

Mating status	Gene name	Deletion disagreement length (bp)	FPKM
Mated	H12D21.6	4	0.00
Mated	<i>vit-1</i>	4	4.49
Mated	<i>cand-1</i>	4	92.47
Mated	<i>scrm-5</i>	4	0.00
Mated	<i>alh-1</i>	23	220.58
Mated	<i>abu-11</i>	4	0.00
Mated	F41E6.15	4	0.00
Mated	<i>gly-11</i>	7	0.00
Mated	<i>phf-10</i>	8	65.65
Mated	C31G12.4	6	0.00
Mated	<i>trr-1</i>	7	23.18
Mated	F53A10.2	19	0.00
Unmated	B0205.1	4	42.27
Mated	F56C11.6	10	0.26
Mated	<i>rha-1</i>	6	74.41
Mated	<i>srw-35</i>	5	0.00
Unmated	F49C5.4	6	0.00
Mated	F14D1.2	49	0.00
Mated	R04E5.8	11	2.00
Mated	Y47H9C.20	15	0.00
Mated	<i>lgo-35</i>	11	0.00
Mated	ZC123.10	4	0.00

Supplemental Table S12. Germline expression levels at loci of duplex deletion disagreements longer than 3 bp and at loci of deletions (selecting disagreements that fall in gene regions).