SC4 Genome wide normalized coverage plot – L strains 1.5 -LOCALE MEDIAN_COVERAGE **STRAIN** 1.2 -C30.13LM ● C4.9L1 C5.9L2 C6.9L3 C7.9L4 ctl3.C9L1 0.9 -0.6 -20 0 10 40 30 CHROM_WINDOW