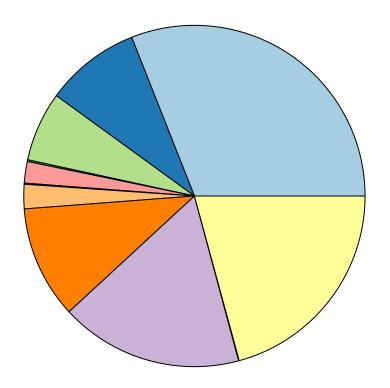


## **Annotation Summary**

Annotated peaks generated by ChIPseeker 1820/1821 peaks were annotated Genomic Annotation Summary:

Feature Frequency

- 9 Promoter (<=1kb) 30.98901099
- 10 Promoter (1-2kb) 9.01098901
- 11 Promoter (2-3kb) 6.59340659
- 4 5' UTR 0.16483516
- 3 3' UTR 2.03296703
- 1 1st Exon 0.10989011
- 7 Other Exon 2.30769231
- 2 1st Intron 10.65934066
- 8 Other Intron 17.30769231
- 6 Downstream (<=300) 0.05494505
- 5 Distal Intergenic 20.76923077



- □ Promoter (<=1kb) (30.99%)
- Promoter (1–2kb) (9.01%)
- Promoter (2–3kb) (6.59%)
- 5' UTR (0.16%)
- 3' UTR (2.03%)
- 1st Exon (0.11%)
- Other Exon (2.31%)
- 1st Intron (10.66%)
- ☐ Other Intron (17.31%)
- Downstream (<=300) (0.05%)
- □ Distal Intergenic (20.77%)