

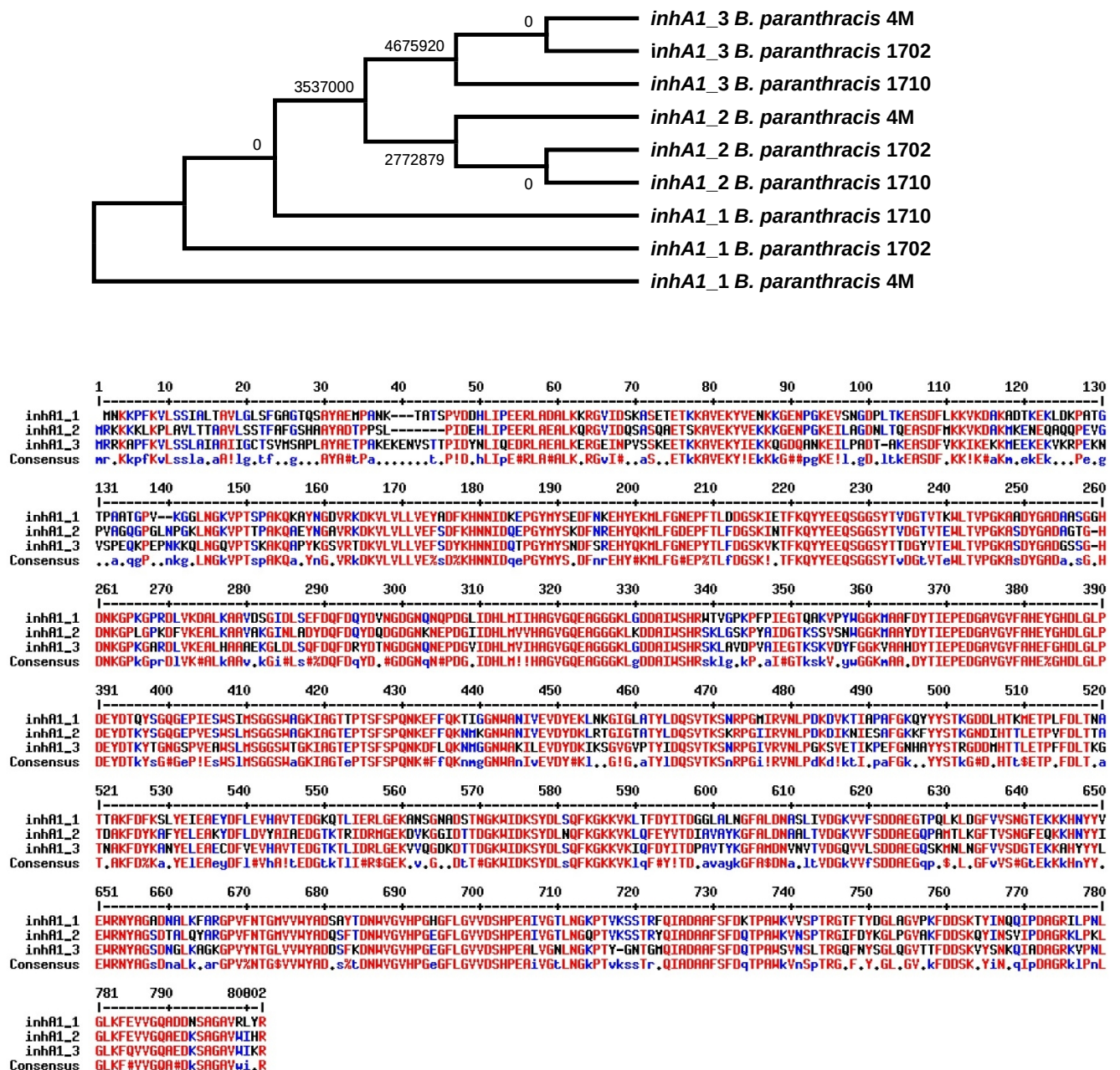


Fig. s1. Distance tree and multiple alignment of the *B. paranthracis* *inhA1* genes.

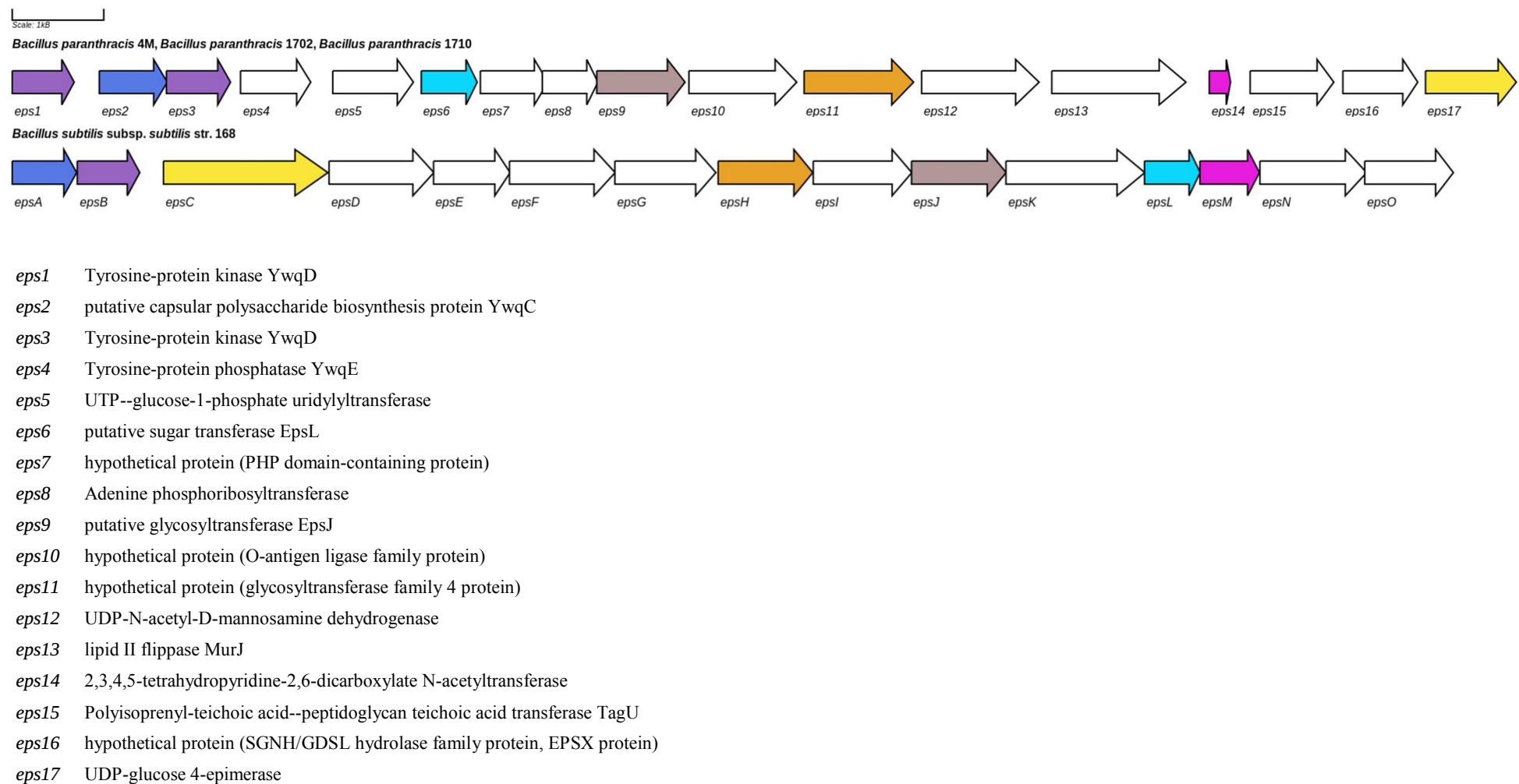


Fig. s2. Comparison of the genetic structures of the *B. paranthracis* *eps1* region and the *B. subtilis* *epsA-O* operon. Orthologous genes are in the same colors. Automatic annotation of the genes in the *eps1* region is shown.