SC5 Genome wide normalized coverage plot – U strains 1.6 **-**1.4 -**STRAIN** B22.134E B23.133H MEDIAN_COVERAGE B24.133F 1.2 -B25.135C B26.131A B27.136A B28.13UM 1.0 -LOCALE • U • • • 0.8 -10 30 20 0 CHROM_WINDOW