SC4 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 20 40 0 30

CHROM_WINDOW