

Samtools idxstats: Mapped reads per contig

log10 (Observed over expected counts)

SPT5_INPUT_R1 SPT5_T0_R1 SPT5_T0_R2 SPT5_T15_R1 SPT5_T15_R2 SPT5_INPUT_R2

Chromosome Name

I II III IV IX MT V VI VII VIII X XI XII XIII XIV XV XVI

