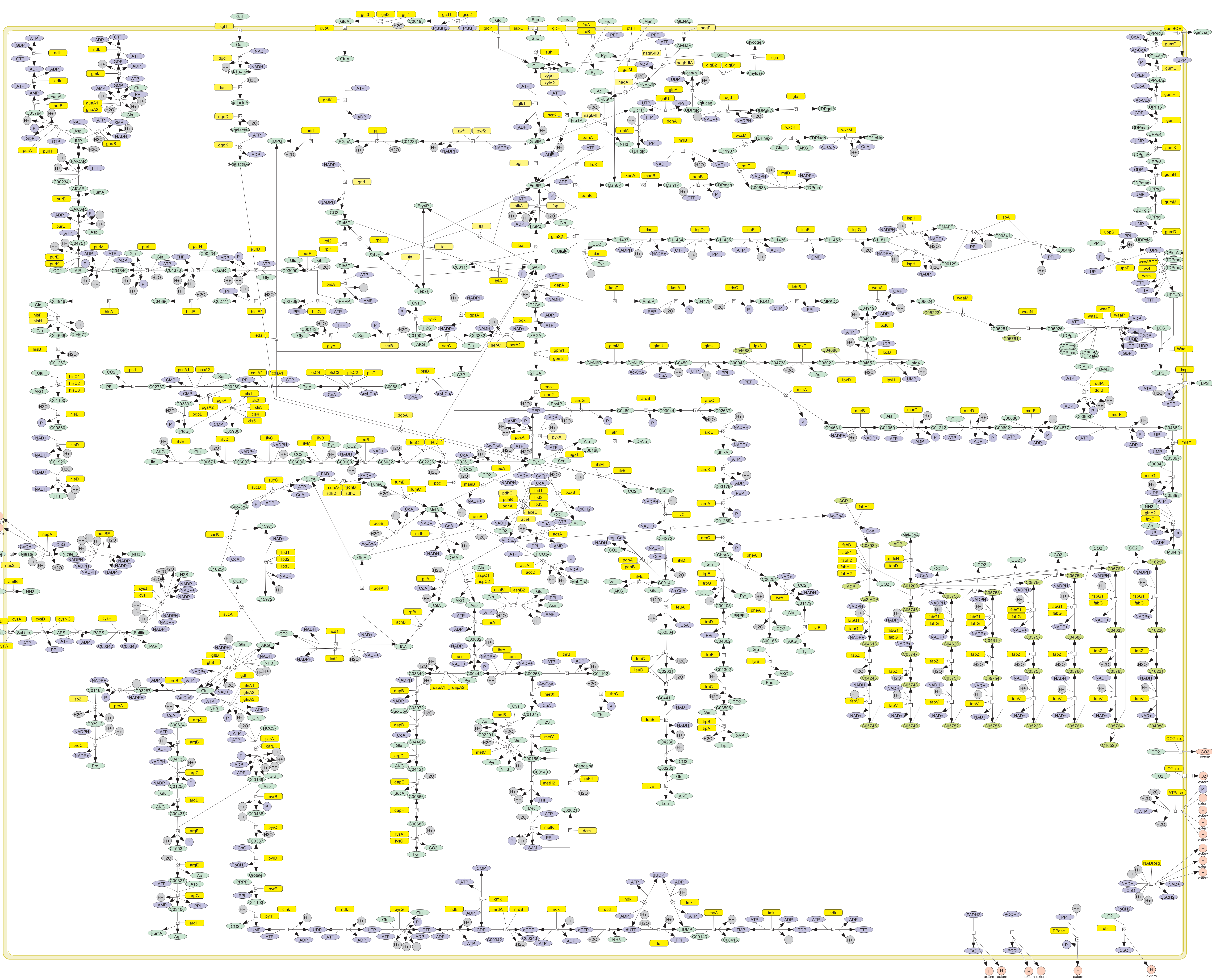


# Supplementary Figure 2. Genome-based reconstruction of the central metabolism and the polysaccharide biosynthesis pathways of *Xanthomonas campestris* pv. *campestris* B100 in systems biology markup language (SBML)



Additional metabolites are identified by their Compound IDs that reference the KEGG database <http://www.genome.jp/kegg/>