

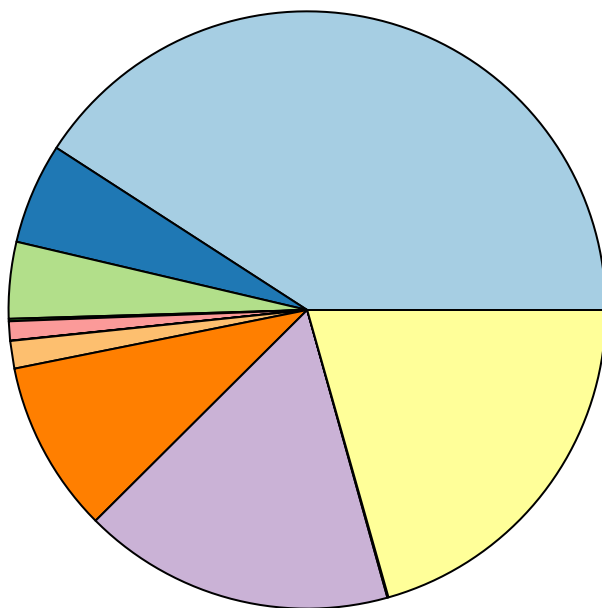
# Annotation Summary

Annotated peaks generated by ChIPseeker

28911/29552 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	40.86333921
10	Promoter (1–2kb)	5.46504791
11	Promoter (2–3kb)	4.14375151
4	5' UTR	0.13143786
3	3' UTR	1.03766732
1	1st Exon	0.01383556
7	Other Exon	1.49424095
2	1st Intron	9.31479368
8	Other Intron	16.86555290
6	Downstream (<=300)	0.05880115
5	Distal Intergenic	20.61153194



- Promoter (<=1kb) (40.86%)
- Promoter (1-2kb) (5.47%)
- Promoter (2-3kb) (4.14%)
- 5' UTR (0.13%)
- 3' UTR (1.04%)
- 1st Exon (0.01%)
- Other Exon (1.49%)
- 1st Intron (9.31%)
- Other Intron (16.87%)
- Downstream (<=300) (0.06%)
- Distal Intergenic (20.61%)