

Samtools idxstats: Mapped reads per contig

log10 ( Observed over expected counts )

1.6

1.4

1.2

1.0

0.8

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

Chromosome Name

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

