

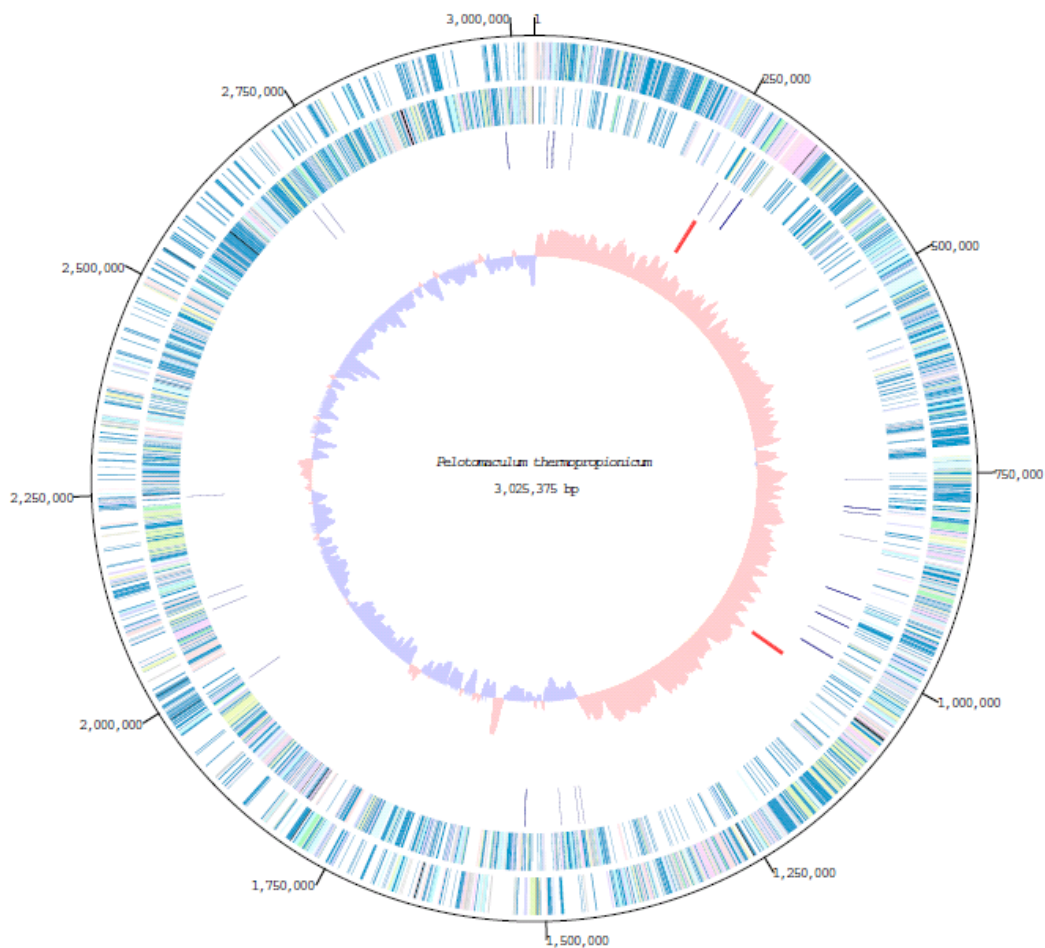


Fig. S1. Schematic circular diagram of the *P. thermopropionicum* genome. The outermost circle, predicted protein-coding regions on the plus strand (colors represent functional categories according to the COG database [<http://www.ncbi.nlm.nih.gov/COG>]). The second circle, predicted protein-coding regions on the minus strands. The third circle, tRNAs. The forth circle, rRNAs. The fifth circle, GC skew (red, positive; blue negative).