

Table 13. Quantitation of genes that affect Dscam exon 17 splicing

dsRNA	cryptic	17.2	17.1	skipping
Untreated	0.00%	9.33%	84.78%	5.10%
CG10279	3.28%	25.60%	36.64%	34.44%
CG11266	0.00%	46.02%	48.87%	4.77%
CG12085	0.00%	48.60%	42.56%	7.60%
CG1658	8.68%	32.37%	51.34%	7.61%
CG4602	0.00%	43.98%	48.88%	6.72%
CG5931	0.00%	49.48%	45.36%	4.46%
CG6999	0.00%	46.02%	49.53%	4.08%
CG7269	0.00%	31.39%	55.22%	12.57%