





- Promoter (<=1kb) (12.53%)
- Promoter (1-2kb) (5.46%)
- Promoter (2-3kb) (4.54%)
- 5' UTR (0.13%)
- 3' UTR (1.44%)
- 1st Exon (0.02%)
- Other Exon (2.13%)
- 1st Intron (13.2%)
- Other Intron (28.62%)
- Downstream (<=300) (0.11%)
- Distal Intergenic (31.82%)

Annotation Summary

Annotated peaks generated by ChIPseeker

32641/33704 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	12.52718973
10	Promoter (1–2kb)	5.45939156
11	Promoter (2–3kb)	4.54030207
4	5' UTR	0.13479979
3	3' UTR	1.43684323
1	1st Exon	0.02144542
7	Other Exon	2.13228761
2	1st Intron	13.19812506
8	Other Intron	28.61738305
6	Downstream (<=300)	0.11335437
5	Distal Intergenic	31.81887810