

Samtools: idxstats: Mapped reads per contig

Observed over Expected Counts, 3 lines

log10 (Observed over expected counts)

4000
3000
2000
1000
0

Chromosome name

chr1 chr10 chr11 chr12 chr13 chr14 chr15 chr16 chr17 chr18 chr19 chr2 chr20 chr21 chr22 chr22_KI270736v1_random chr3 chr4 chr5 chr6 chr7 chr8 chr9 chrM chrUn_KN707896v1_decoy chrX

