

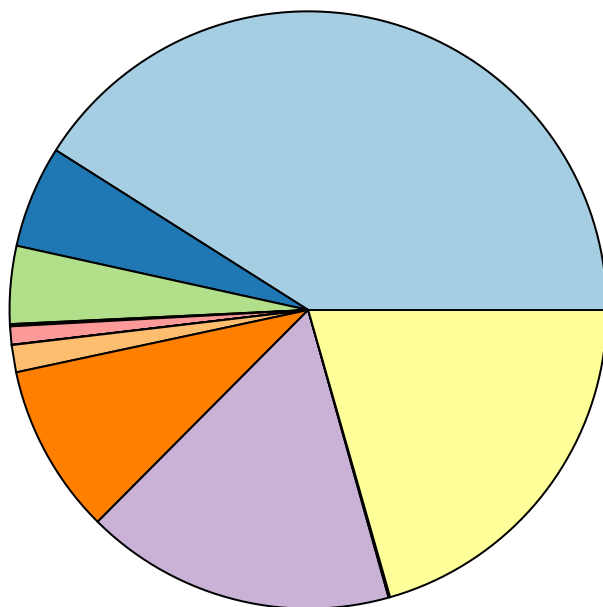
Annotation Summary

Annotated peaks generated by ChIPseeker

27614/28221 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	41.01542696
10	Promoter (1–2kb)	5.52980372
11	Promoter (2–3kb)	4.21163178
4	5' UTR	0.11950460
3	3' UTR	0.98500760
1	1st Exon	0.01448541
7	Other Exon	1.47389006
2	1st Intron	9.19823278
8	Other Intron	16.79220685
6	Downstream (<=300)	0.08329108
5	Distal Intergenic	20.57651916



- Promoter (<=1kb) (41.02%)
- Promoter (1-2kb) (5.53%)
- Promoter (2-3kb) (4.21%)
- 5' UTR (0.12%)
- 3' UTR (0.99%)
- 1st Exon (0.01%)
- Other Exon (1.47%)
- 1st Intron (9.2%)
- Other Intron (16.79%)
- Downstream (<=300) (0.08%)
- Distal Intergenic (20.58%)