

表1. 524条Tos17标签在水稻基因组中各区间内的分布(以SR值>2.33为显著水平)

Table 1 Distribution of the 524 Tos17 insertions in different regions of the rice genome.

比较组	区域	插入标签数	期望值	标准差	区段大小
Test	Region	Number of insertion	Expected	SR	Chromosome size (Mb)
1	Genic	389	226	10.85	160.4
	TE-related	55	90	-3.68	63.8
	Intergenic	80	208	-8.88	147.8
2	Coding	344	269	4.55	36.6
	1kb upstream	66	116	-4.62	58.7
	500bp downstream	34	59	-3.25	60.4
3	Intron	154	185	-2.29	37.6
	Exon	190	159	2.48	35.5

