

# Samtools idxstats: Mapped reads per contig

log10 ( Observed over expected counts )

10000  
8000  
6000  
4000  
2000  
0

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

Chromosome Name

LIVER\_RNA\_HET\_1\_S4\_L001

LIVER\_RNA\_SMA\_1\_S1\_L001

LIVER\_RNA\_SMA\_2\_S2\_L001

P7\_LIVER\_RNA\_HET\_3\_S6\_L001

P7\_LIVER\_RNA\_SMA\_3\_S3\_L001

LIVER\_RNA\_HET\_2\_S5\_L001

