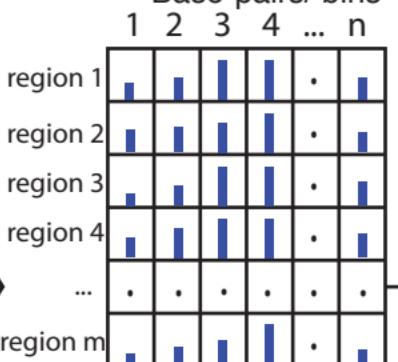


Genomic Intervals

BAM
BigWig
BED
GFF
Tab txt
GRanges

Summarize

ScoreMatrix/ScoreMatrixList object
Base-pairs/ bins



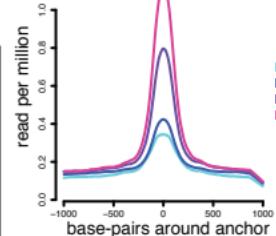
Annotation

BED
GFF
Tab txt
GRanges

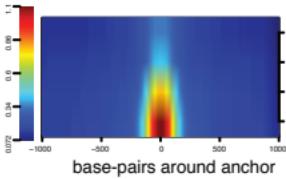
Annotate

Visualize

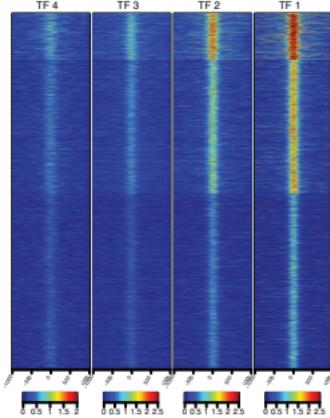
meta-region plots



meta-region heatmaps



heatmaps for genomic interval sets



Piecharts for annotation

