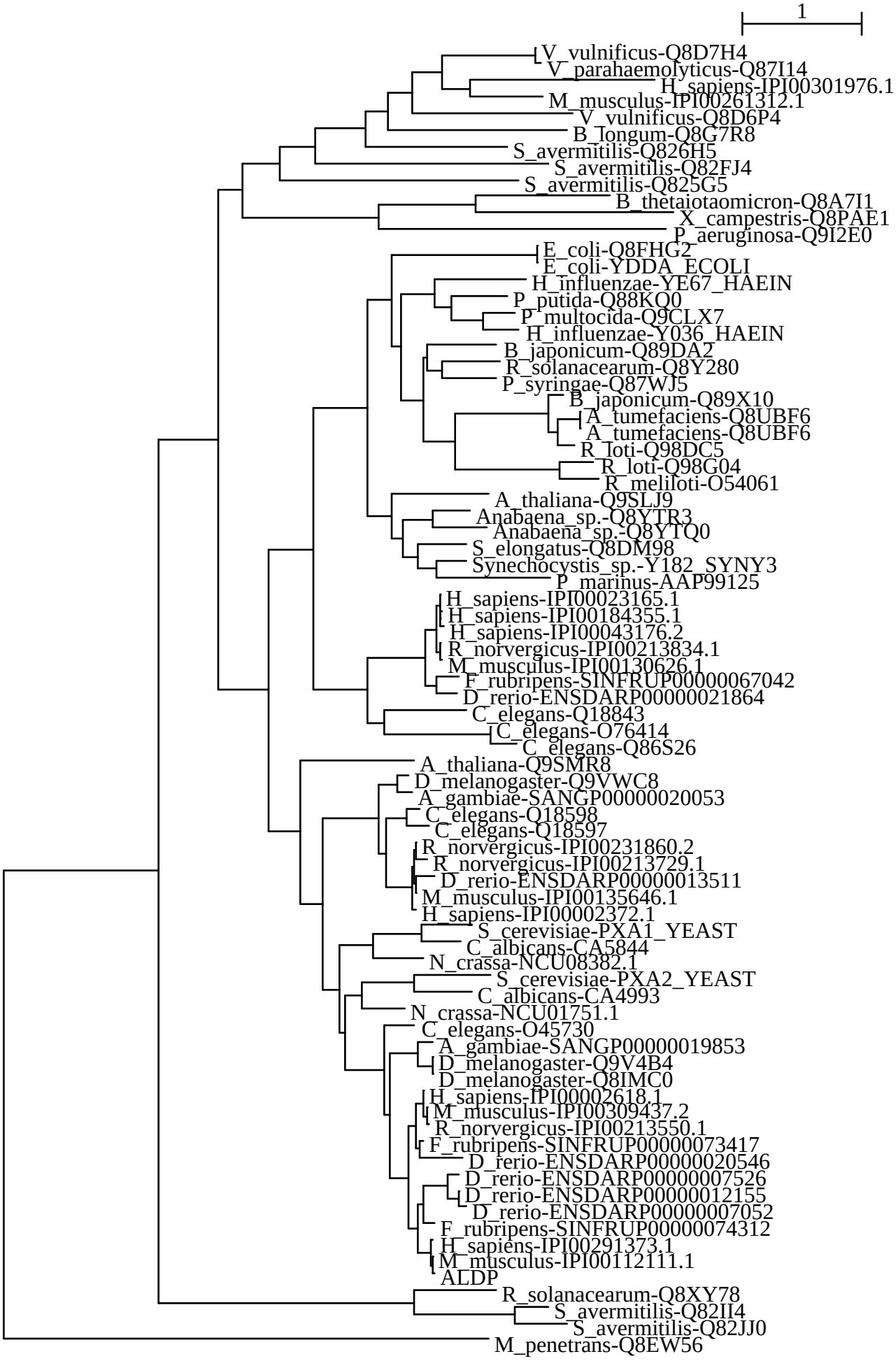
1
V_vulnificus-Q8D7H4
V_parahaemolyticus-Q87I14
H_sapiens-IPI00301976.1
M_musculus-IPI00261312.1
V_vulnificus-Q8D6P4
B_longum-Q8G7R8
S_avermitilis-Q826H5
S_avermitilis-Q82FJ4
S_avermitilis-Q825G5
B_thetaiotaomicron-Q8A7I1
X_campestris-Q8PAE1
P_aeruginosa-Q9I2E0
E_coli-Q8FHG2
E_coli-YDDA_ECOLI
H_influenzae-YE67_HAEIN
P_putida-Q88KQ0
P_multocida-Q9CLX7
H_influenzae-Y036_HAEIN
B_japonicum-Q89DA2
R_solanacearum-Q8Y280
P_syringae-Q87WJ5
B_japonicum-Q89X10
A_tumefaciens-Q8UBF6
A_tumefaciens-Q8UBF6
R_loti-Q98DC5
R_loti-Q98G04
R_meliloti-O54061
A_thaliana-Q9SLJ9
Anabaena_sp.-Q8YTR3
Anabaena_sp.-Q8YTQ0
S_elongatus-Q8DM98
Synechocystis_sp.-Y182_SYNY3
P_marinus-AAP99125
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M_musculus-IPI00130626.1
F_rubripens-SINFRUP00000067042
D_rerio-ENSDARP00000021864
C_elegans-Q18843
C_elegans-O76414
C_elegans-Q86S26
A_thaliana-Q9SMR8
D_melanogaster-Q9VWC8
A_gambiae-SANGP00000020053
C_elegans-Q18598
C_elegans-Q18597
R_norvergicus-IPI00231860.2
R_norvergicus-IPI00213729.1
D_rerio-ENSDARP00000013511
M_musculus-IPI00135646.1
H_sapiens-IPI00002372.1
S_cerevisiae-PXA1_YEAST
C_albicans-CA5844
N_crassa-NCU08382.1
S_cerevisiae-PXA2_YEAST
C_albicans-CA4993
N_crassa-NCU01751.1
C_elegans-O45730
A_gambiae-SANGP00000019853
D_melanogaster-Q9V4B4
D_melanogaster-Q8IMC0
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M_musculus-IPI00309437.2
R_norvergicus-IPI00213550.1
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H_sapiens-IPI00291373.1
M_musculus-IPI00112111.1
ALDP
R_solanacearum-Q8XY78
S_avermitilis-Q82II4
S_avermitilis-Q82JJ0
M_penetrans-Q8EW56