

Samtools idxstats: Mapped reads per contig

log10 (Observed over expected counts)

1.04

1.02

1.00

0.98

0.96

I

Chromosome Name

RAP1_IAA_30M_REP1

RAP1_UNINDUCED_REP1

RAP1_UNINDUCED_REP2

WT_REP1

WT_REP2