

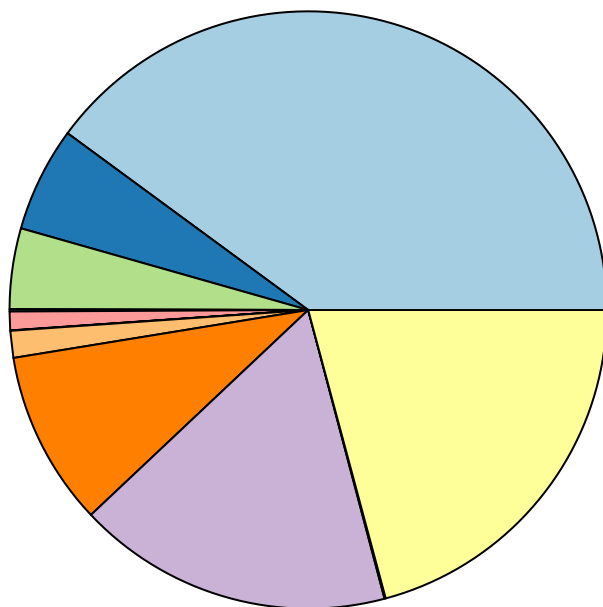
# Annotation Summary

Annotated peaks generated by ChIPseeker

32152/32891 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	39.91664593
10	Promoter (1–2kb)	5.67305300
11	Promoter (2–3kb)	4.36364767
4	5' UTR	0.12751928
3	3' UTR	1.01082359
1	1st Exon	0.01866136
7	Other Exon	1.45869619
2	1st Intron	9.46441901
8	Other Intron	17.08136352
6	Downstream (<=300)	0.06531475
5	Distal Intergenic	20.81985569



- Promoter (<=1kb) (39.92%)
- Promoter (1-2kb) (5.67%)
- Promoter (2-3kb) (4.36%)
- 5' UTR (0.13%)
- 3' UTR (1.01%)
- 1st Exon (0.02%)
- Other Exon (1.46%)
- 1st Intron (9.46%)
- Other Intron (17.08%)
- Downstream (<=300) (0.07%)
- Distal Intergenic (20.82%)