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 USNA. Mapping of *tuh-3* between *pbx* and
fl in *D.melanogaster*.

The tumorous-head mutant gene (*tuh-3*) was mapped in partial aneuploids and by recombination studies distal to *pbx* (3.58.8+) and presumably is one of the most posterior bi-thorax-complex genes (Kuhn et al. 1981; Kuhn and Woods 1982). These studies did not tell us how distal *tuh-3* was to *pbx*. We report here the results of our mapping *tuh-3* between

postbithorax (*pbx*) and fluted (*fl*). We mapped *fl* to 3-59.7+. Female tumorous-head flies were mated to *sbd*² *bx*³ *pbx fl*/TMI males. F₁ females were backcrossed to *sbd*² *bx*³ *pbx fl*/TMI males. Male offspring that were *fl* only were isolated for further testing. Twenty males were shown to carry a 3rd chromosome in which a cross-over occurred somewhere between *pbx* and *fl*. Identity matings were necessary to determine presence or absence of *tuh-3*. From *sbd*² *bx*³ *pbx tuh-3*⁺ *fl*/ *sbd*⁺ *bx*⁺ *pbx*⁺ *tuh-3* *fl*⁺ females, cross-over bearing males that were either *tuh-3 fl* or *tuh-3*⁺ *fl* were selected. The recombinant chromosomes were balanced over TMI.

Table 1. Localization of *tuh-3* by analyzing cross-overs between *pbx* and *fl* in 3R.

Strain	Tumorous-head defect				Male genital disc defect			
	with	without	total	% penetrance	with	without	total	% penetrance
<i>pbx tuh-3</i> ⁺ <i>fl</i>								
X	→ <i>pbx</i> ⁺ <i>tuh-3 fl</i>							
<i>pbx</i> ⁺ <i>tuh-3 fl</i> ⁺								
#3	65	38	103	63.1	32	70	102	31.4
#6	74	28	102	72.5	78	25	103	75.7
#11	69	30	99	69.7	80	21	101	79.2
#17	54	51	105	51.4	36	66	102	35.3
#23	83	20	103	80.6	55	70	125	44.0
#25	59	38	97	60.8	85	19	104	81.7
#26	67	32	99	67.7	24	83	107	22.4
#47	74	34	108	68.5	35	64	99	35.4
#49	76	51	127	59.8	54	47	101	53.5
#50	61	18	79	77.2	2	37	39	5.1
#81	70	46	116	60.3	59	42	101	58.4
#86	93	9	102	91.2	53	49	102	52.0
#90	47	16	63	74.6	72	55	127	56.7
#92	68	36	104	65.4	42	68	110	38.2
#94	74	26	100	74.0	59	48	107	55.1
Totals	1034	473	1507	68.6	766	764	1530	50.1
<i>pbx tuh-3</i> ⁺ <i>fl</i>								
X	→ <i>pbx</i> ⁺ <i>tuh-3</i> ⁺ <i>fl</i>							
<i>pbx</i> ⁺ <i>tuh-3 fl</i> ⁺								
#10	0	100	100	0.0	0	100	100	0.0
#21	0	111	111	0.0	0	118	118	0.0
#57	0	108	108	0.0	1	44	45	2.2
#73	0	102	102	0.0	0	106	106	0.0
#96	0	131	131	0.0	0	101	101	0.0
Totals	0	552	552	0.0	1	469	470	0.2

#57/Df(3R)P9 uncovers *tuh-3*

Two different genetic tests were needed to detect presence of *tuh-3*. The mutant gene acts as a semidominant in the presence of an X-linked maternal effect gene, symbolized *tuh-1h*, to cause growths of abdominal and/or genital tissue in the head. *Tuh-3* acts as a simple recessive in the presence of a second maternal effect gene, symbolized *tuh-1g*, which causes the defect to switch from the head to the posterior of the fly where internal and external genitalia may be completely absent. The maternal effect genes *tuh-1h* (head defects) and *tuh-1g* (genital defects) are naturally occurring alleles. The semidominant head defect phenotype was detected among F₁ progeny resulting from mating males of each cross-over strain to females that carried an attached X-chromosome homozygous for *tuh-1h* and a 3rd chromosome gene that enhances the penetrance and expression of the head defects. The recessive phenotype was found by mating the appropriate males to females possessing an attached X-chromosome homozygous for *tuh-1g* and backcrossing the attached X-chromosome bearing female offspring to the males from each strain being tested.

Table 1 shows that 15 of the 20 cross-overs between *pbx* and *f1* occurred distal to or right of *tuh-3*. The remaining 5 cross-overs were between *pbx* and *tuh-3*. *Tuh-3* was mapped much closer to *pbx* at 58.8+ than to *f1* at 59.7+. This places *tuh-3* at about 59.0+. However, it must be emphasized that an exact placement of *tuh-3* was not possible because the viability of flies with recombinant chromosomes was quite low. A total of 93 males was originally selected as potential cross-over types. Of these, 40 either were sterile or did not survive even light etherization. The 15 *tuh-3 f1* cross-over types showed both the head defect with *tuh-1h* and the genital defect with *tuh-1g*. The tumorous-head phenotype was seen in 68.6% of the flies examined with average penetrance ranging from a low of 51.4% to a high of 91.2%, while the genital disc defect was found in 50.0% of the males examined with a low of 5.1% penetrance to a high of 81.7% penetrance. The five *tuh-3⁺ f1* strains of flies lacked a mutant phenotype when confronted with either maternal effect gene. Strain #57 had 1 of 45 males showing the trait. However, when the *tuh-3⁺ f1* chromosome was made hemizygous for *tuh-3⁺* by placing it with *Df(3R)P9*, which uncovers *tuh-3*, none of the 100 males tested showed the phenotype.

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References: Kuhn, D.T., D.F. Woods & D.J. Andrew 1981, *Genetics* 99:99; Kuhn, D.T. & D.F. Woods 1982, *DIS* 58:96.

Larochelle, C., J. Côté, and F. Garcin.
Laval University, Quebec, Canada. The ethanol metabolic pathway in *D.melanogaster* and *D.simulans*.

The two cosmopolitan sibling species *D.melanogaster* and *D.simulans* are able to use environmental ethanol as source of energy; however, *D.melanogaster* exhibits better capacity to handle high ethanol concentrations than *D.simulans* (Parsons et al. 1979).

This differential tolerance to ethanol could be explained at the biochemical level by a much higher alcohol dehydrogenase (ADH) activity in *D.melanogaster* than in *D.simulans*. Nevertheless the ADH-mediated oxidation of ethanol generates acetaldehyde, a highly toxic product. For fly survival it is essential that acetaldehyde be rapidly oxidized into acetate.

In most animal species this critical step involves and NAD⁺-dependent aldehyde dehydrogenase (ALDH). In *Drosophila*, aldehyde oxidase (ALDOX) a flavine enzyme using many aldehydes as substrates, was assumed to be the active enzyme (see Courtright 1967; Dickinson 1970, 1971). Recently we have provided evidence that ALDH is present in *Drosophila* and that its physico-chemical properties are very similar to those of other animal species (Garcin et al. 1981, 1983).

In these experiments we have carried out a comparative study of ADH, ALDH, and ALDOX activities in *D.melanogaster* and *D.simulans*. We show that ADH and ALDH activities are directly correlated with the level of tolerance to ethanol and acetaldehyde whereas ALDOX activities are inversely correlated.

Drosophila melanogaster collected from Colmar (France) and *Drosophila simulans* collected from Villeurbanne (France) were kindly made available by Prof. J. David Laboratoire de Biologie et de Genetique Evolutive Gif-sur-Yvette (France). The flies were grown in low density populations on *Drosophila* medium (Carolina medium 4-24) and maintained in an incubator providing a constant temperature (25±1°C) and an 18 hr light/6 hr dark photoperiod with fluorescent light. Adult flies, 5 to 6 days old were frozen in liquid nitrogen, homogenized and centrifuged as described in Garcin et al. 1983. The resultant supernatants were kept