

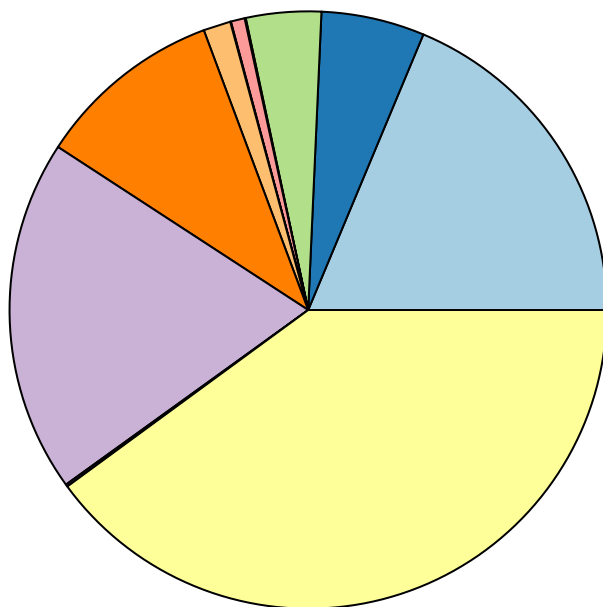
Annotation Summary

Annotated peaks generated by ChIPseeker

8118/8459 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	18.69918699
10	Promoter (1–2kb)	5.59251047
11	Promoter (2–3kb)	4.06504065
4	5' UTR	0.06159153
3	3' UTR	0.75141661
1	1st Exon	0.03695492
7	Other Exon	1.47819660
2	1st Intron	10.12564671
8	Other Intron	19.13032767
6	Downstream (<=300)	0.12318305
5	Distal Intergenic	39.93594481



- Promoter (<=1kb) (18.7%)
- Promoter (1-2kb) (5.59%)
- Promoter (2-3kb) (4.07%)
- 5' UTR (0.06%)
- 3' UTR (0.75%)
- 1st Exon (0.04%)
- Other Exon (1.48%)
- 1st Intron (10.13%)
- Other Intron (19.13%)
- Downstream (<=300) (0.12%)
- Distal Intergenic (39.94%)