

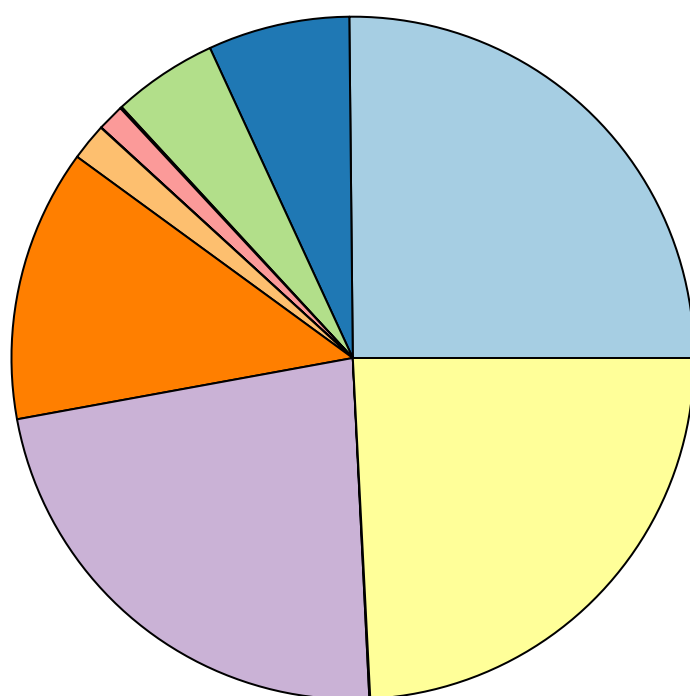
Annotation Summary

Annotated peaks generated by ChIPseeker

4904/4906 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	25.16313214
10	Promoter (1–2kb)	6.70880914
11	Promoter (2–3kb)	4.97553018
4	5' UTR	0.08156607
3	3' UTR	1.26427406
1	1st Exon	0.02039152
7	Other Exon	1.75367047
2	1st Intron	12.90783034
8	Other Intron	22.89967374
6	Downstream (<=300)	0.04078303
5	Distal Intergenic	24.18433931



- Promoter (<=1kb) (25.16%)
- Promoter (1-2kb) (6.71%)
- Promoter (2-3kb) (4.98%)
- 5' UTR (0.08%)
- 3' UTR (1.26%)
- 1st Exon (0.02%)
- Other Exon (1.75%)
- 1st Intron (12.91%)
- Other Intron (22.9%)
- Downstream (<=300) (0.04%)
- Distal Intergenic (24.18%)