SC1 Genome wide normalized coverage plot – U strains 1.5 -**STRAIN** A2.U10A A22.UpperP MEDIAN_COVERAGE A3.U3G A4.U3B 1.2 **-**A5.U10D ● A6.U5C A7.U3E A8.U2E LOCALE • U 0.9 -0.6 -20 40 60 0 CHROM_WINDOW