

# Samtools idxstats: Mapped reads per contig

log10 ( Observed over expected counts )

— SPT5\_INPUT\_R1\_T1  
— SPT5\_INPUT\_R2\_T1  
— SPT5\_T0\_R1\_T1  
— SPT5\_T0\_R2\_T1  
— SPT5\_T15\_R1\_T1  
— SPT5\_T15\_R2\_T1

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

Chromosome Name

