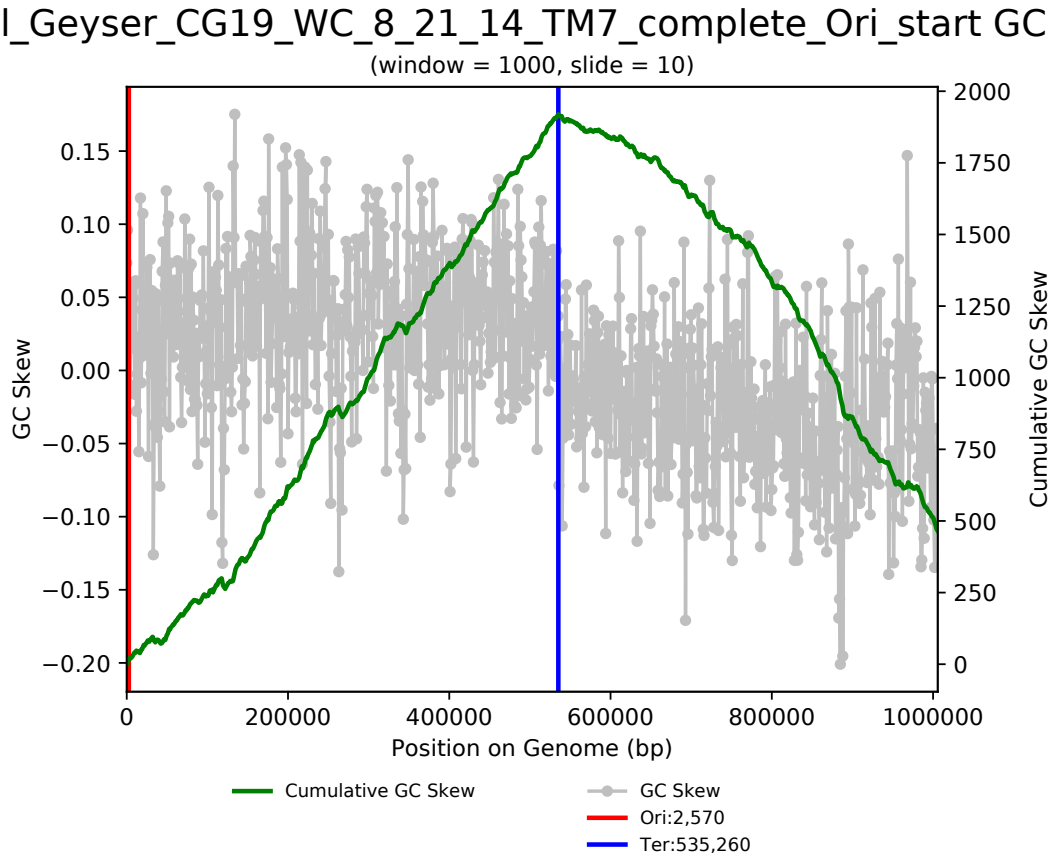

Supplemental Figure S1 GC skew plot of the circularized and curated Saccharibacteria genome from CG_WC. iRep (Brown et al. 2016) was used to predict the position of the origin and terminus as well as calculate and plot the GC skew across the genome. Predicted origin is represented by the red line and predicted terminus is represented by the blue line.