SC8 Genome wide normalized coverage plot – U strains 3.0 -**STRAIN** 2.5 **-**A2.U10A A22.UpperP A3.U3G A4.U3B A5.U10D 2.0 -A6.U5C A7.U3E A8.U2E **LOCALE** 1.5 -U 1.0 -

10

CHROM_WINDOW

15

5

MEDIAN_COVERAGE

0