

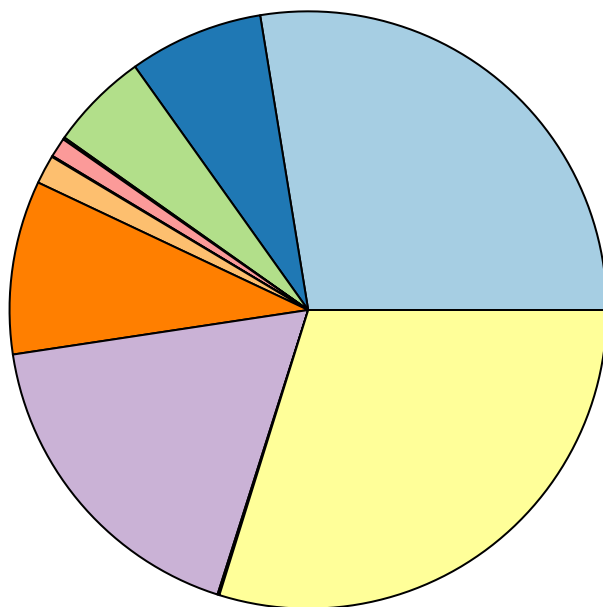
Annotation Summary

Annotated peaks generated by ChIPseeker

15453/16027 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	27.57393386
10	Promoter (1–2kb)	7.27366854
11	Promoter (2–3kb)	5.35171164
4	5' UTR	0.11648224
3	3' UTR	1.05481136
1	1st Exon	0.05824112
7	Other Exon	1.55309649
2	1st Intron	9.40270498
8	Other Intron	17.70529994
6	Downstream (<=300)	0.10353977
5	Distal Intergenic	29.80651006



- Promoter ($\leq 1\text{kb}$) (27.57%)
- Promoter (1–2kb) (7.27%)
- Promoter (2–3kb) (5.35%)
- 5' UTR (0.12%)
- 3' UTR (1.05%)
- 1st Exon (0.06%)
- Other Exon (1.55%)
- 1st Intron (9.4%)
- Other Intron (17.71%)
- Downstream (≤ 300) (0.1%)
- Distal Intergenic (29.81%)