



Supplemental Figure 1. Domains and transmembrane helix predictions of GBS LCP proteins. (A) *brpA* (W903_0426, accession: WP_000708159.1), (B) *cpsA* (W903_1258, accession: WP_000064987.1), and (C) putative *psr* (W903_0744, WP_000089333.1) conserved LCP domains were predicted using Pfam 31.0 (top images). Transmembrane helices were predicted using TMHMM Server v. 2.0 (bottom images).