

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

Chromosome Name

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

