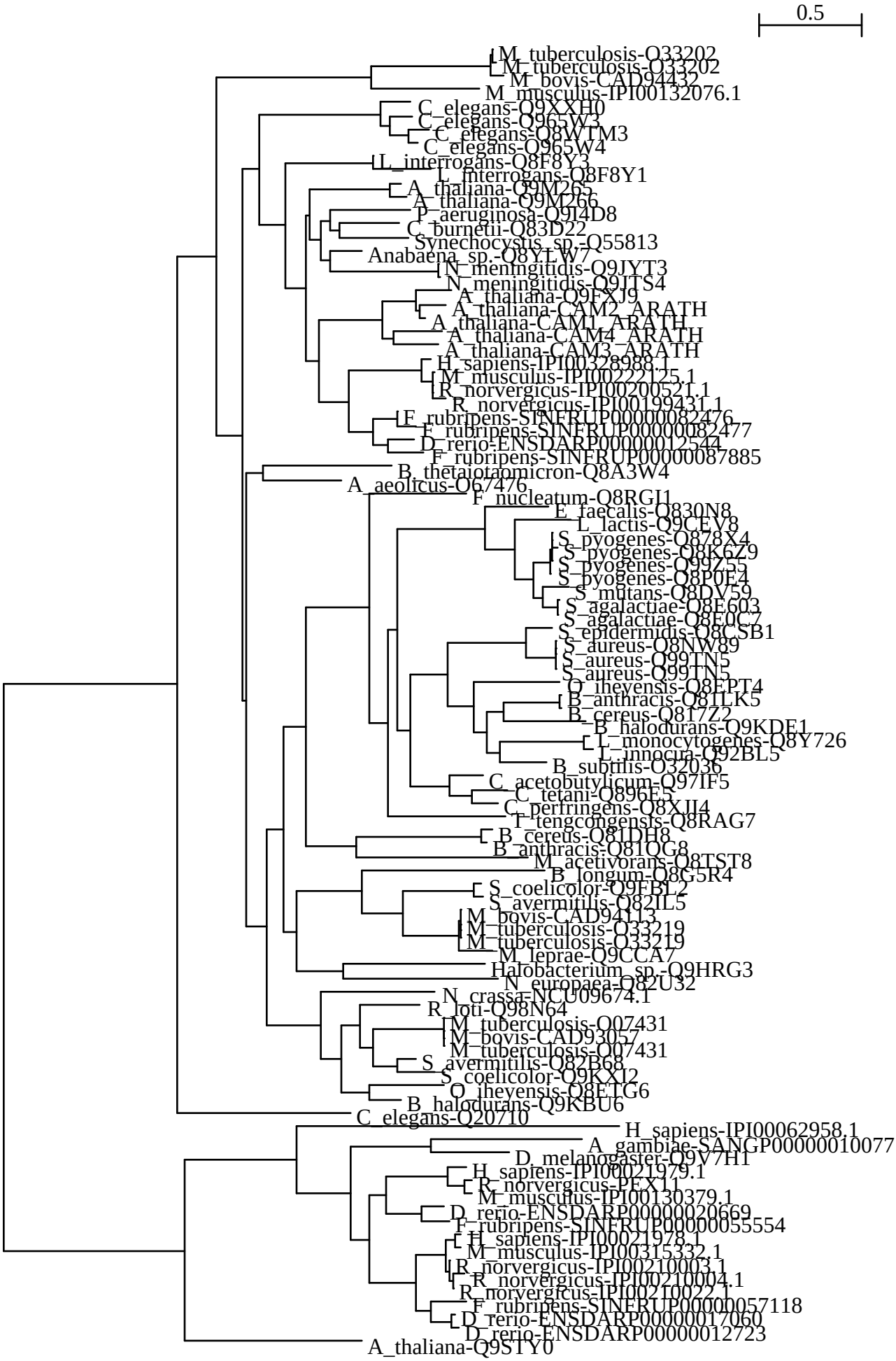
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M_tuberculosis-O33202
M_tuberculosis-O33202
M_bovis-CAD94432
M_musculus-IPI00132076.1
C_elegans-Q9XXH0
C_elegans-Q965W3
C_elegans-Q8WTM3
C_elegans-Q965W4
L_interrogans-Q8F8Y3
L_interrogans-Q8F8Y1
A_thaliana-Q9M265
A_thaliana-Q9M266
P_aeruginosa-Q9I4D8
C_burnetii-Q83D22
Synechocystis_sp.-Q55813
Anabaena_sp.-Q8YLW7
N_meningitidis-Q9JYT3
N_meningitidis-Q9JTS4
A_thaliana-Q9FXJ9
A_thaliana-CAM2_ARATH
A_thaliana-CAM1_ARATH
A_thaliana-CAM4_ARATH
A_thaliana-CAM3_ARATH
H_sapiens-IPI00328988.1
M_musculus-IPI00222125.1
R_norvergicus-IPI00200521.1
R_norvergicus-IPI00199431.1
F_rubripens-SINFRUP00000082476
F_rubripens-SINFRUP00000082477
D_rerio-ENSDARP00000012544
F_rubripens-SINFRUP00000087885
B_thetaiotaomicron-Q8A3W4
A_aeolicus-O67476
F_nucleatum-Q8RGI1
E_faecalis-Q830N8
L_lactis-Q9CEV8
S_pyogenes-Q878X4
S_pyogenes-Q8K6Z9
S_pyogenes-Q99Z55
S_pyogenes-Q8P0E4
S_mutans-Q8DV59
S_agalactiae-Q8E603
S_agalactiae-Q8E0C7
S_epidermidis-Q8CSB1
S_aureus-Q8NW89
S_aureus-Q99TN5
S_aureus-Q99TN5
O_iheyensis-Q8EPT4
B_anthracis-Q81LK5
B_cereus-Q817Z2
B_halodurans-Q9KDE1
L_monocytogenes-Q8Y726
L_innocua-Q92BL5
B_subtilis-O32036
C_acetobutylicum-Q97IF5
C_tetani-Q896E5
C_perfringens-Q8XJI4
T_tengcongensis-Q8RAG7
B_cereus-Q81DH8
B_anthracis-Q81QG8
M_acetivorans-Q8TST8
B_longum-Q8G5R4
S_coelicolor-Q9FBL2
S_avermitilis-Q82IL5
M_bovis-CAD94113
M_tuberculosis-O33219
M_tuberculosis-O33219
M_leprae-Q9CCA7
Halobacterium_sp.-Q9HRG3
N_europaea-Q82U32
N_crassa-NCU09674.1
R_loti-Q98N64
M_tuberculosis-O07431
M_bovis-CAD93057
M_tuberculosis-O07431
S_avermitilis-Q82B68
S_coelicolor-Q9KXI2
O_iheyensis-Q8ETG6
B_halodurans-Q9KBU6
C_elegans-Q20710
H_sapiens-IPI00062958.1
A_gambiae-SANGP00000010077
D_melanogaster-Q9V7H1
H_sapiens-IPI00021979.1
R_norvergicus-PEX11
M_musculus-IPI00130379.1
D_rerio-ENSDARP00000020669
F_rubripens-SINFRUP00000055554
H_sapiens-IPI00021978.1
M_musculus-IPI00315332.1
R_norvergicus-IPI00210003.1
R_norvergicus-IPI00210004.1
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F_rubripens-SINFRUP00000057118
D_rerio-ENSDARP00000017060
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A_thaliana-Q9STY0