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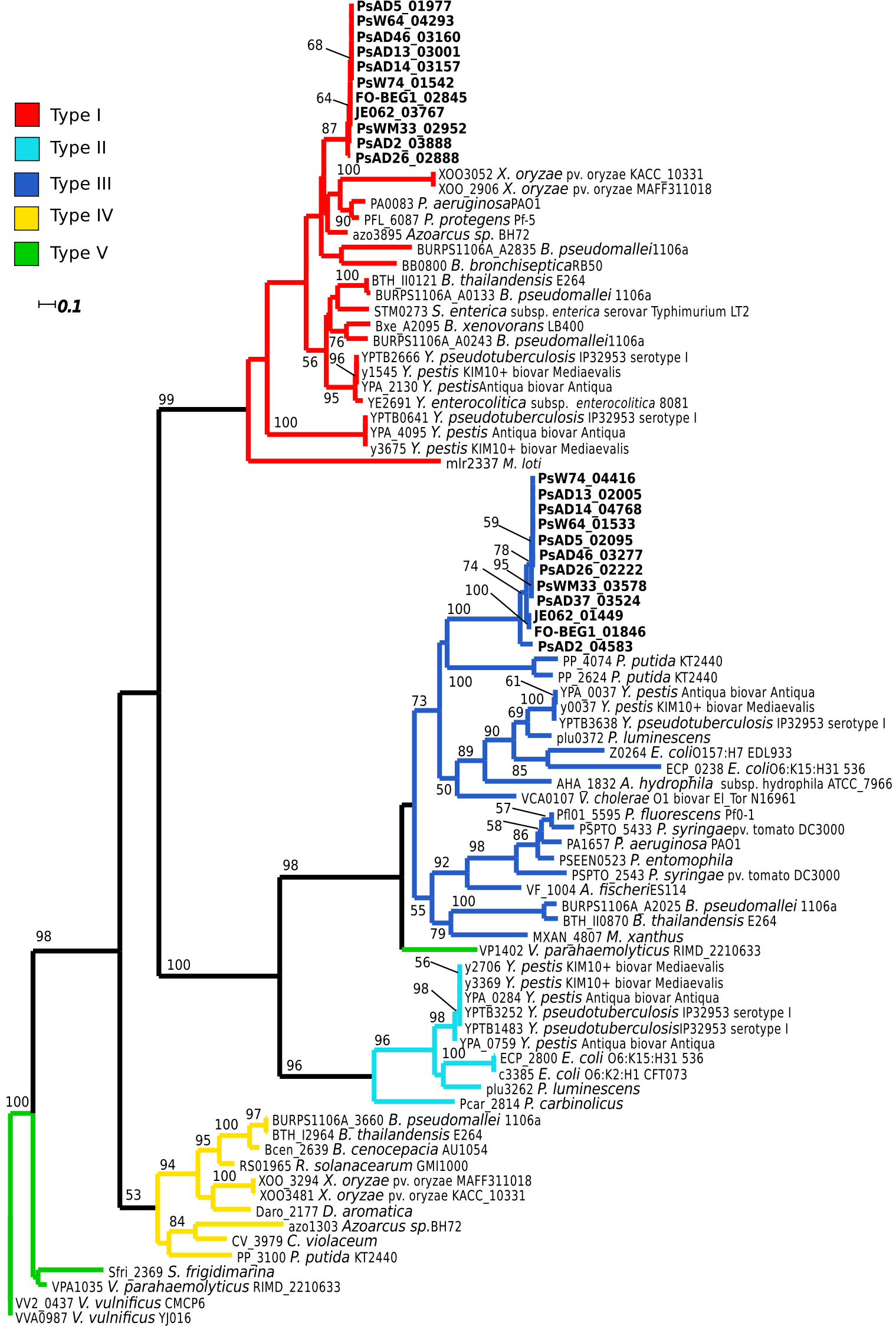


Figure S8 Unrooted phylogenetic tree reconstructed using protein homologous to IgIA belonging to the T6SS KEGG identifiers for each protein are reported. Only Bootstrap values higher than 50 are shown. Color of the branches are based on the T6SS classification reported in Boyer et al. (Boyer et al., 2009).