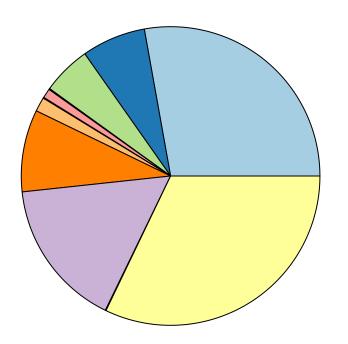


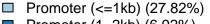
## **Annotation Summary**

Annotated peaks generated by ChIPseeker 14169/14715 peaks were annotated Genomic Annotation Summary:

Feature Frequency

- 9 Promoter (<=1kb) 27.82130002
- 10 Promoter (1-2kb) 6.92356553
- 11 Promoter (2–3kb) 5.30030348
- 4 5' UTR 0.11292258
- 3 3' UTR 1.01630320
- 1 1st Exon 0.09174959
- 7 Other Exon 1.54562778
- 2 1st Intron 8.89265297
- 8 Other Intron 16.09852495
- 6 Downstream (<=300) 0.07763427
- 5 Distal Intergenic 32.11941563





- Promoter (1–2kb) (6.92%)Promoter (2–3kb) (5.3%)
- 5' UTR (0.11%)
- 3' UTR (1.02%)
- 1st Exon (0.09%)
- Other Exon (1.55%)1st Intron (8.89%)
- ☐ Other Intron (16.1%)
- Downstream (<=300) (0.08%)</li>
- □ Distal Intergenic (32.12%)