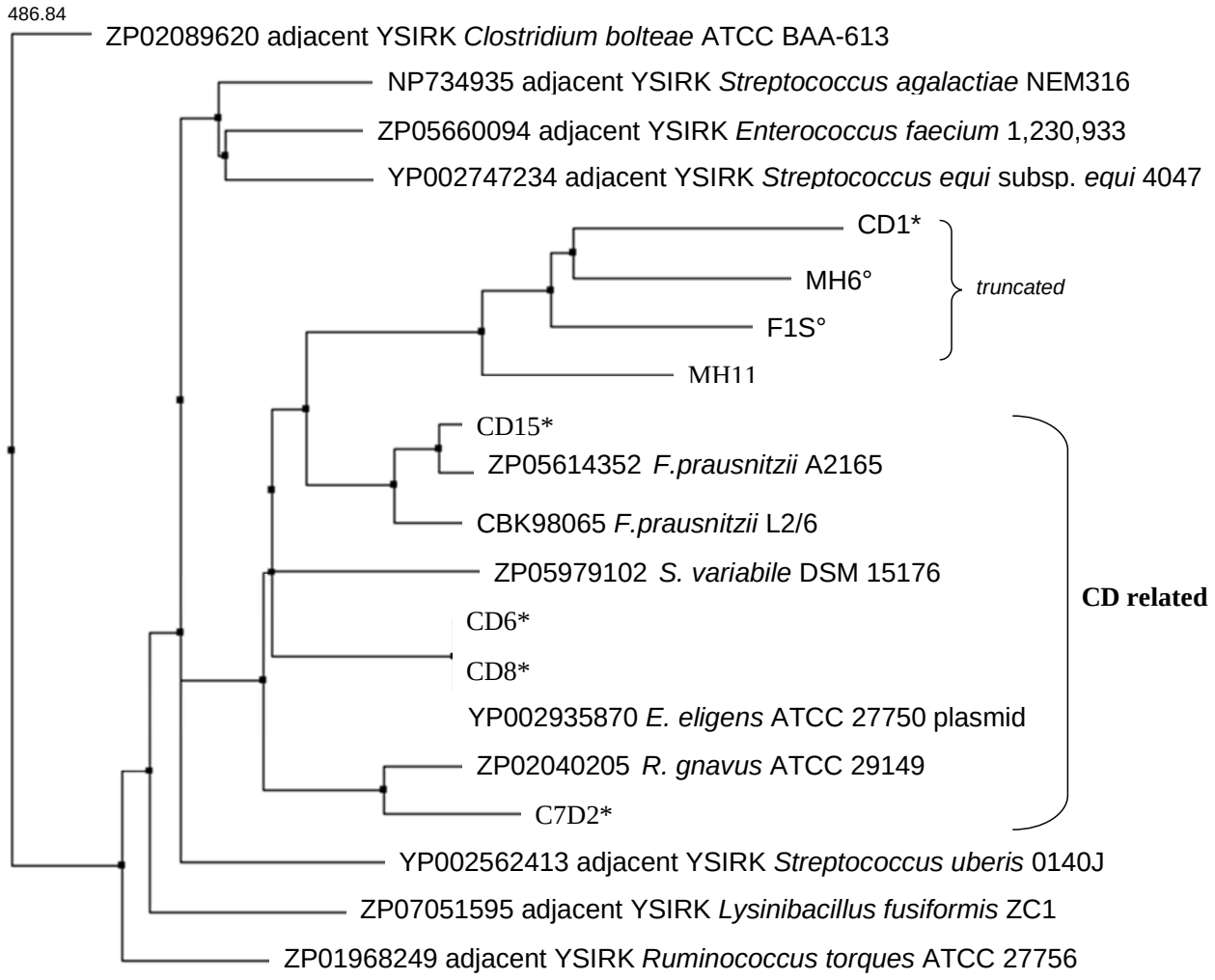
486.84
ZP02089620 adjacent YSIRK *Clostridium bolteae* ATCC BAA-613
NP734935 adjacent YSIRK *Streptococcus agalactiae* NEM316
ZP05660094 adjacent YSIRK *Enterococcus faecium* 1,230,933
YP002747234 adjacent YSIRK *Streptococcus equi* subsp. *equi* 4047
CD1*
MH6°
F1S°
MH11
truncated
CD15*
ZP05614352 *F.prausnitzii* A2165
CBK98065 *F.prausnitzii* L2/6
ZP05979102 *S. variabile* DSM 15176
CD6*
CD8*
YP002935870 *E. eligens* ATCC 27750 plasmid
ZP02040205 *R. gnavus* ATCC 29149
C7D2*
CD related
YP002562413 adjacent YSIRK *Streptococcus uberis* 0140J
ZP07051595 adjacent YSIRK *Lysinibacillus fusiformis* ZC1
ZP01968249 adjacent YSIRK *Ruminococcus torques* ATCC 27756

S6 Figure