

# Samtools idxstats: Mapped reads per contig

log10 ( Observed over expected counts )

OSMOTIC\_STRESS\_T0\_PE\_REP1 OSMOTIC\_STRESS\_T100\_SE\_REP1 OSMOTIC\_STRESS\_T100\_SE\_REP2 OSMOTIC\_STRESS\_T150\_SE\_REP1 OSMOTIC\_STRESS\_T15\_PE\_REP1 OSMOTIC\_STRESS\_T0\_PE\_REP2

Chromosome Name

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

