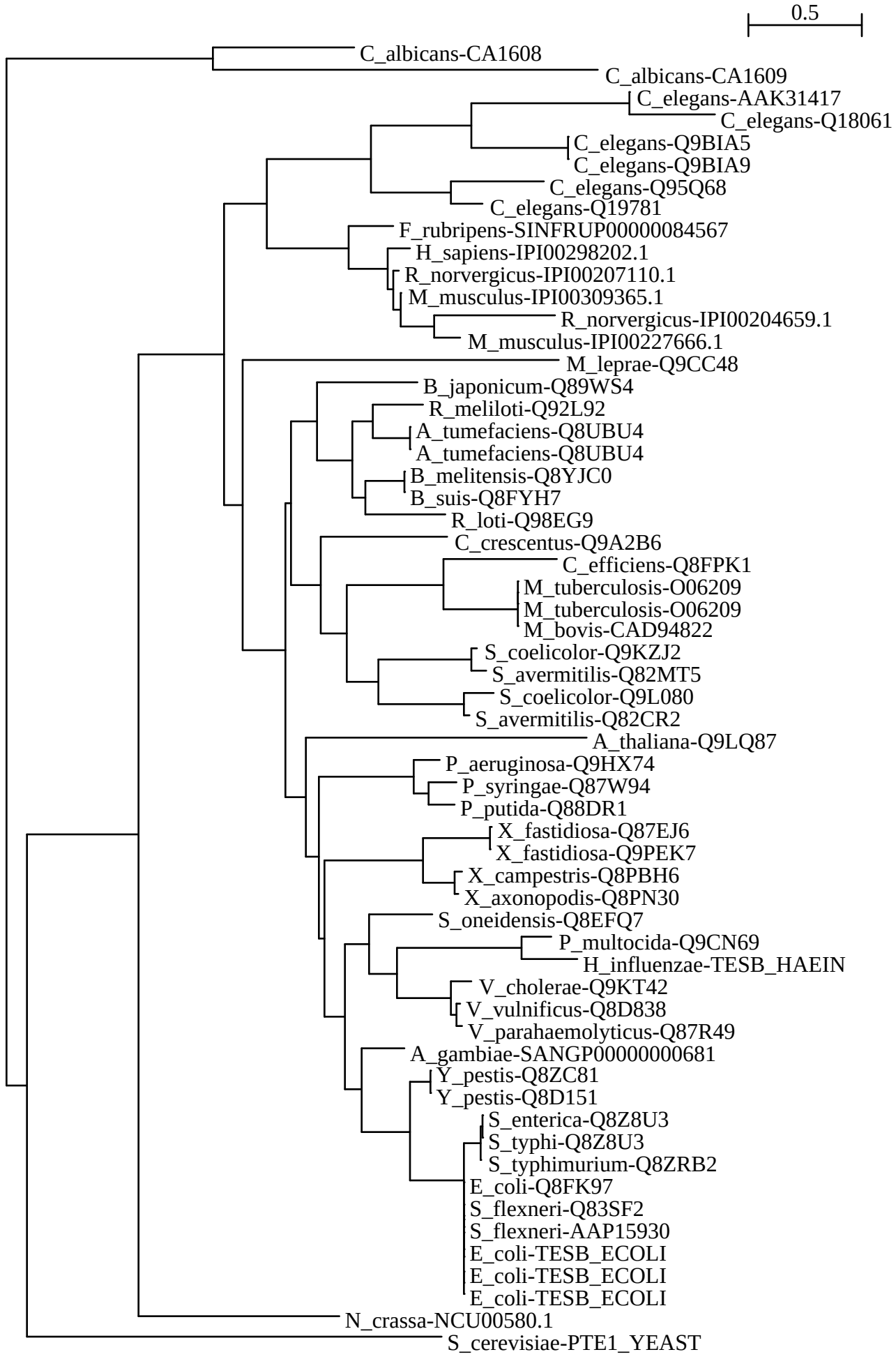
0.5
C_albicans-CA1608
C_albicans-CA1609
C_elegans-AAK31417
C_elegans-Q18061
C_elegans-Q9BIA5
C_elegans-Q9BIA9
C_elegans-Q95Q68
C_elegans-Q19781
F_rubripens-SINFRUP00000084567
H_sapiens-IPI00298202.1
R_norvergicus-IPI00207110.1
M_musculus-IPI00309365.1
R_norvergicus-IPI00204659.1
M_musculus-IPI00227666.1
M_leprae-Q9CC48
B_japonicum-Q89WS4
R_meliloti-Q92L92
A_tumefaciens-Q8UBU4
A_tumefaciens-Q8UBU4
B_melitensis-Q8YJC0
B_suis-Q8FYH7
R_loti-Q98EG9
C_crescentus-Q9A2B6
C_efficiens-Q8FPK1
M_tuberculosis-O06209
M_tuberculosis-O06209
M_bovis-CAD94822
S_coelicolor-Q9KZJ2
S_avermitilis-Q82MT5
S_coelicolor-Q9L080
S_avermitilis-Q82CR2
A_thaliana-Q9LQ87
P_aeruginosa-Q9HX74
P_syringae-Q87W94
P_putida-Q88DR1
X_fastidiosa-Q87EJ6
X_fastidiosa-Q9PEK7
X_campestris-Q8PBH6
X_axonopodis-Q8PN30
S_oneidensis-Q8EFQ7
P_multocida-Q9CN69
H_influenzae-TESB_HAEIN
V_cholerae-Q9KT42
V_vulnificus-Q8D838
V_parahaemolyticus-Q87R49
A_gambiae-SANGP00000000681
Y_pestis-Q8ZC81
Y_pestis-Q8D151
S_enterica-Q8Z8U3
S_typhi-Q8Z8U3
S_typhimurium-Q8ZRB2
E_coli-Q8FK97
S_flexneri-Q83SF2
S_flexneri-AAP15930
E_coli-TESB_ECOLI
E_coli-TESB_ECOLI
E_coli-TESB_ECOLI
N_crassa-NCU00580.1
S_cerevisiae-PTE1_YEAST