

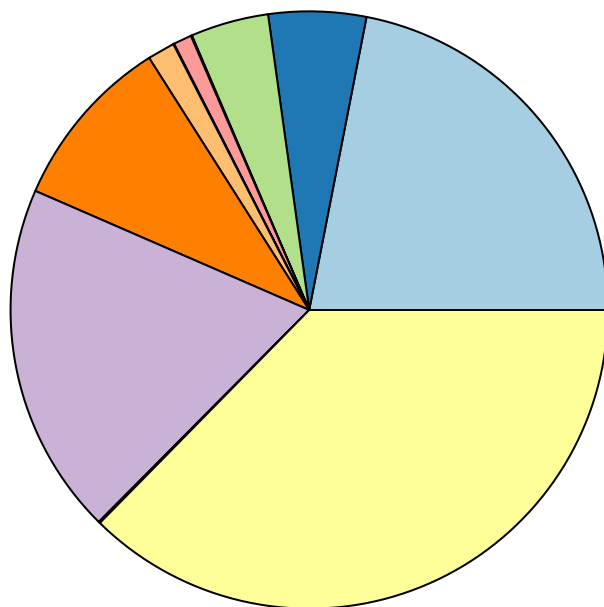
Annotation Summary

Annotated peaks generated by ChIPseeker

12183/12663 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	21.92399245
10	Promoter (1–2kb)	5.27784618
11	Promoter (2–3kb)	4.22720184
4	5' UTR	0.05745711
3	3' UTR	0.99318723
1	1st Exon	0.04104079
7	Other Exon	1.48567676
2	1st Intron	9.46400722
8	Other Intron	19.08396947
6	Downstream (<=300)	0.08208159
5	Distal Intergenic	37.36353936



- Promoter (<=1kb) (21.92%)
- Promoter (1-2kb) (5.28%)
- Promoter (2-3kb) (4.23%)
- 5' UTR (0.06%)
- 3' UTR (0.99%)
- 1st Exon (0.04%)
- Other Exon (1.49%)
- 1st Intron (9.46%)
- Other Intron (19.08%)
- Downstream (<=300) (0.08%)
- Distal Intergenic (37.36%)