

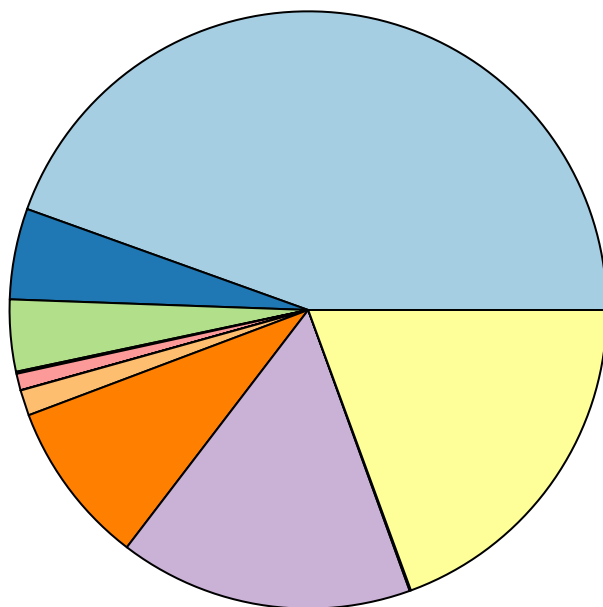
Annotation Summary

Annotated peaks generated by ChIPseeker

25499/25998 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	44.51154947
10	Promoter (1–2kb)	4.92176164
11	Promoter (2–3kb)	3.88642692
4	5' UTR	0.10980823
3	3' UTR	0.90199616
1	1st Exon	0.01176517
7	Other Exon	1.38828974
2	1st Intron	8.88270128
8	Other Intron	15.90650614
6	Downstream (<=300)	0.07059100
5	Distal Intergenic	19.40860426



- Promoter (<=1kb) (44.51%)
- Promoter (1-2kb) (4.92%)
- Promoter (2-3kb) (3.89%)
- 5' UTR (0.11%)
- 3' UTR (0.9%)
- 1st Exon (0.01%)
- Other Exon (1.39%)
- 1st Intron (8.88%)
- Other Intron (15.91%)
- Downstream (<=300) (0.07%)
- Distal Intergenic (19.41%)