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Stocks
Notes: 1)TM6B, Hu e Tb ca tends to lose Tb
2)I suspect that the deficiency chromosomes may carry other marker mutations, but I have listed what was on the labels.

dev^{1}/ TM6C, cu Sb e ca

dev^{2} p^{K4}/ TM6C, cu Sb e ca

Df(3L)fzGF3b/ TM6B, Hu e Tb ca

Df(3L)fzCAL5/ TM3

Df(3L)fzGS1a/ TM3

Df(3L)fzD21/ TM6B, Hu e Tb ca

Df(3L)fzM21, th st/ TM9, p^{p} Ubx

D^{3} Sb ca^{2}/ TM6B, Hu e Tb ca

In(3LR)CxD, D/ Ly

In(3L)P In(3R)C, Mc e l(3)c/ In(3LR)DcxP, ru h D Sb e^{s}

Df(3L)fzGF3b	Df(3L)70B?; 70D6	
Df(3L)fzCAL5	Df(3L)70C2-7; 70E1-3	called Df(3L) in DIS62
Df(3L)fzGS1a	Df(3L)70C6-15; 70E4-6	
Df(3L)fzD21	Df(3L)70D1-3; 70E3-8	(70D; 71F in Indiana list)
Df(3L)fzM21	Df(3L)70D2-3; 71E4-5	

D^1 complements all five deficiencies for viability.

D^3 complements Df(3L)fzM21, but fails to complement Df(3L)fzD21 for viability.

dev fails to complement all deficiencies except Df(3L)fzM21.

dev^1/dev^2:Dp(3;Y)ST1 is lethal

map position of dev is 40.9 based on 10 recombinants between dev^1 and th in a total of 433 progeny.

Gene	Salivary band	Meiotic map position	
Ly	70A	40.5	
D	70C13-D1	40.7	(associated with inversion)
dev	70D1-2	40.9	
fz	70D6-7	41.7	
nc16	71D-F	42.8	
brm	72AB	43.0	
th	72D	43.2	
Gl	70C2	41.4	(map position inconsistent with others)