SC1 Genome wide normalized coverage plot – U strains 1.5 -**STRAIN** A2.U10A A3.U3G MEDIAN_COVERAGE A4.U3B ▶ A5.U10D 1.2 -A6.U5C ▶ A7.U3E A8.U2E LOCALE • U 0.9 -0.6 -20 25 5 15 0 CHROM_WINDOW