





- Promoter (<=1kb) (17.84%)
- Promoter (1-2kb) (7.23%)
- Promoter (2-3kb) (4.56%)
- 5' UTR (0.04%)
- 3' UTR (1.14%)
- Other Exon (1.65%)
- 1st Intron (13.1%)
- Other Intron (24.43%)
- Downstream (<=300) (0.08%)
- Distal Intergenic (29.92%)

# Annotation Summary

Annotated peaks generated by ChIPseeker

2366/2419 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
8	Promoter (<=1kb)	17.83601014
9	Promoter (1–2kb)	7.22738800
10	Promoter (2–3kb)	4.56466610
3	5' UTR	0.04226543
2	3' UTR	1.14116653
6	Other Exon	1.64835165
1	1st Intron	13.10228233
7	Other Intron	24.42941674
5	Downstream (<=300)	0.08453085
4	Distal Intergenic	29.92392223