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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<sup>2</sup> bx<sup>3</sup> pbx fl*/TM1 males. F<sub>1</sub> females were backcrossed to *sbd<sup>2</sup> bx<sup>3</sup> pbx fl*/TM1 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<sup>2</sup> bx<sup>3</sup> pbx tuh-3<sup>+</sup> fl*/ *sbd<sup>+</sup> bx<sup>+</sup> pbx<sup>+</sup> tuh-3 fl<sup>+</sup>* females, cross-over bearing males that were either *tuh-3 fl* or *tuh-3<sup>+</sup> fl* were selected. The recombinant chromosomes were balanced over TM1.

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<sup>+</sup> fl</i>								
X	→ <i>pbx<sup>+</sup> tuh-3 fl</i>							
<i>pbx<sup>+</sup> tuh-3 fl<sup>+</sup></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<sup>+</sup> fl</i>								
X	→ <i>pbx<sup>+</sup> tuh-3<sup>+</sup> fl</i>							
<i>pbx<sup>+</sup> tuh-3 fl<sup>+</sup></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-lh*, 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-lg*, 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-lh* (head defects) and *tuh-lg* (genital defects) are naturally occurring alleles. The semidominant head defect phenotype was detected among F<sub>1</sub> progeny resulting from mating males of each cross-over strain to females that carried an attached X-chromosome homozygous for *tuh-lh* 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-lg* 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-lh* and the genital defect with *tuh-lg*. 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<sup>+</sup> 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<sup>+</sup> f1* chromosome was made hemizygous for *tuh-3<sup>+</sup>* 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.  
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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<sup>+</sup>-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