SC8 Genome wide normalized coverage plot – U strains 3.0 -2.5 -**STRAIN** A2.U10A A22.UpperP MEDIAN_COVERAGE A3.U3G A4.U3B A5.U10D A6.U5C A7.U3E A8.U2E 1.5 -**LOCALE** U 1.0 -5 10 15 0

CHROM_WINDOW