



CHANCE: ChIP-seq Analytics and Confidence Estimation

Main Output:

Export main output to text file

View IP strength

Compare with ENCODE

Multi IP normalization

Validate region list

View nucleotide frequency

View Phred score map

View read density bias

Welcome to CHANCE!

Reading wgEncodeBroadHistoneH1heschH3k4me1StdAlnRep1.bam... this may take a few minutes.
wgEncodeBroadHistoneH1heschH3k4me1StdAlnRep1.bam processed successfully, 8951369 tags processed

Samples:

Create sample

Delete sample

HESC H3K4me3 - # reads: 8951369 - genome: hg19 - wgEncodeBroadHistoneH1heschH3k4me1StdAlnRep1

Save samples to file

Restore saved samples