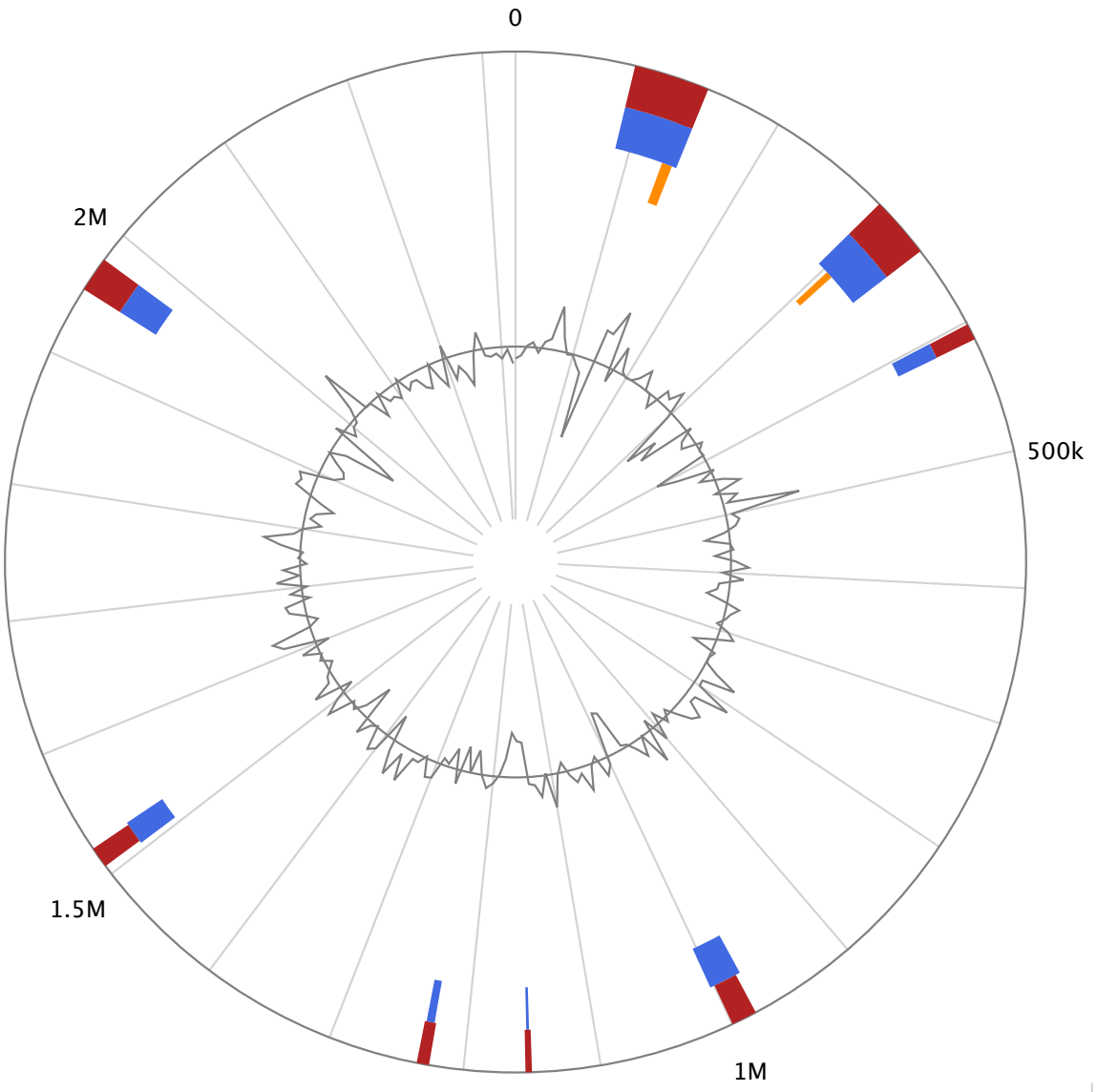


Figure S1. Distribution of genomic islands in the *B. campestris* DSM 4712 chromosome. Red, blue, and orange rectangles represent genomic islands predicted by the "Integrated", "IslandPath-DIMOB," and "SIGI-HMM" methods, respectively. The solid line in the center represents GC content percentage. Chromosomal positions are marked at 0.1 Mb intervals.