SC5 Genome wide normalized coverage plot **STRAIN** C10.9M3 C11.9M41 C12.9U1 C13.9U2 C14.9U3 1.5 - $\triangle$ C15.9U4s 0 C30.13LM MEDIAN\_COVERAGE C31.13MP C4.9L1 C5.9L2 C6.9L3 + C7.9L4 C8.9M1 C9.9M2 1.0 -ctl3.C9L1 LOCALE O L  $\triangle$  M ☐ Stock +  $\upsilon$  $^{\triangle}\,_{\triangle}\, \triangleq$ 10 30 20 0 CHROM\_WINDOW