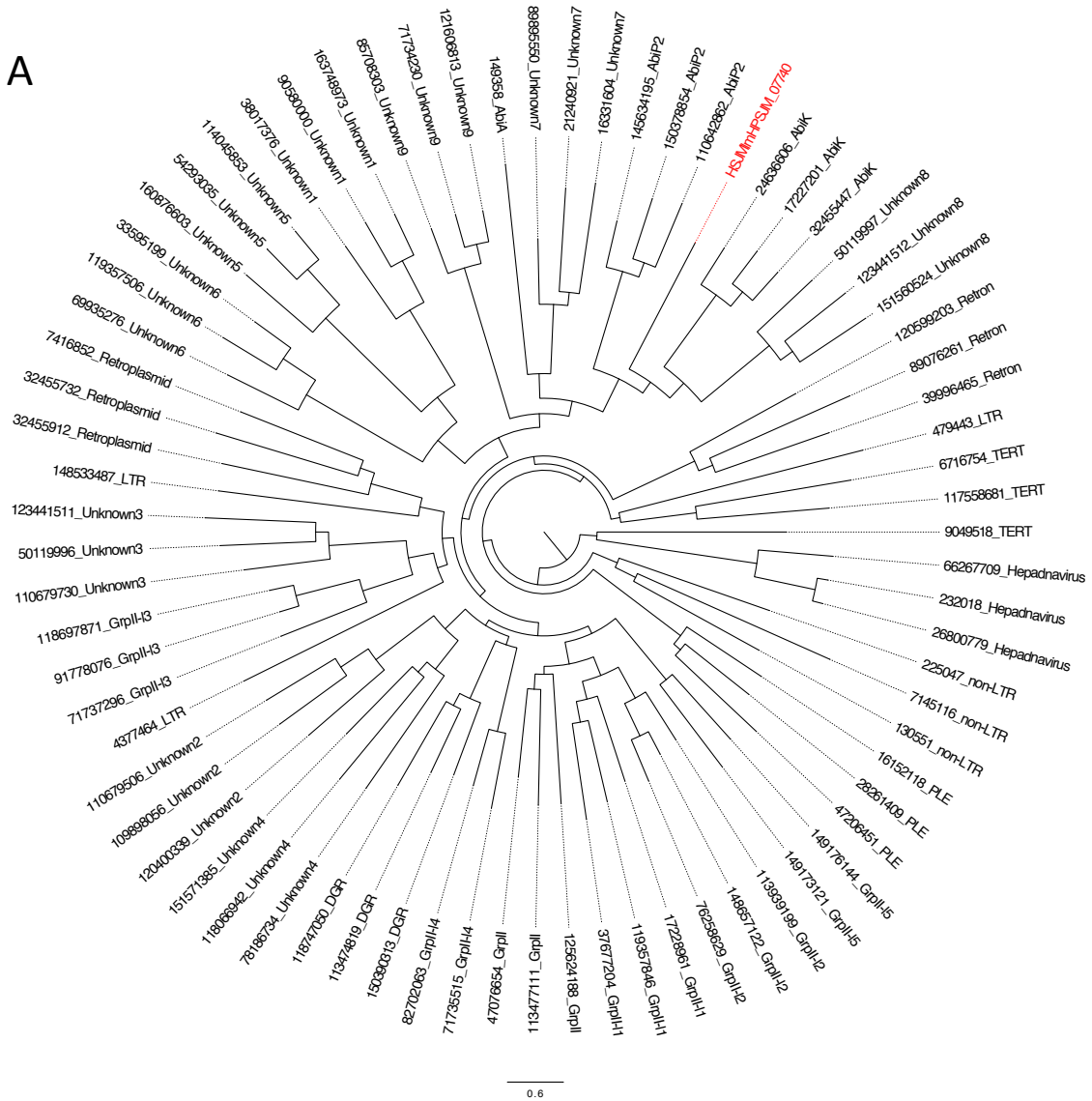
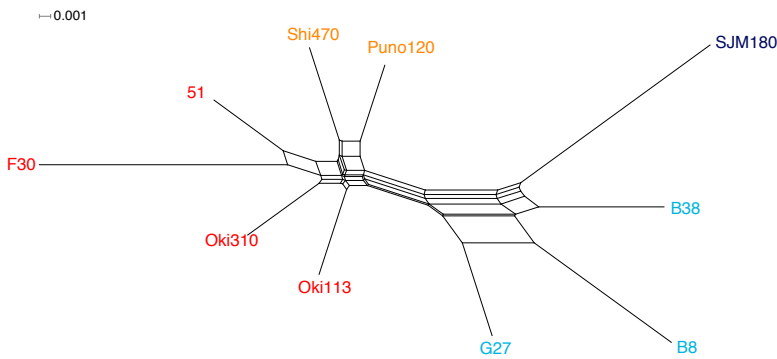


A



B



S4 Fig. (A) Phylogenetic tree of reverse transcriptase homologs including the HPSJM_07740 gene of *H. pylori* SJM180. (B) Phylogenetic network based on the alignment of nucleotide sequences of CGC-10 genes. Strain names are assigned colors according to the phylogeographical groups as in Fig 4.