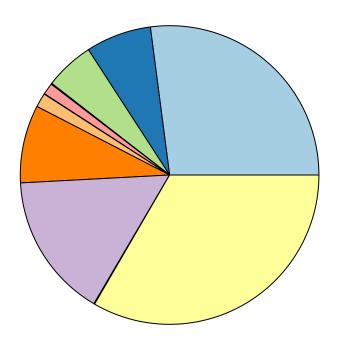


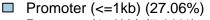
## **Annotation Summary**

Annotated peaks generated by ChIPseeker 9420/9828 peaks were annotated Genomic Annotation Summary:

Feature Frequency

- 9 Promoter (<=1kb) 27.05944798
- 10 Promoter (1–2kb) 7.09129512
- 11 Promoter (2–3kb) 5.32908705
- 4 5' UTR 0.10615711
- 3 3' UTR 1.25265393
- 1 1st Exon 0.04246285
- 7 Other Exon 1.52866242
- 2 1st Intron 8.43949045
- 8 Other Intron 15.69002123
- 6 Downstream (<=300) 0.08492569
- 5 Distal Intergenic 33.37579618





- Promoter (1–2kb) (7.09%)
- Promoter (2–3kb) (5.33%)5' UTR (0.11%)
- 3' UTR (1.25%)
- 1st Exon (0.04%)
- Other Exon (1.53%)
- 1st Intron (8.44%)Other Intron (15.69%)
- Downstream (<=300) (0.08%)
- □ Distal Intergenic (33.38%)