

Samtools idxstats: Mapped reads per contig

log10 (Observed over expected counts)

400
300
200
100
0

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 Xhsb107.05 chr1 chr2 chr3 chr4 chr5 chr6 chr7 chr8 chr9 chr10 chr11 chr12 chr13 chr14 chr15 chr16 chr17 chr18 chr19 chr20 chr21 chr22 chrM chrX

Chromosome Name

First_SmallTest_Paired.bam.idxstats JK2782_TGGCCGATCAACGA_L008_R1_001.fastq.gz.tengrand.fq.combined.fq.mapped.bam.idxstats Second_SmallTest_Paired.bam.idxstats wgEncodeUwRepliSeqK562G1AInRep1.bam.idxstats

