

Samtools idxstats: Mapped reads per contig

log10 (Observed over expected counts)

— SPT5_INPUT_R1
— SPT5_INPUT_R2
— SPT5_T0_R1
— SPT5_T0_R2
— SPT5_T15_R1
— SPT5_T15_R2

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

Chromosome Name

