SC8 Genome wide normalized coverage plot – U strains 1.2 -1.1 -**STRAIN** A2.U10A MEDIAN_COVERAGE A3.U3G A4.U3B A5.U10D A6.U5C A7.U3E A8.U2E **LOCALE** • U 0.9 -0.8 -2 0 6 CHROM_WINDOW