SC5 Genome wide normalized coverage plot 0 1.75 **-**8 **LOCALE** 0 O L  $\triangle$  U 1.50 -**STRAIN** B16.1310D MEDIAN\_COVERAGE B17.1311G B18.1312G 0 B19.1310A 1.25 -B20.1311A 0 B21.1312A B22.134E B23.133H B24.133F B25.135C 1.00 -B26.131A 会 B27.136A B28.13UM 0.75 -10 20 30 CHROM\_WINDOW