

Genomic Feature

Fold Enrichment

2.0
1.5
1.0
0.5
0.0

1stExon

OtherExon

1stIntron

OtherIntron

3'UTR

5'UTR

IGR

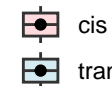
TSS1000

TSS2000

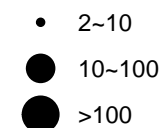
TSS3000

Region

Type



$-\log_{10}(\text{P-value})$



Cancer Type

