

TABLE S4

Estimates of the fraction of adaptive substitutions using the Fay, Wycoff and Wu method on the dataset restricted to the two-thirds of loci with the highest K_A/K_S values

<i>D. miranda</i>		<i>D. pseudoobscura</i>	
X chromosome	Autosome	X chromosome	Autosome
-0.180	-1.112	0.398	0.534
(-1.434 / 0.550)	(-2.906 / -0.042)	(0.050 / 0.669)	(0.193 / 0.796)

90% confidence intervals in parentheses.