

	feature	observed	expected	fc	test	sig	p_val
3UTR	3UTR	9	8.386	1.1	enrichment	F	0.463
5UTR	5UTR	6	3.838	1.6	enrichment	F	0.188
intron	intron	86	73.052	1.2	enrichment	T	0.0256
ncRNA	ncRNA	6	5.612	1.1	enrichment	F	0.492
exon	exon	14	29.845	0.5	depletion	T	0.000393
intergenic	intergenic	33	42.709	0.8	depletion	T	0.0478