SC1 Genome wide normalized coverage plot – L strains 1.50 -**LOCALE** • L 1.25 -**STRAIN** A10.L4C A11.L11G A12.L10B A13.L3B A14.L4E 1.00 -A15.L4G A16.L7C A17.L1B A23.LowerP A9.L3D ctl1.AL1B ctl2.AL1B 0.75 -

CHROM\_WINDOW

40

60

20

MEDIAN\_COVERAGE