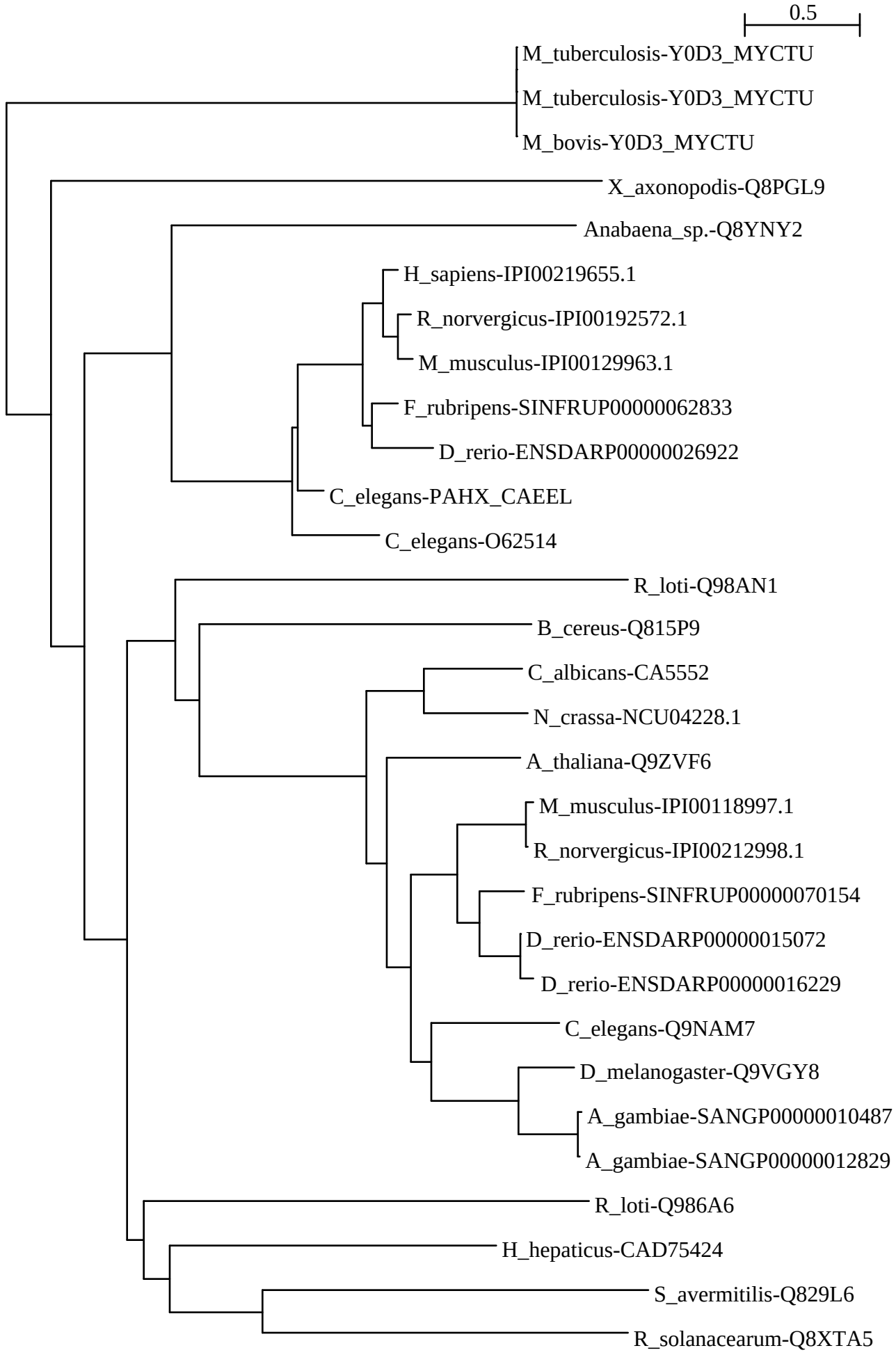
0.5
M_tuberculosis-Y0D3_MYCTU
M_tuberculosis-Y0D3_MYCTU
M_bovis-Y0D3_MYCTU
X_axonopodis-Q8PGL9
Anabaena_sp.-Q8YNY2
H_sapiens-IPI00219655.1
R_norvergicus-IPI00192572.1
M_musculus-IPI00129963.1
F_rubripens-SINFRUP00000062833
D_rerio-ENSDARP00000026922
C_elegans-PAHX_CAEEL
C_elegans-O62514
R_loti-Q98AN1
B_cereus-Q815P9
C_albicans-CA5552
N_crassa-NCU04228.1
A_thaliana-Q9ZVF6
M_musculus-IPI00118997.1
R_norvergicus-IPI00212998.1
F_rubripens-SINFRUP00000070154
D_rerio-ENSDARP00000015072
D_rerio-ENSDARP00000016229
C_elegans-Q9NAM7
D_melanogaster-Q9VGY8
A_gambiae-SANGP00000010487
A_gambiae-SANGP00000012829
R_loti-Q986A6
H_hepaticus-CAD75424
S_avermitilis-Q829L6
R_solanacearum-Q8XTA5