

**Table S6 Comparison of expected and observed types of mutations**

	<i>dw3</i>		<i>rht1</i>	
	Expected	Observed	Expected	Observed
Missense	50.36%	70.18%	57.14%	50.00%
Silent	46.59%	27.19%	41.35%	50.00%
Stop codon	3.05%	2.63%	1.50%	