





- Promoter (<=1kb) (41.28%)
- Promoter (1-2kb) (5.26%)
- Promoter (2-3kb) (3.99%)
- 5' UTR (0.1%)
- 3' UTR (1%)
- 1st Exon (0.01%)
- Other Exon (1.49%)
- 1st Intron (9.65%)
- Other Intron (16.94%)
- Downstream (<=300) (0.07%)
- Distal Intergenic (20.21%)

Annotation Summary

Annotated peaks generated by ChIPseeker

15253/15556 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	41.277125811
10	Promoter (1–2kb)	5.257982036
11	Promoter (2–3kb)	3.986101095
4	5' UTR	0.104897397
3	3' UTR	1.003081361
1	1st Exon	0.006556087
7	Other Exon	1.494787911
2	1st Intron	9.650560545
8	Other Intron	16.940929653
6	Downstream (<=300)	0.072116961
5	Distal Intergenic	20.205861142