

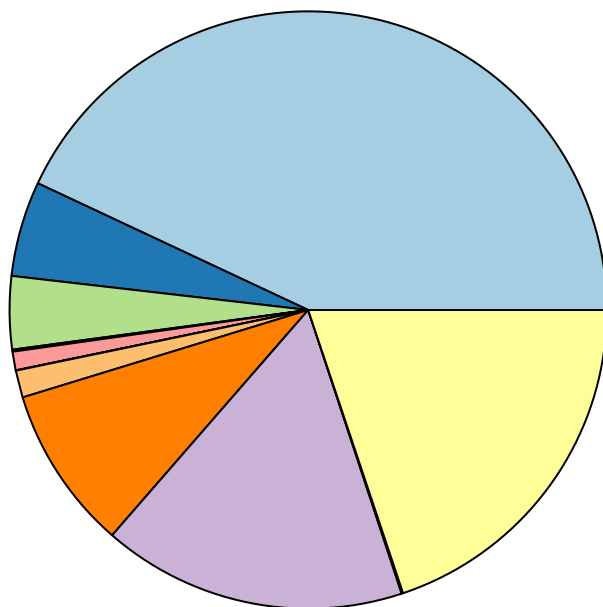
Annotation Summary

Annotated peaks generated by ChIPseeker

30039/30700 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
9	Promoter (<=1kb)	43.02406871
10	Promoter (1–2kb)	5.15330071
11	Promoter (2–3kb)	3.94487167
4	5' UTR	0.10985719
3	3' UTR	1.00868870
1	1st Exon	0.01664503
7	Other Exon	1.47142049
2	1st Intron	8.87179999
8	Other Intron	16.46859083
6	Downstream (<=300)	0.07656713
5	Distal Intergenic	19.85418955



- Promoter (<=1kb) (43.02%)
- Promoter (1-2kb) (5.15%)
- Promoter (2-3kb) (3.94%)
- 5' UTR (0.11%)
- 3' UTR (1.01%)
- 1st Exon (0.02%)
- Other Exon (1.47%)
- 1st Intron (8.87%)
- Other Intron (16.47%)
- Downstream (<=300) (0.08%)
- Distal Intergenic (19.85%)