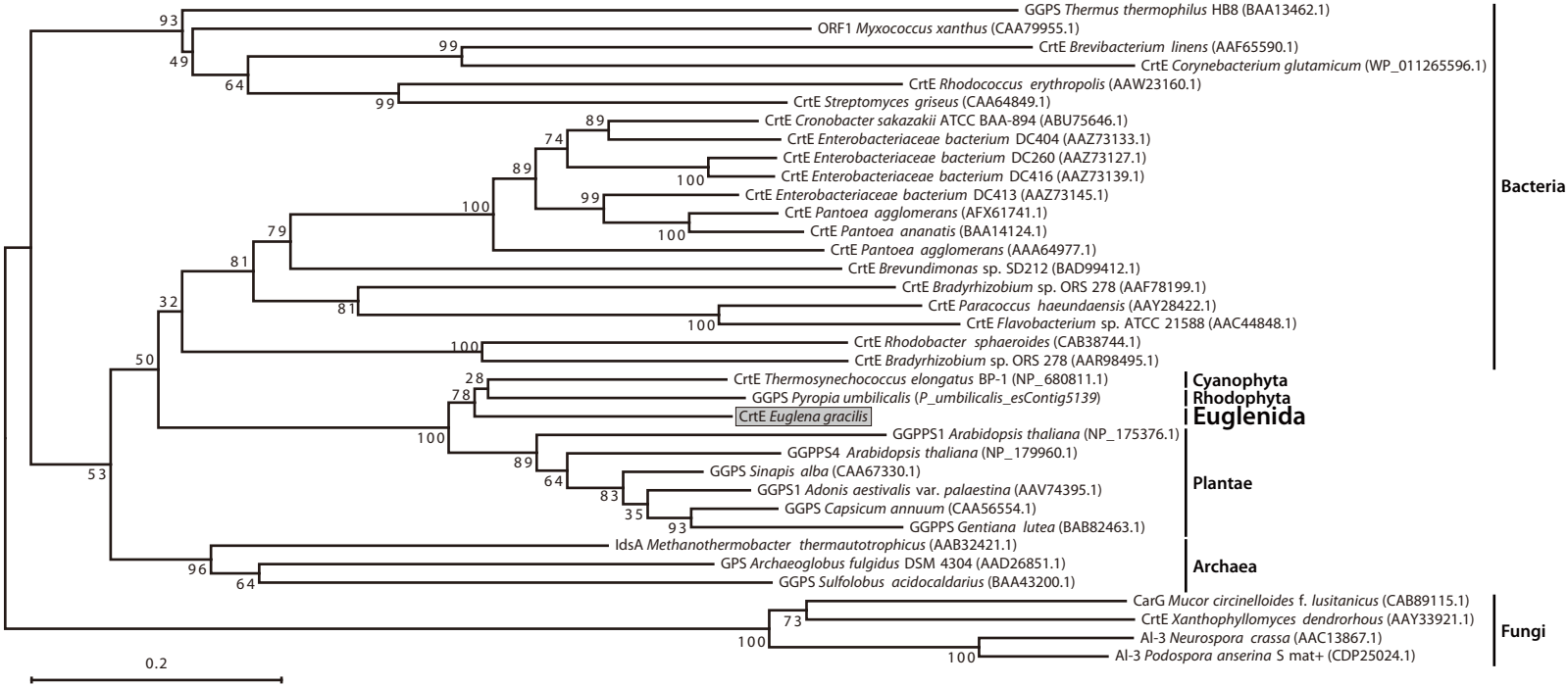
GGPS Thermus thermophilus HB8 (BAA13462.1)
ORF1 Myxococcus xanthus (CAA79955.1)
CrtE Brevibacterium linens (AAF65590.1)
CrtE Corynebacterium glutamicum (WP_011265596.1)
CrtE Rhodococcus erythropolis (AAW23160.1)
CrtE Streptomyces griseus (CAA64849.1)
CrtE Cronobacter sakazakii ATCC BAA-894 (ABU75646.1)
CrtE Enterobacteriaceae bacterium DC404 (AAZ73133.1)
CrtE Enterobacteriaceae bacterium DC260 (AAZ73127.1)
CrtE Enterobacteriaceae bacterium DC416 (AAZ73139.1)
CrtE Enterobacteriaceae bacterium DC413 (AAZ73145.1)
CrtE Pantoea agglomerans (AFX61741.1)
CrtE Pantoea ananatis (BAA14124.1)
CrtE Pantoea agglomerans (AAA64977.1)
CrtE Brevundimonas sp. SD212 (BAD99412.1)
CrtE Bradyrhizobium sp. ORS 278 (AAF78199.1)
CrtE Paracoccus haeundaensis (AAY28422.1)
CrtE Flavobacterium sp. ATCC 21588 (AAC44848.1)
CrtE Rhodobacter sphaeroides (CAB38744.1)
CrtE Bradyrhizobium sp. ORS 278 (AAR98495.1)
CrtE Thermosynechococcus elongatus BP-1 (NP_680811.1)
GGPS Pyropia umbilicalis (P_umbilicalis_esContig5139)
CrtE Euglena gracilis
GGPPS1 Arabidopsis thaliana (NP_175376.1)
GGPPS4 Arabidopsis thaliana (NP_179960.1)
GGPS Sinapis alba (CAA67330.1)
GGPS1 Adonis aestivalis var. palaestina (AAV74395.1)
GGPS Capsicum annuum (CAA56554.1)
GGPPS Gentiana lutea (BAB82463.1)
IdsA Methanothermobacter thermautotrophicus (AAB32421.1)
GPS Archaeoglobus fulgidus DSM 4304 (AAD26851.1)
GGPS Sulfolobus acidocaldarius (BAA43200.1)
CarG Mucor circinelloides f. lusitanicus (CAB89115.1)
CrtE Xanthophyllomyces dendrorhous (AAY33921.1)
Al-3 Neurospora crassa (AAC13867.1)
Al-3 Podospora anserina S mat+ (CDP25024.1)
Bacteria
Cyanophyta
Rhodophyta
Euglenida
Plantae
Archaea
Fungi
0.2