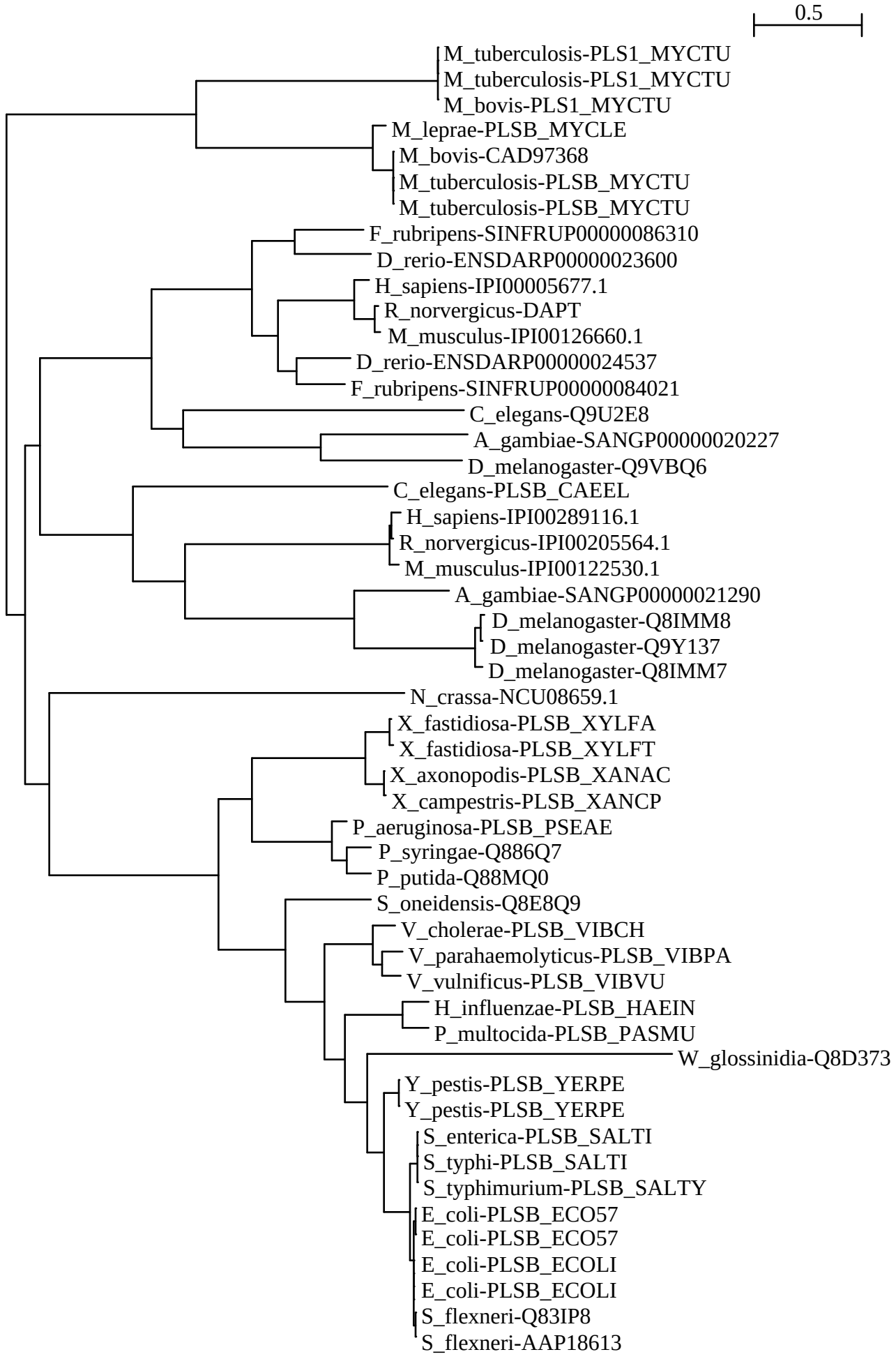
0.5
M_tuberculosis-PLS1_MYCTU
M_tuberculosis-PLS1_MYCTU
M_bovis-PLS1_MYCTU
M_leprae-PLSB_MYCLE
M_bovis-CAD97368
M_tuberculosis-PLSB_MYCTU
M_tuberculosis-PLSB_MYCTU
F_rubripens-SINFRUP00000086310
D_rerio-ENSDARP00000023600
H_sapiens-IPI00005677.1
R_norvergicus-DAPT
M_musculus-IPI00126660.1
D_rerio-ENSDARP00000024537
F_rubripens-SINFRUP00000084021
C_elegans-Q9U2E8
A_gambiae-SANGP00000020227
D_melanogaster-Q9VBQ6
C_elegans-PLSB_CAEEL
H_sapiens-IPI00289116.1
R_norvergicus-IPI00205564.1
M_musculus-IPI00122530.1
A_gambiae-SANGP00000021290
D_melanogaster-Q8IMM8
D_melanogaster-Q9Y137
D_melanogaster-Q8IMM7
N_crassa-NCU08659.1
X_fastidiosa-PLSB_XYLFA
X_fastidiosa-PLSB_XYLFT
X_axonopodis-PLSB_XANAC
X_campestris-PLSB_XANCP
P_aeruginosa-PLSB_PSEAE
P_syringae-Q886Q7
P_putida-Q88MQ0
S_oneidensis-Q8E8Q9
V_cholerae-PLSB_VIBCH
V_parahaemolyticus-PLSB_VIBPA
V_vulnificus-PLSB_VIBVU
H_influenzae-PLSB_HAEIN
P_multocida-PLSB_PASMU
W_glossinidia-Q8D373
Y_pestis-PLSB_YERPE
Y_pestis-PLSB_YERPE
S_enterica-PLSB_SALTI
S_typhi-PLSB_SALTI
S_typhimurium-PLSB_SALTY
E_coli-PLSB_ECO57
E_coli-PLSB_ECO57
E_coli-PLSB_ECOLI
E_coli-PLSB_ECOLI
S_flexneri-Q83IP8
S_flexneri-AAP18613