SC2 Genome wide normalized coverage plot – L strains 1.6 **-**LOCALE 1.4 -• L **STRAIN** A10.L4C MEDIAN_COVERAGE A11.L11G 1.2 -A12.L10B A13.L3B A14.L4E A15.L4G A16.L7C A17.L1B 1.0 -A23.LowerP A9.L3D ctl1.AL1B ctl2.AL1B 0.8 -20 40 0 CHROM_WINDOW