SC3 Genome wide normalized coverage plot 0 **LOCALE** 8 O L 1.50 - $\triangle$  M □ U **STRAIN** B16.1310D MEDIAN\_COVERAGE B17.1311G 1.25 -B18.1312G B19.1310A B20.1311A B21.1312A B22.134E 1.00 -B23.133H B24.133F (0000) (I) B25.135C B26.131A B27.136A 00 B28.13UM 0.75 -B29.13MM 0 10 30 20 40 CHROM\_WINDOW