SC3 Genome wide normalized coverage plot – L strains 1.75 **-**1.50 **-**LOCALE MEDIAN\_COVERAGE **STRAIN** C30.13LM C4.9L1 C5.9L2 C6.9L3 C7.9L4 ctl3.C9L1 0.75 -10 20 30 40 CHROM\_WINDOW