

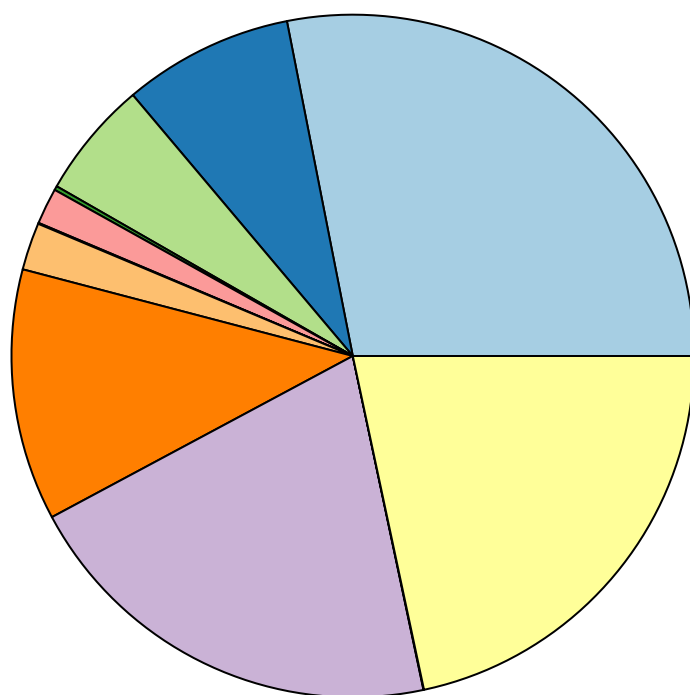
Annotation Summary

Annotated peaks generated by ChIPseeker

4147/4150 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	28.09259706
10	Promoter (1–2kb)	8.05401495
11	Promoter (2–3kb)	5.57029178
4	5' UTR	0.19291054
3	3' UTR	1.71208102
1	1st Exon	0.04822763
7	Other Exon	2.24258500
2	1st Intron	11.91222571
8	Other Intron	20.49674463
6	Downstream (<=300)	0.02411382
5	Distal Intergenic	21.65420786



- Promoter (<=1kb) (28.09%)
- Promoter (1-2kb) (8.05%)
- Promoter (2-3kb) (5.57%)
- 5' UTR (0.19%)
- 3' UTR (1.71%)
- 1st Exon (0.05%)
- Other Exon (2.24%)
- 1st Intron (11.91%)
- Other Intron (20.5%)
- Downstream (<=300) (0.02%)
- Distal Intergenic (21.65%)