SC2 Genome wide normalized coverage plot – U strains 1.5 **-**1.3 -**STRAIN** A2.U10A A3.U3G MEDIAN_COVERAGE A4.U3B A5.U10D A6.U5C A7.U3E A8.U2E **LOCALE** • U 0.9 -10 15 20 5 0 CHROM_WINDOW