Martínez-Sebastián, M.J. & J.L. Mensua. University of Valencia, Spain. Variations of wing dimensions in *Drosophila subobscura* populations selected for abdominal bristle number.

laboratory population. At 17th generation of selection, wing length and wing width were measured.

Table shows the means of wing dimensions in the laboratory population, control lines and selection lines.

Significant differences exist between the laboratory population and the selection and control lines (except wing length males R versus males C₁). Also there are significant

A_{st} is fixed in N₂. In chromosome J, frequency of the J₁ arrangement reaches values close to 100% in all selection lines, except in N₂ where J₁ arrangement shows smaller values than in the initial population.

As in A and J chromosomes, in U chromosome the same arrangement (U₁₊₂) is fixed in the two high selection lines and N₂ shows a different behaviour to N₁. In control lines the two most frequent arrangements (U₁₊₂ and U₁₊₂₊₈) reach values close to 50%.

In chromosome E, the most polymorphic, in no case was homozygosity reached. A different arrangement tends to be increased in each line.

In chromosome O which shows a great number of gene arrangements, homozygosity was reached in several selection lines. The two low selection lines are homozygotic for the O₃₊₄ arrangement and P₁ for O₃₊₄₊₇ arrangement. The rest of the lines remained polymorphic.

On comparing the chromosomal frequencies in the initial population with the frequencies in the selection lines and control lines after 24 generations of artificial selection, it can be seen that the two high selection lines and one low selection line (N₁) tend to reach homozygosity while the control lines and N₂ low line tend to remain polymorphic.

In a laboratory population (R) of *D. subobscura*, the characters of wing length and wing width were measured.

Two replicate selection lines for abdominal bristle number in both, high (P₁ and P₂) and low (N₁ and N₂), directions and two control lines (C₁ and C₂) were established from the