

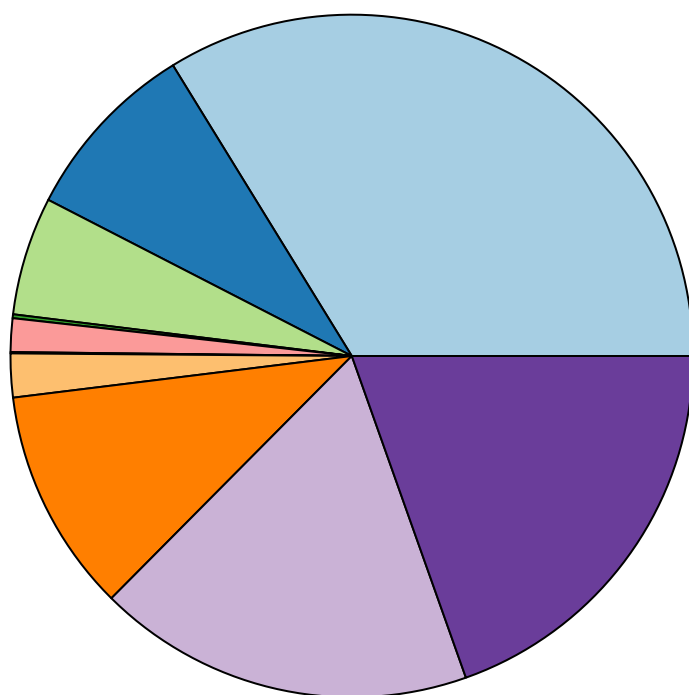
Annotation Summary

Annotated peaks generated by ChIPseeker

1640/1640 peaks were annotated

Genomic Annotation Summary:

	Feature	Frequency
8	Promoter (<=1kb)	33.78048780
9	Promoter (1–2kb)	8.65853659
10	Promoter (2–3kb)	5.60975610
4	5' UTR	0.18292683
3	3' UTR	1.58536585
1	1st Exon	0.06097561
6	Other Exon	2.07317073
2	1st Intron	10.60975610
7	Other Intron	17.86585366
5	Distal Intergenic	19.57317073



- Promoter (<=1kb) (33.78%)
- Promoter (1-2kb) (8.66%)
- Promoter (2-3kb) (5.61%)
- 5' UTR (0.18%)
- 3' UTR (1.59%)
- 1st Exon (0.06%)
- Other Exon (2.07%)
- 1st Intron (10.61%)
- Other Intron (17.87%)
- Distal Intergenic (19.57%)