SC5 Genome wide normalized coverage plot – U strains 1.6 -1.4 -**STRAIN** A2.U10A A22.UpperP MEDIAN_COVERAGE A3.U3G A4.U3B A5.U10D 1.2 -A6.U5C A7.U3E A8.U2E **LOCALE** 1.0 -• U 0.8 -10 20 30 0 CHROM_WINDOW