

Mated expressed gene vs unmated expressed gene	All	(-4.047, 0.000)
	Substitutions	(-3.1917, 0.000)
	Deletions	(-3.9309, 0.000)
Mated non-expressed gene vs unmated non-expressed gene	All	99.1% (-8.195, -0.020), p < 0.036*
	Substitutions	(-1.6898, 2.1997)
	Deletions	99.91% (-7.032, -1.545), p < 0.007**
Mated expressed gene vs mated non-expressed gene	All	(-4.683, 0.556)
	Substitutions	(-1.0654, 2.5748)
	Deletions	(-4.945, 0.281)
Unmated expressed gene vs unmated non-expressed gene	All	(-3.1202, 0.000)
	Substitutions	(-3.0924, 0.000)
	Deletions	N/A

Supplemental Table S11. Bootstrapped confidence intervals for average differences between mated vs unmated, expressed vs non-expressed mutation rates.

Bootstrap confidence levels are 95%, or higher if indicated. Gene expression status was determined as shown in Supplemental Fig S12a. Intervals highlighted in bold do not contain zero.