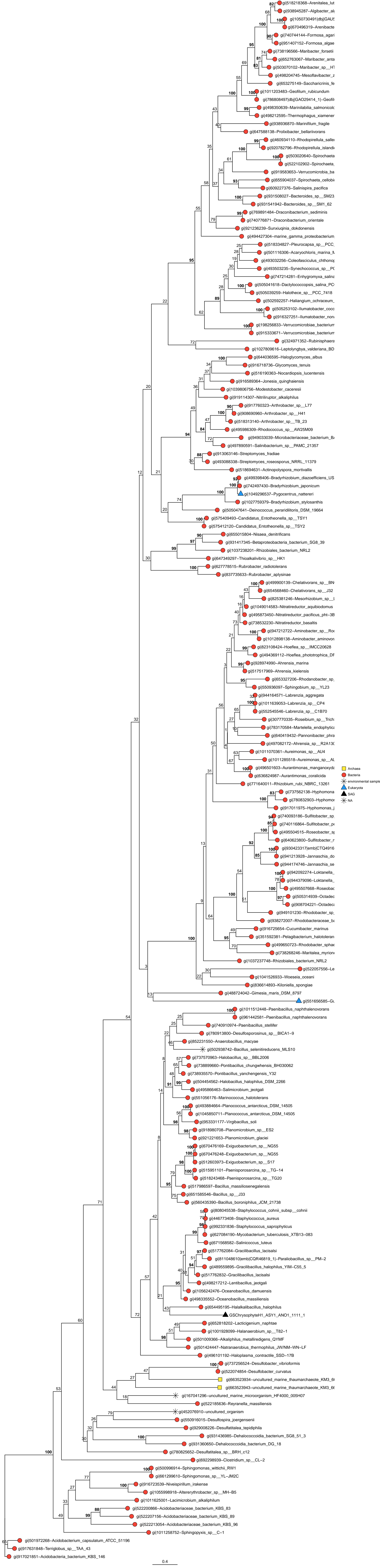
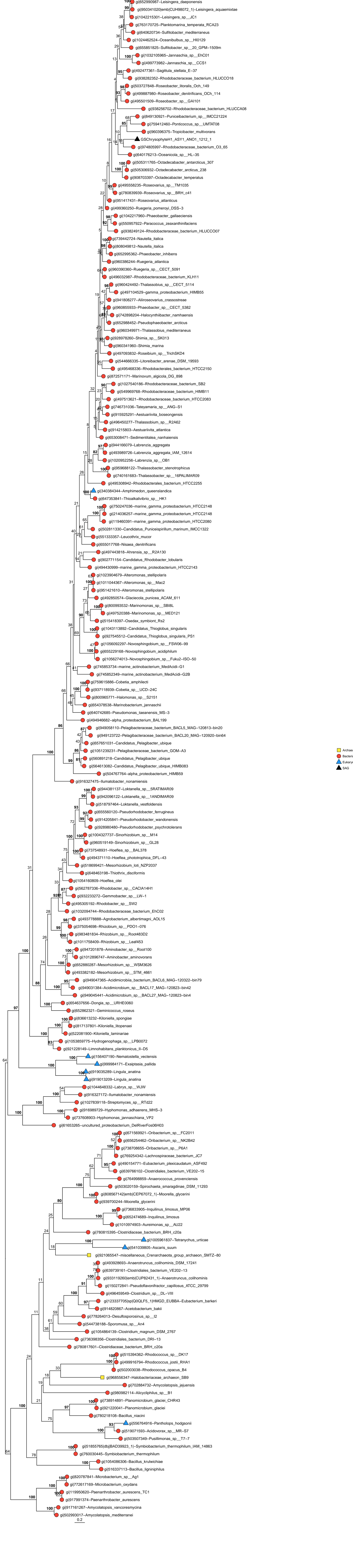
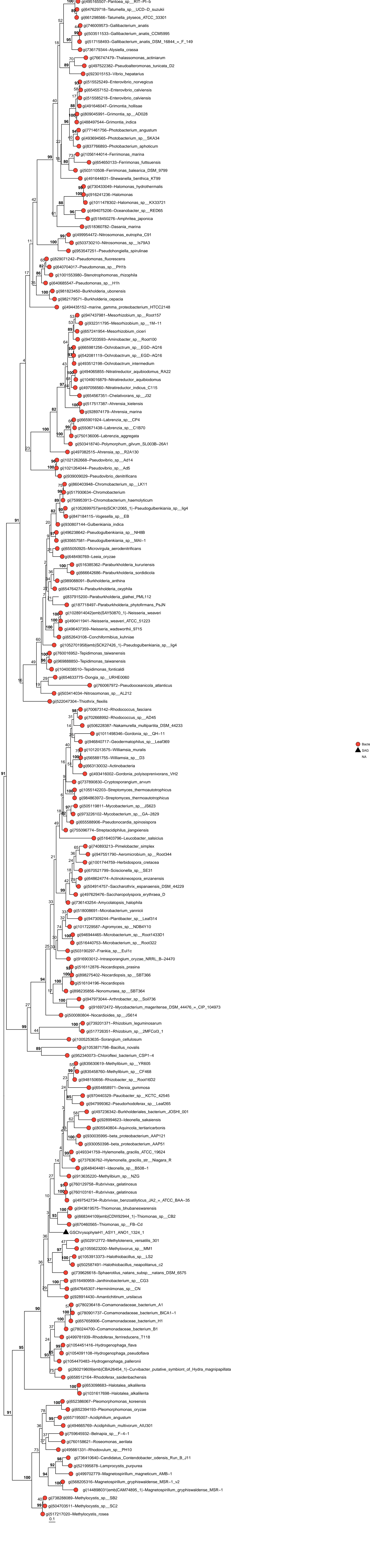


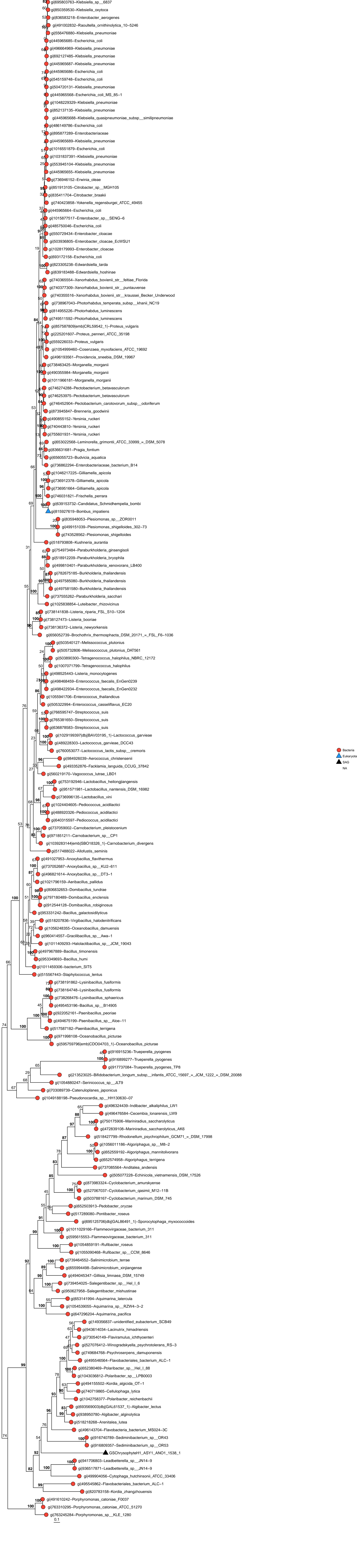
● Bacteria
▲ SAG

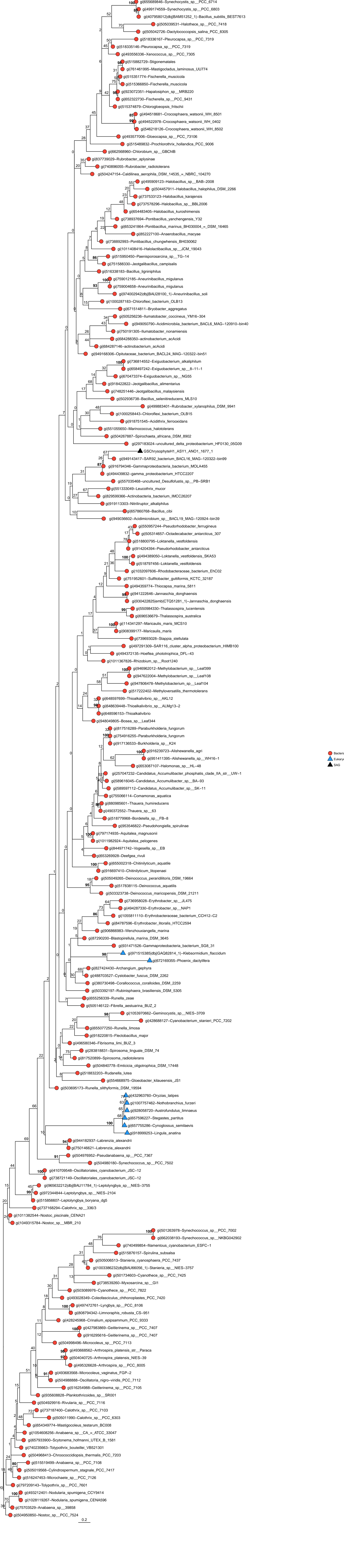
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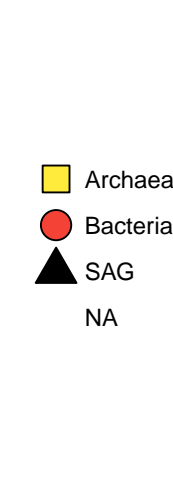
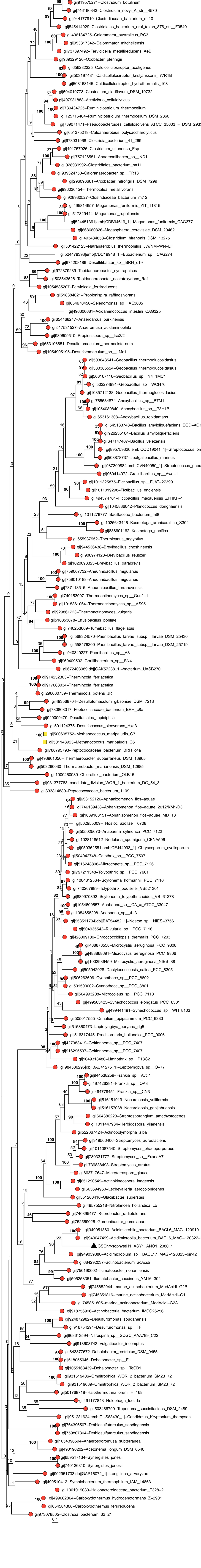


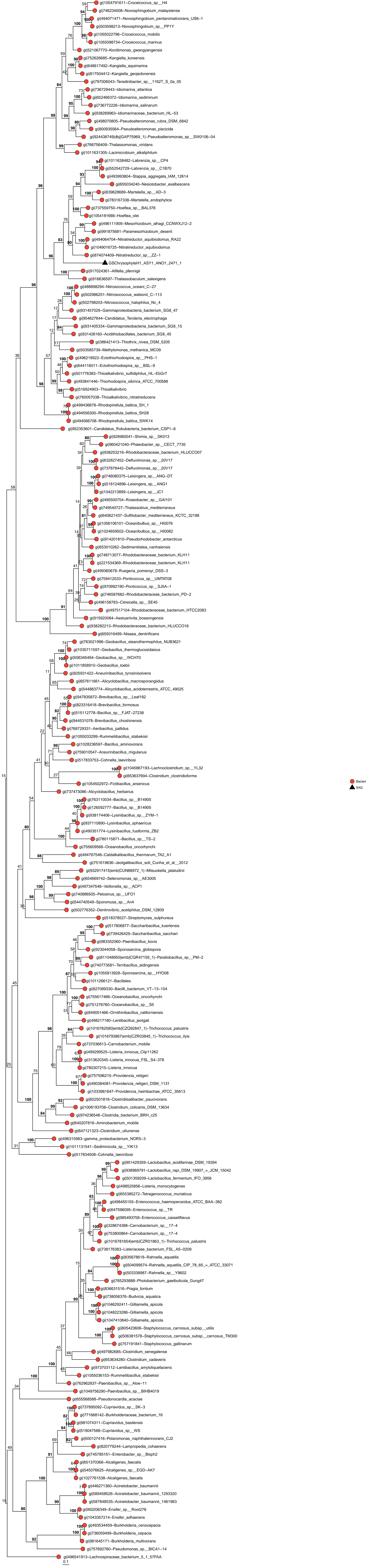


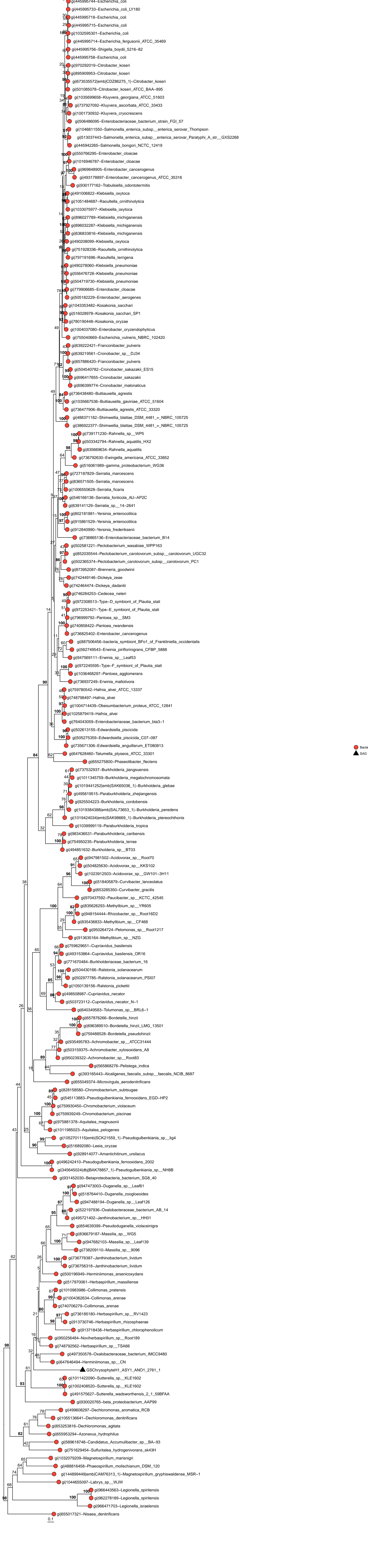


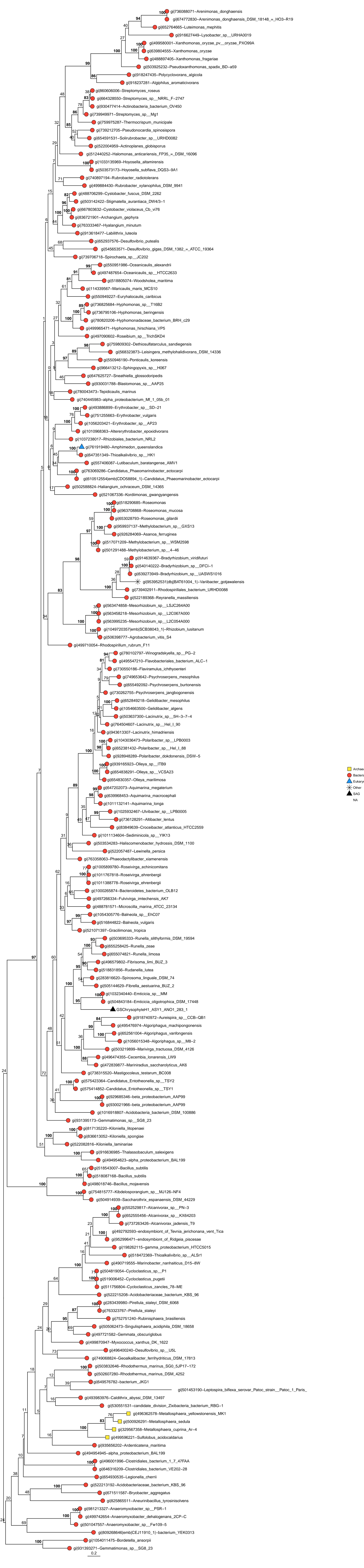


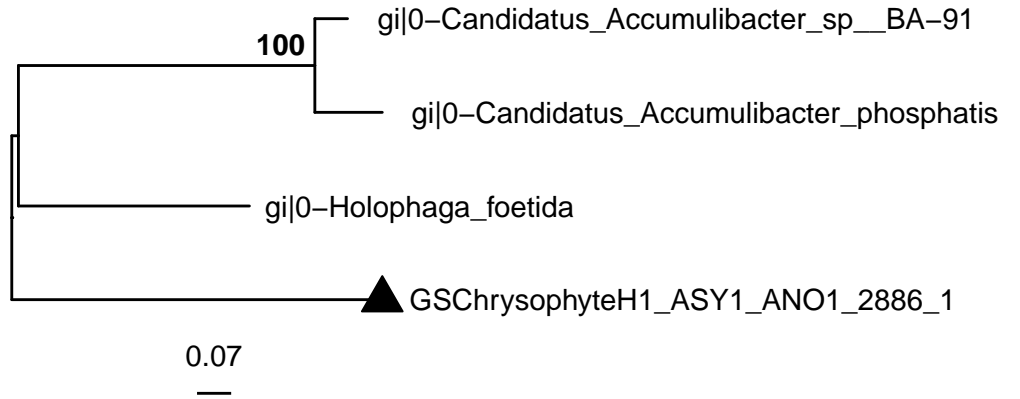




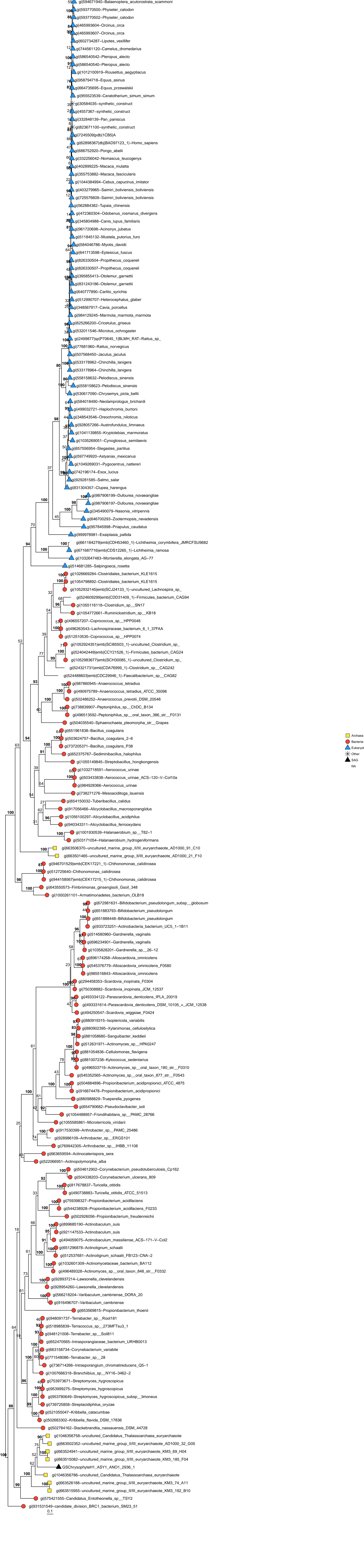


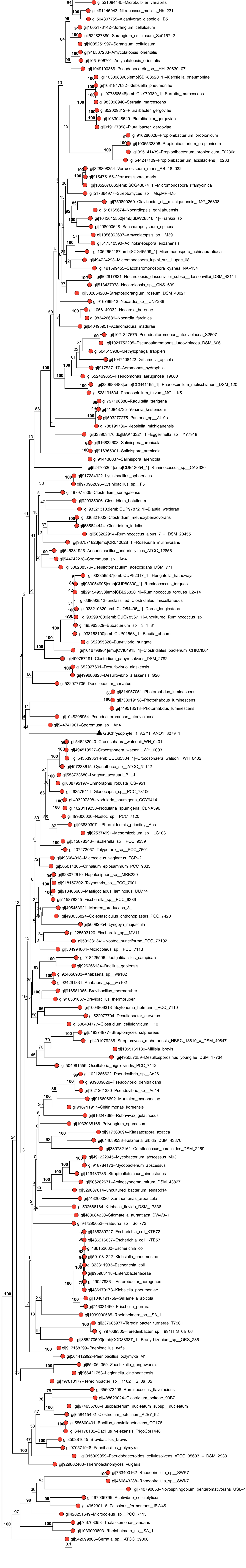




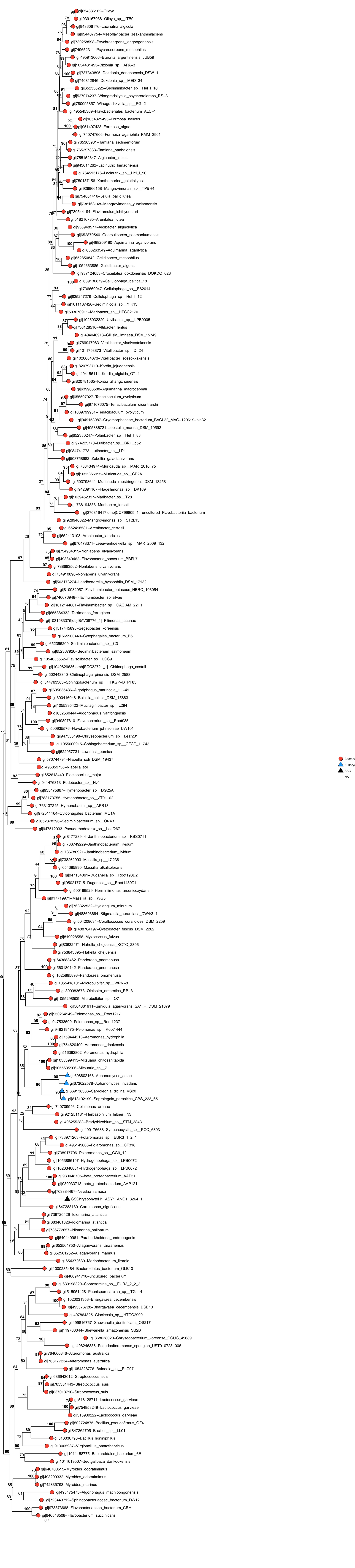


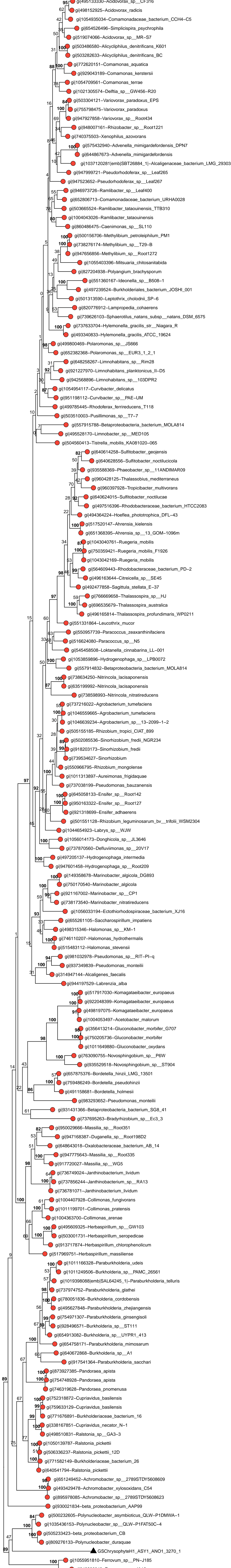
▲ SAG
NA



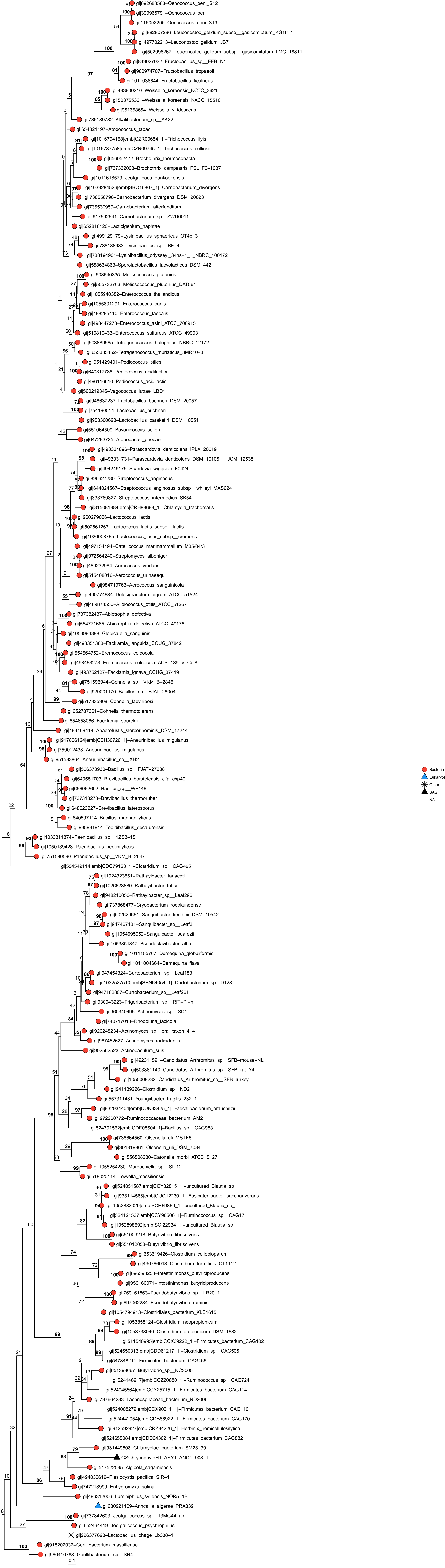


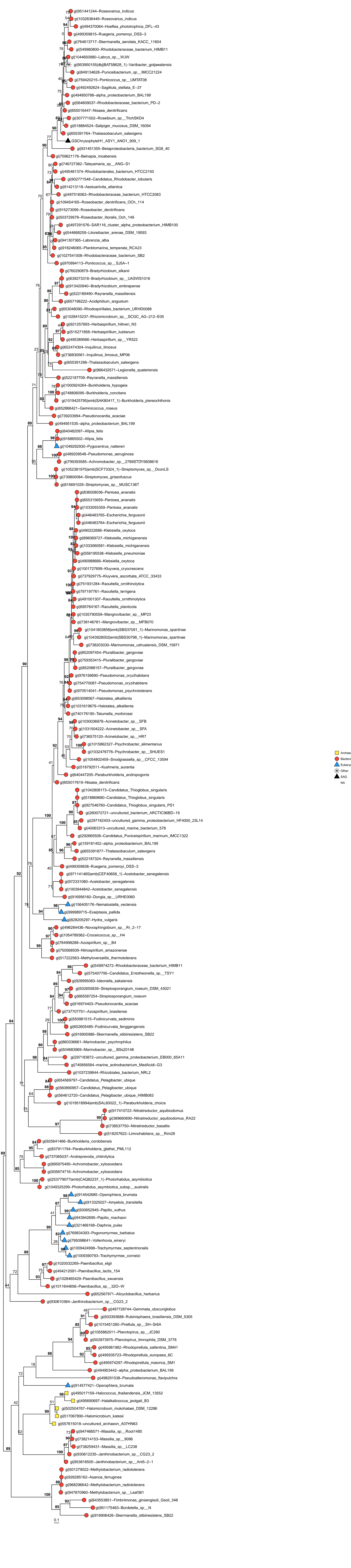
● Bacteria
▲ SAG
NA

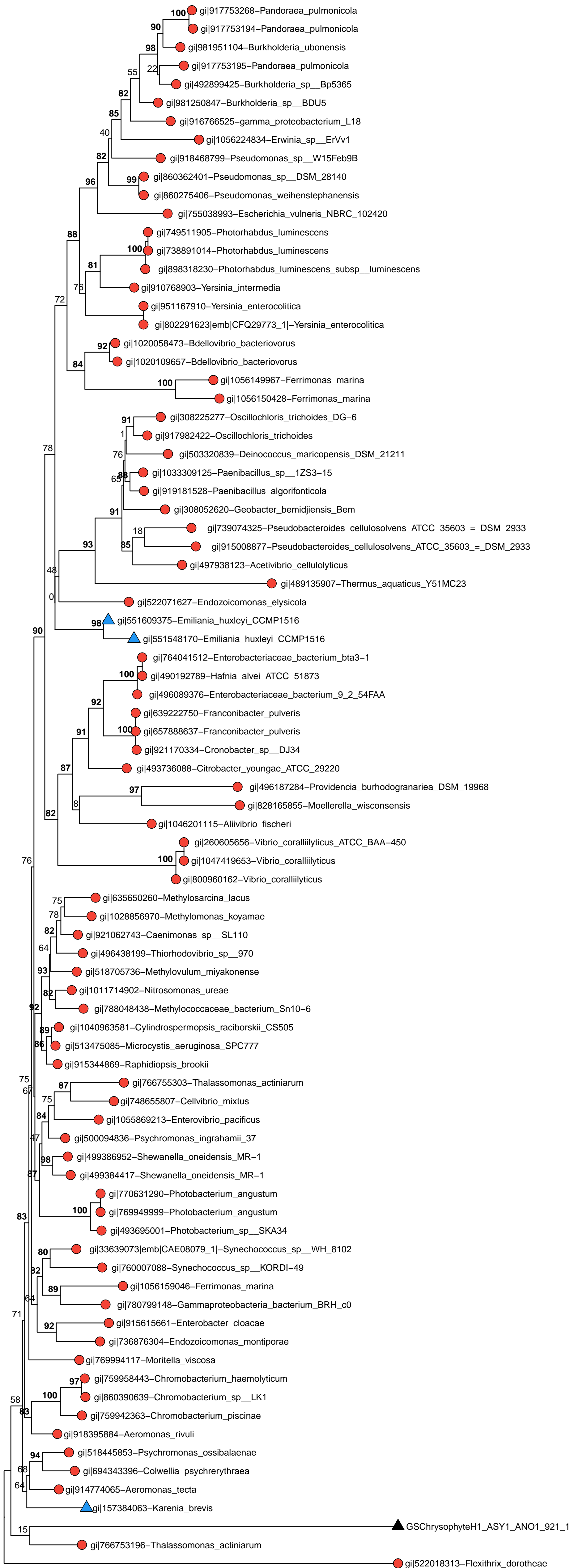




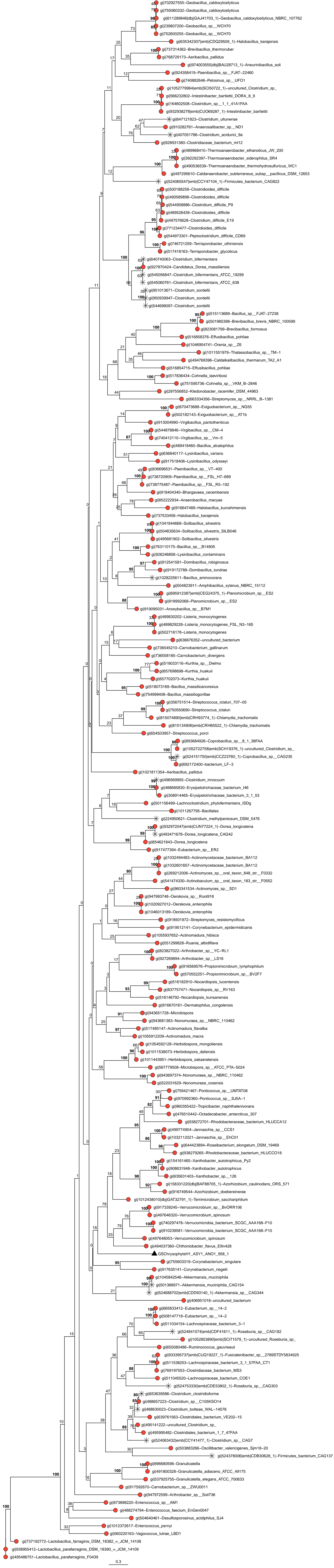
● Bacteria
▲ SAG

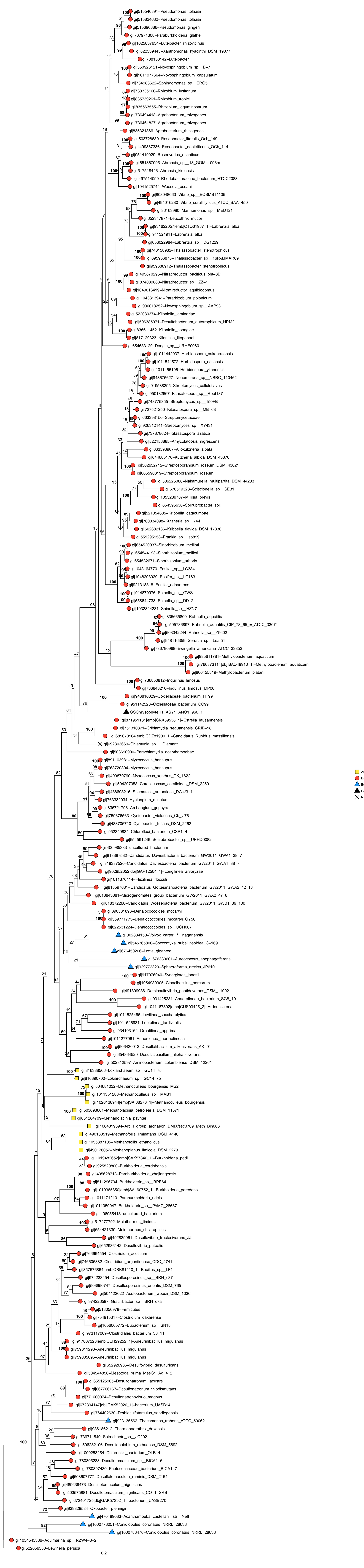


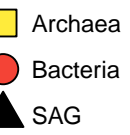
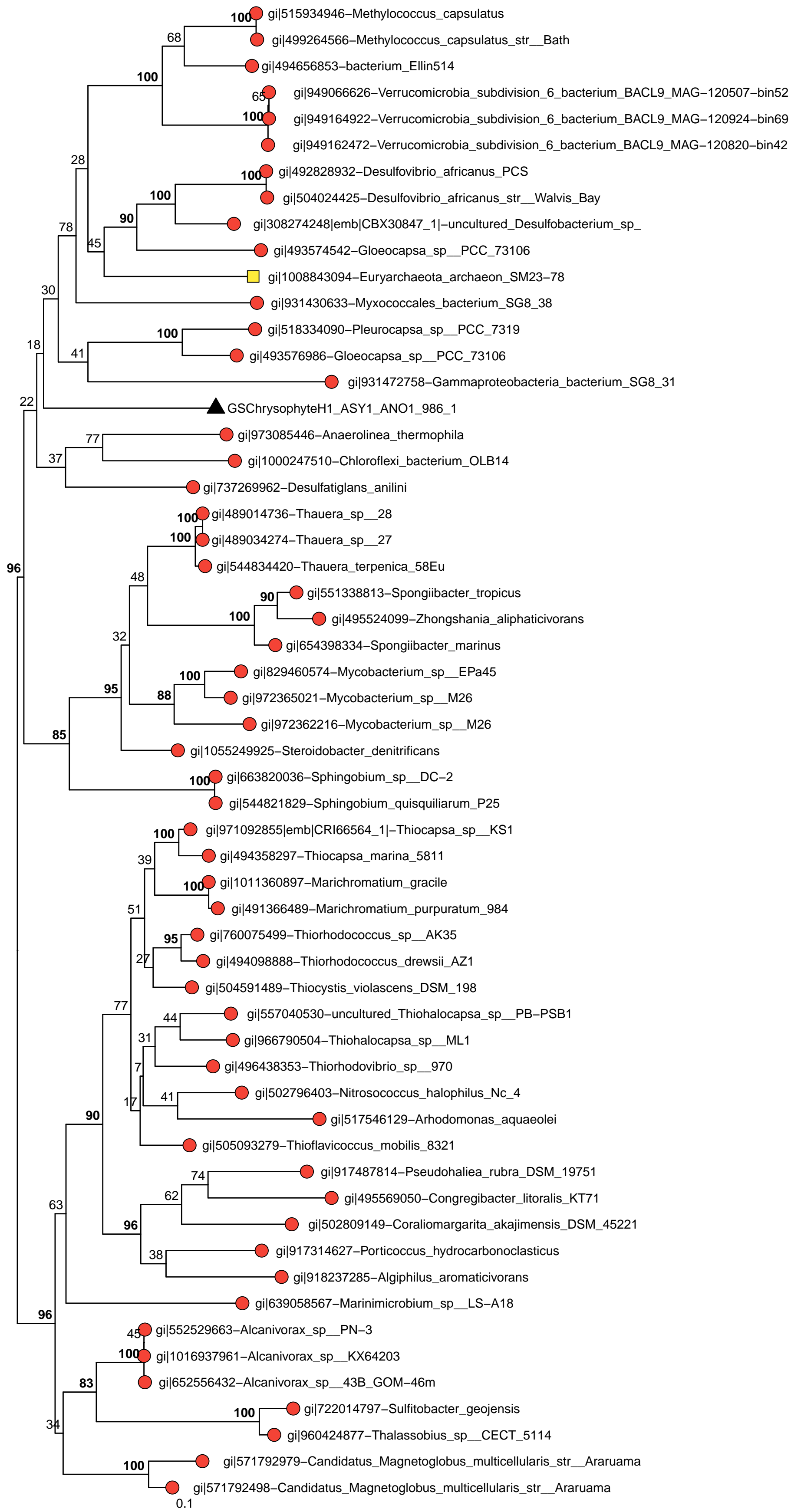


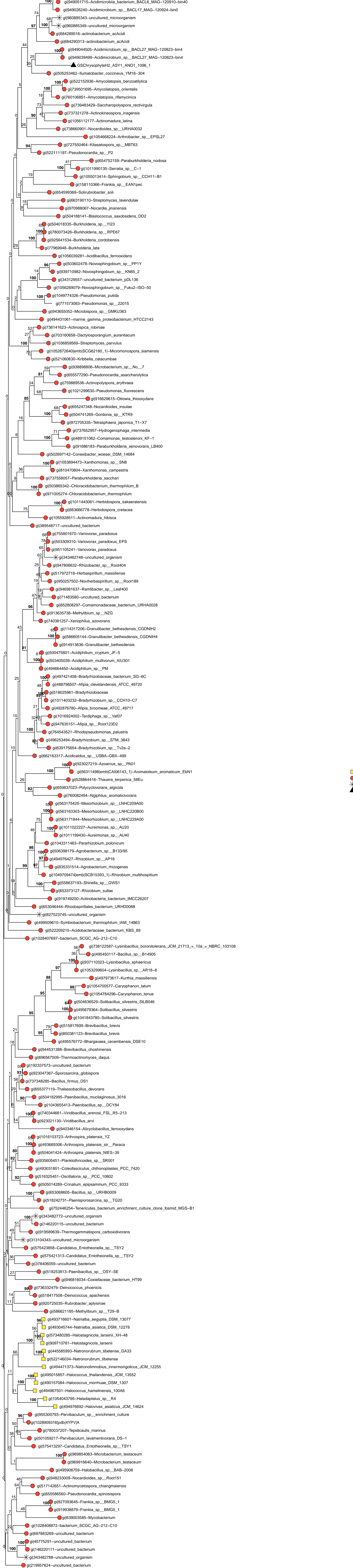


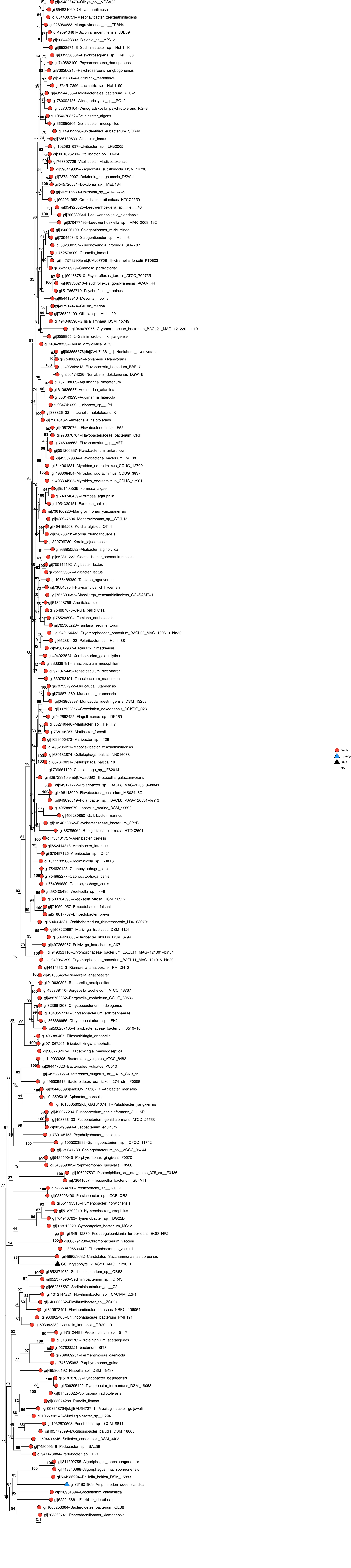
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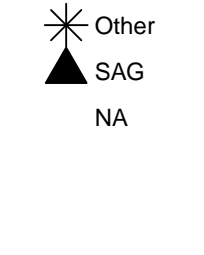
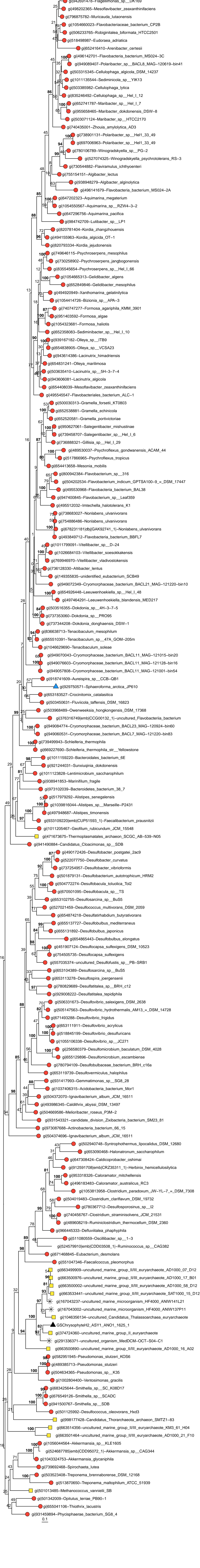


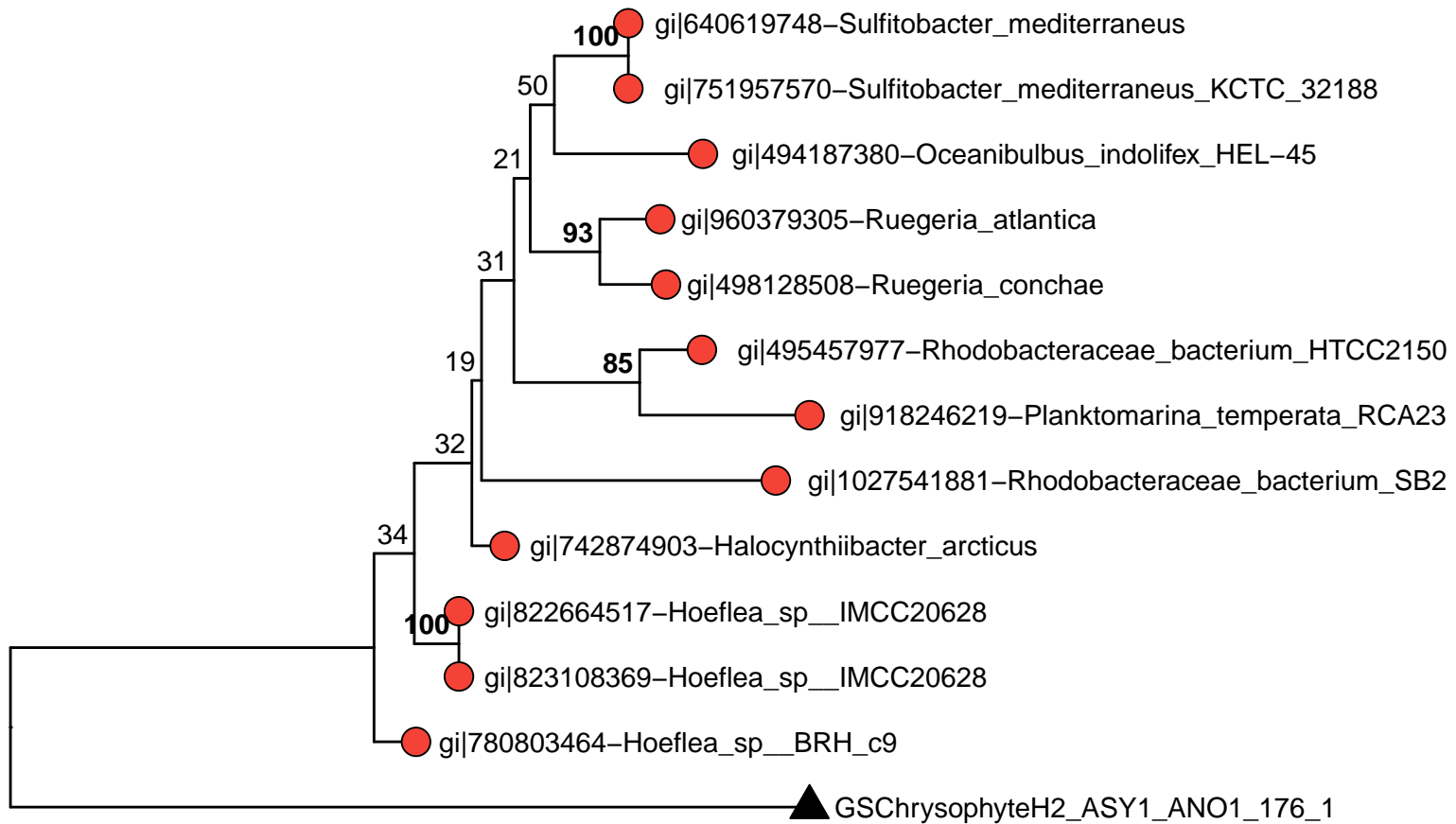




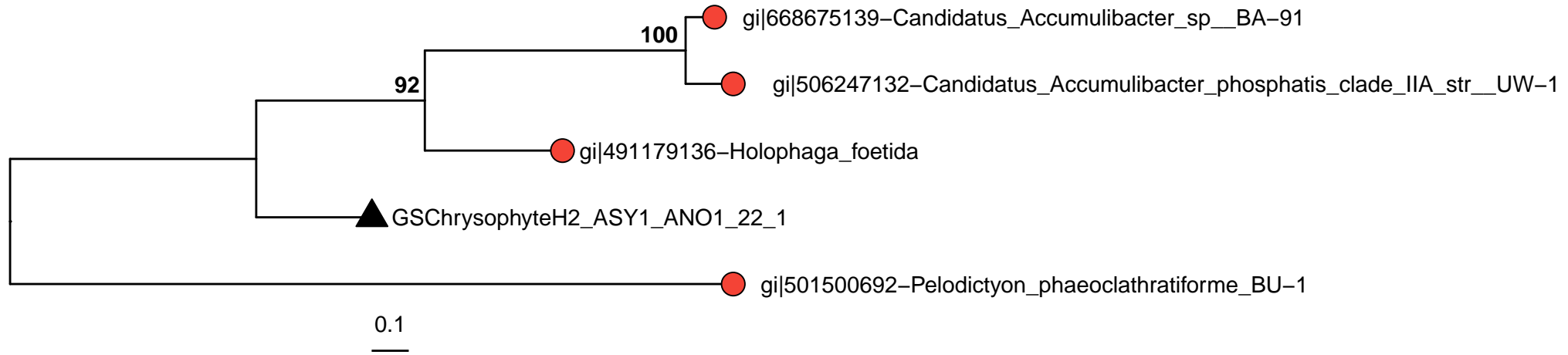




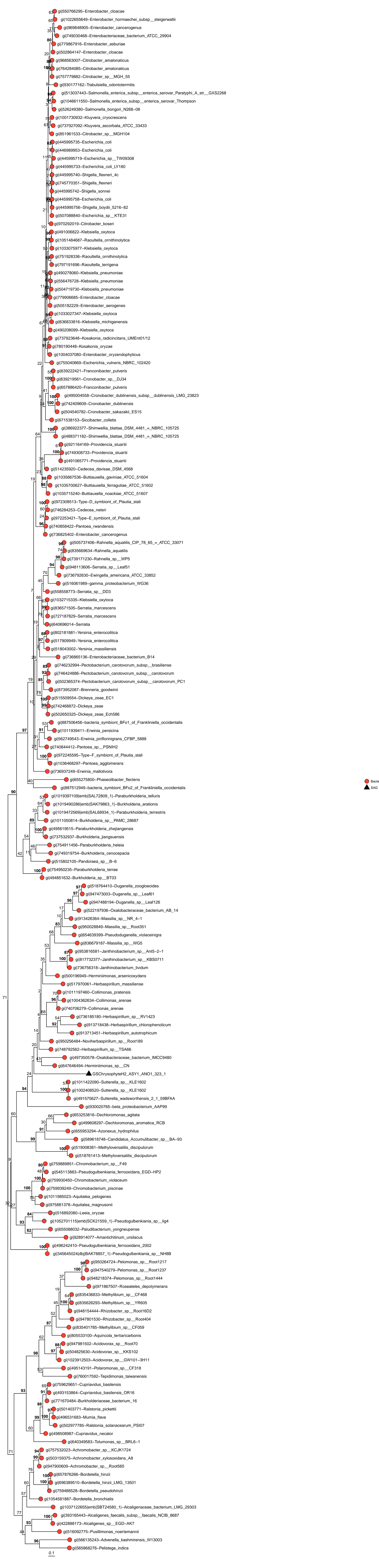




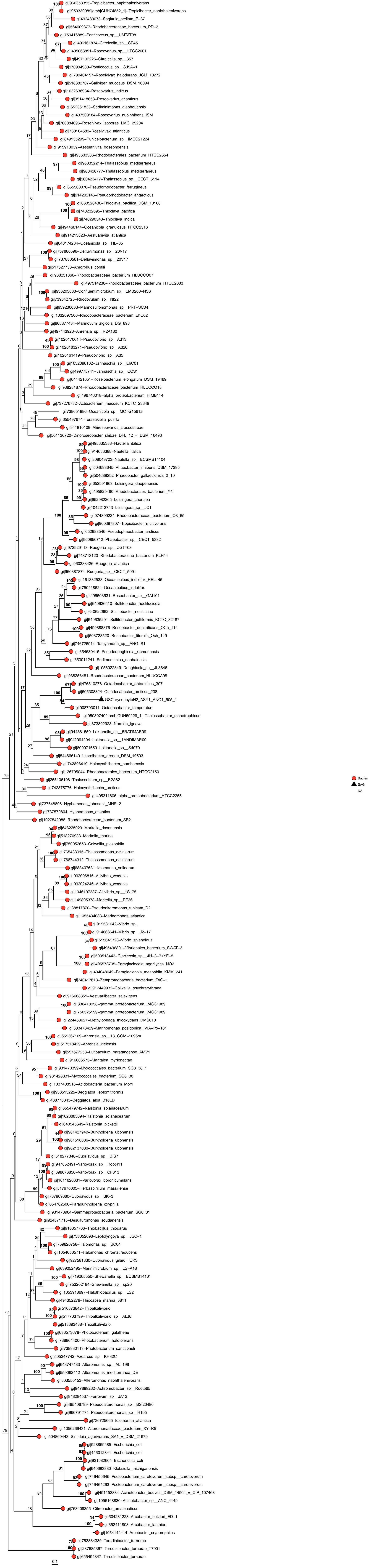
● Bacteria
▲ SAG



● Bacteria
▲ SAG

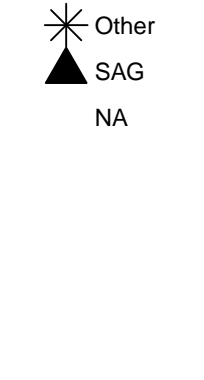
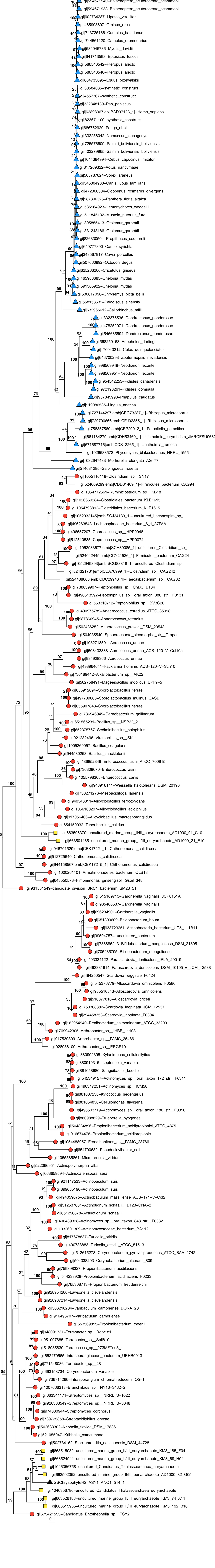


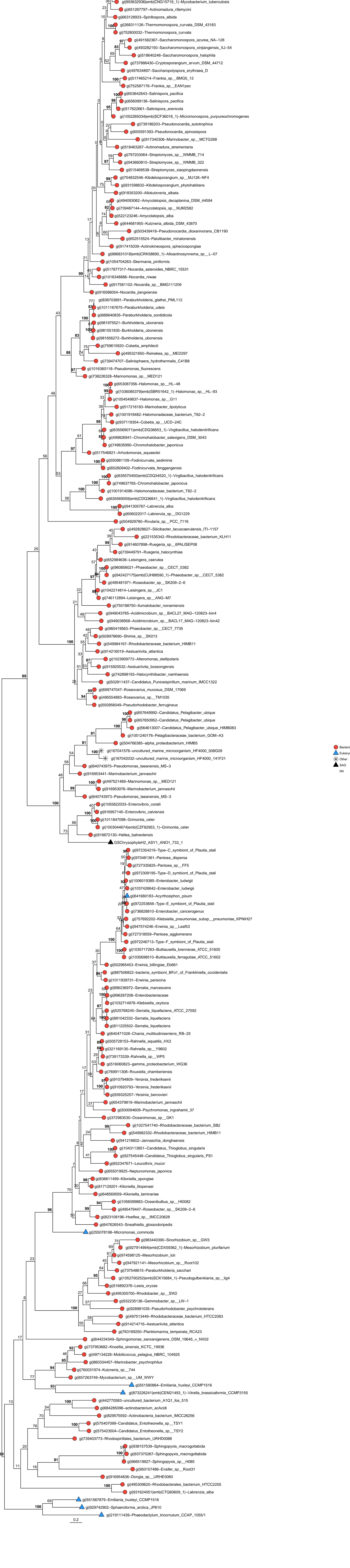
● Bacteria
▲ SAG



● Bacteria
▲ SAG
NA

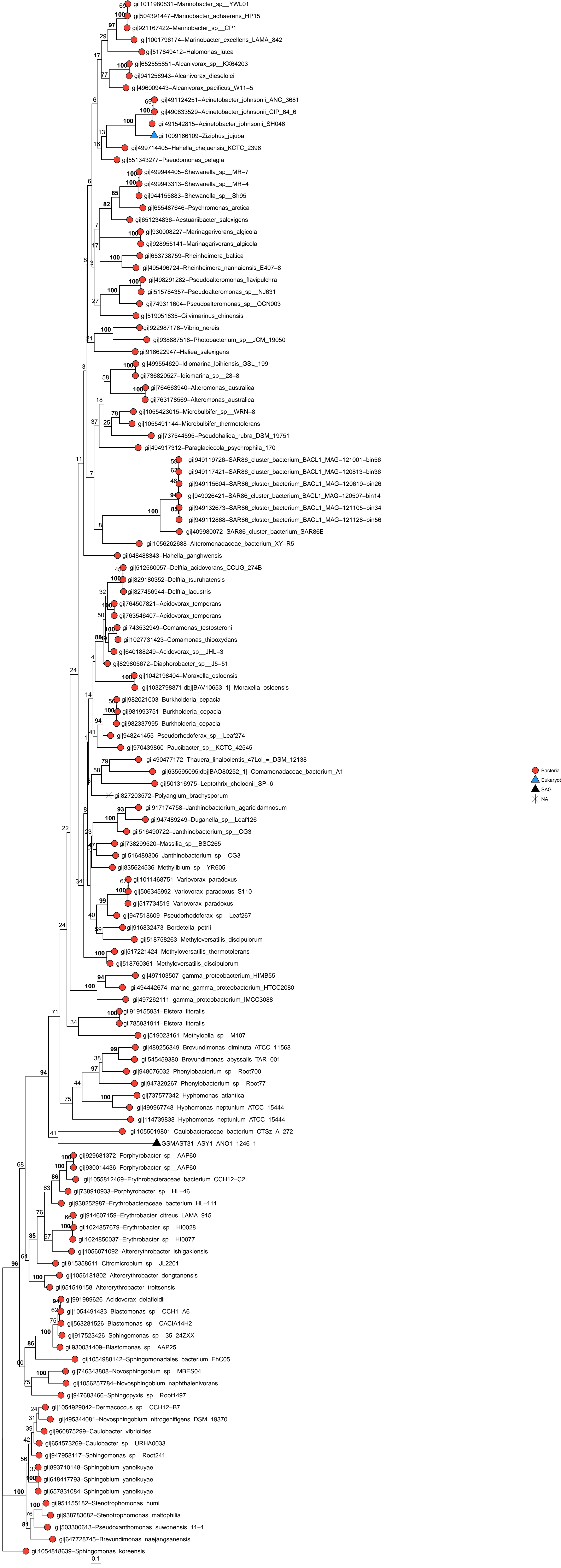
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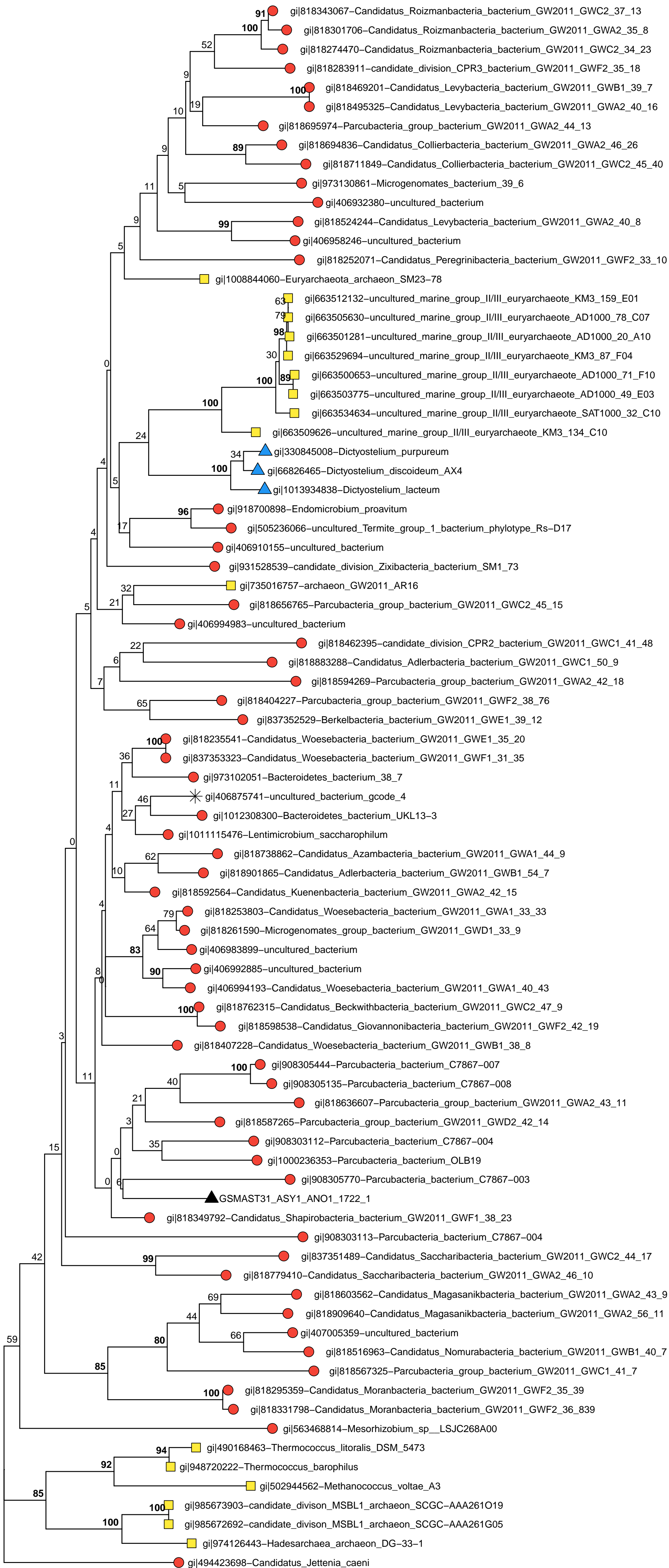
- Bacteria
- ▲ Eukaryota
- ▲ Other
- ★ SAG
- NA

0.2

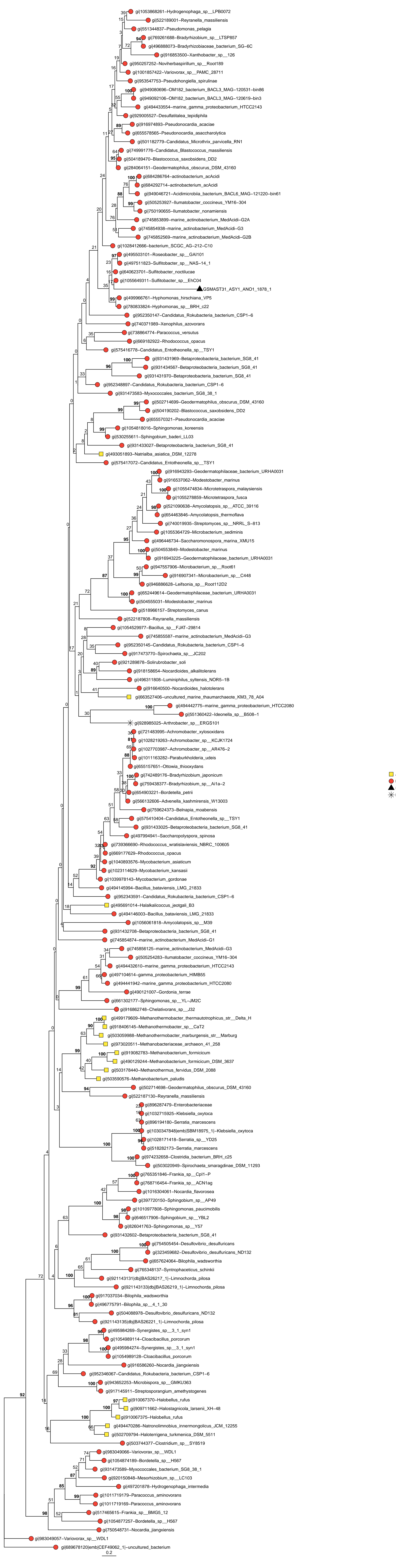


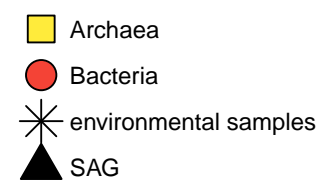
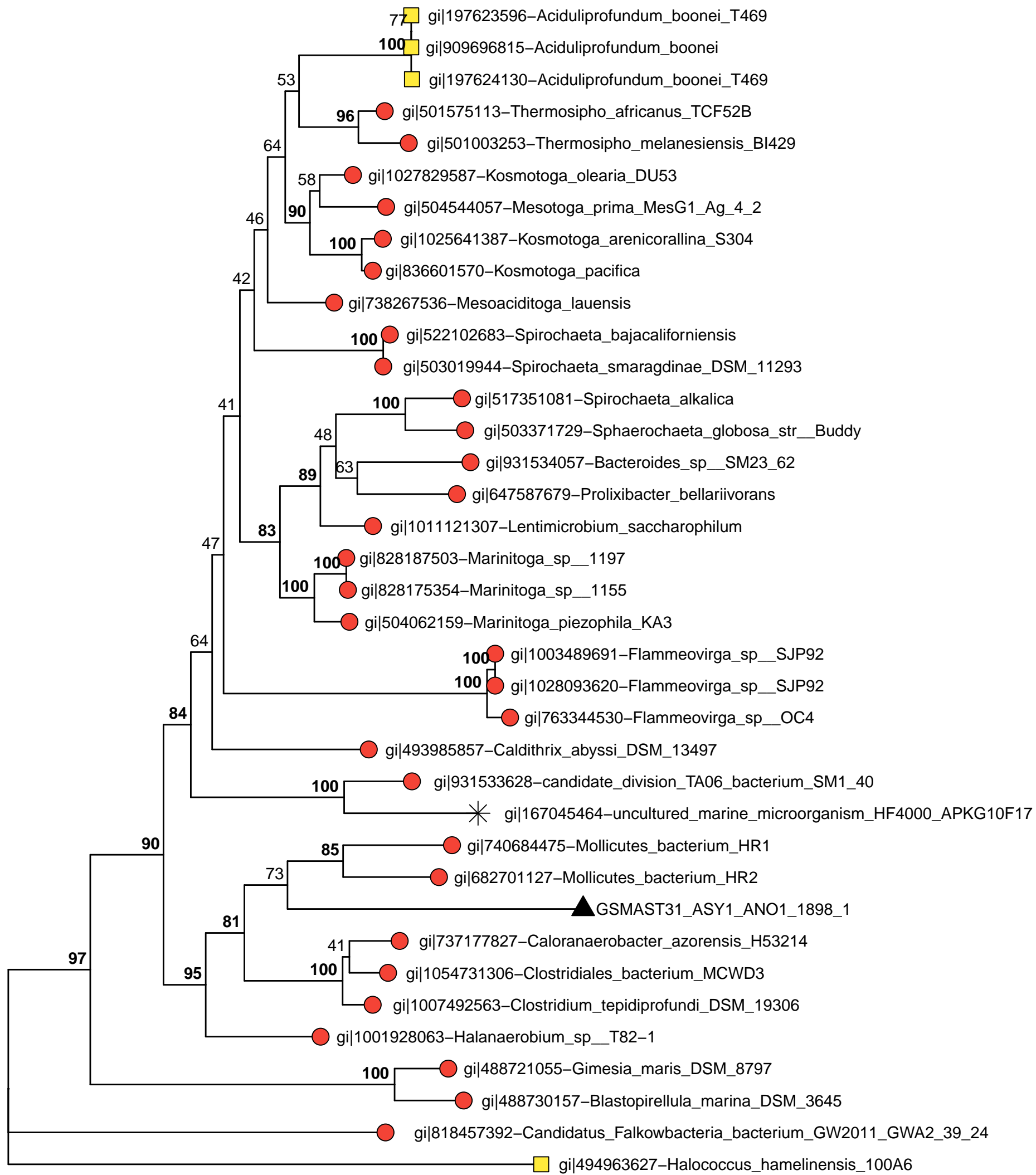
- Bacteria
- ▲ Eukaryota
- ▲ SAG
- * NA

0.1

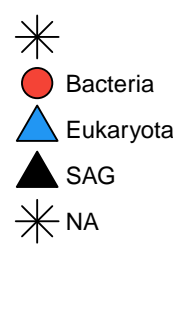
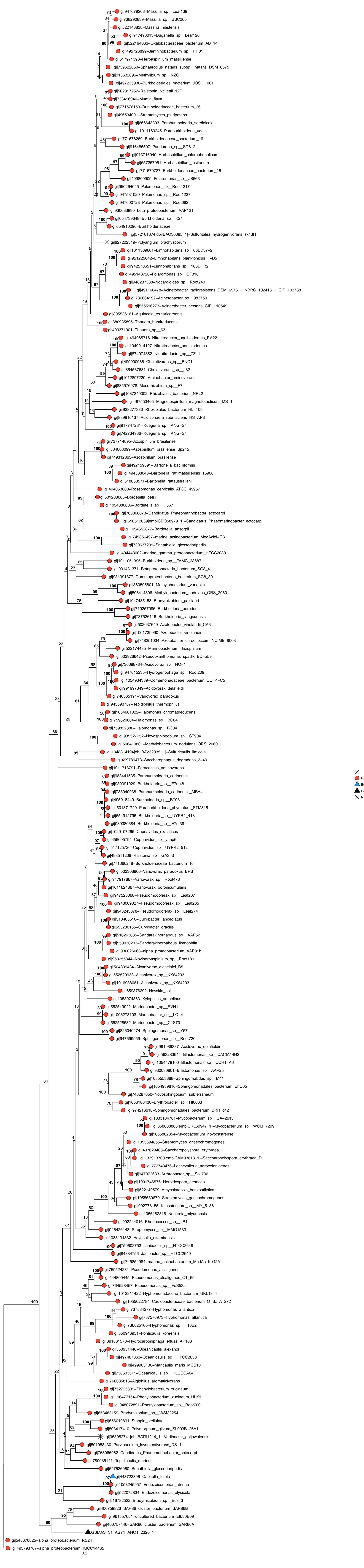


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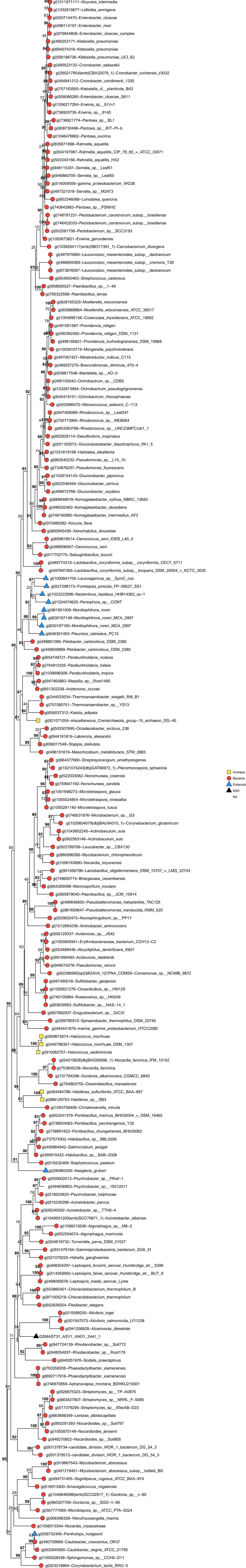


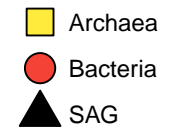
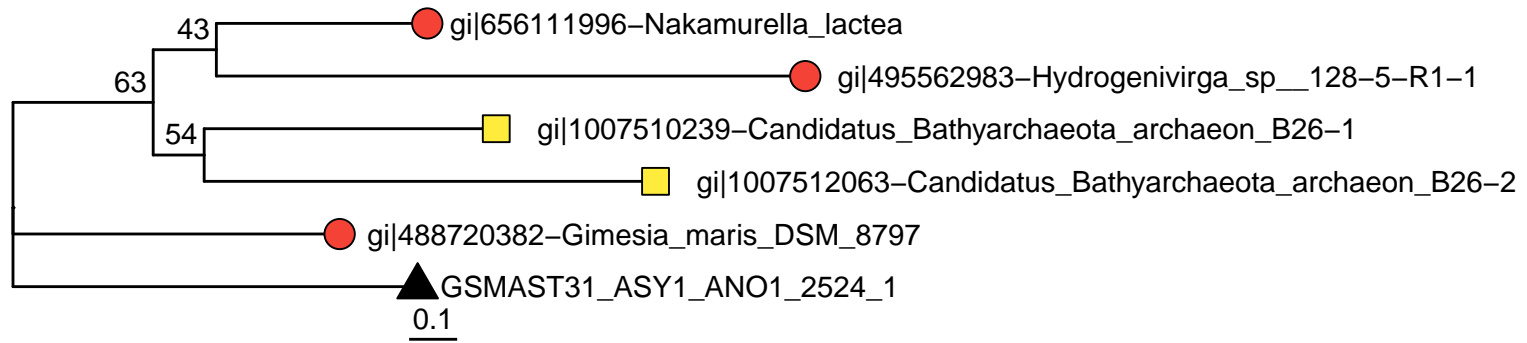


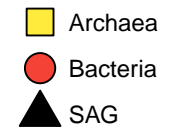
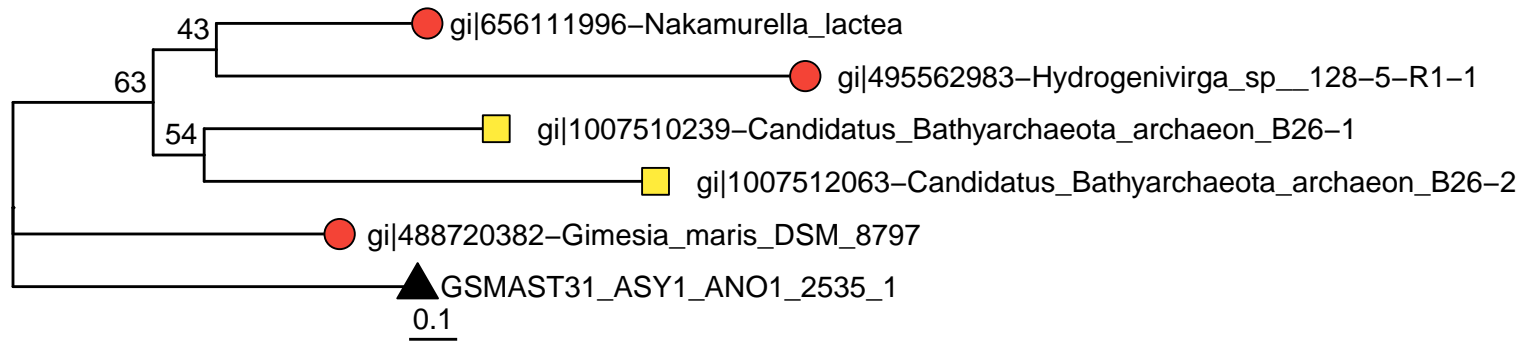
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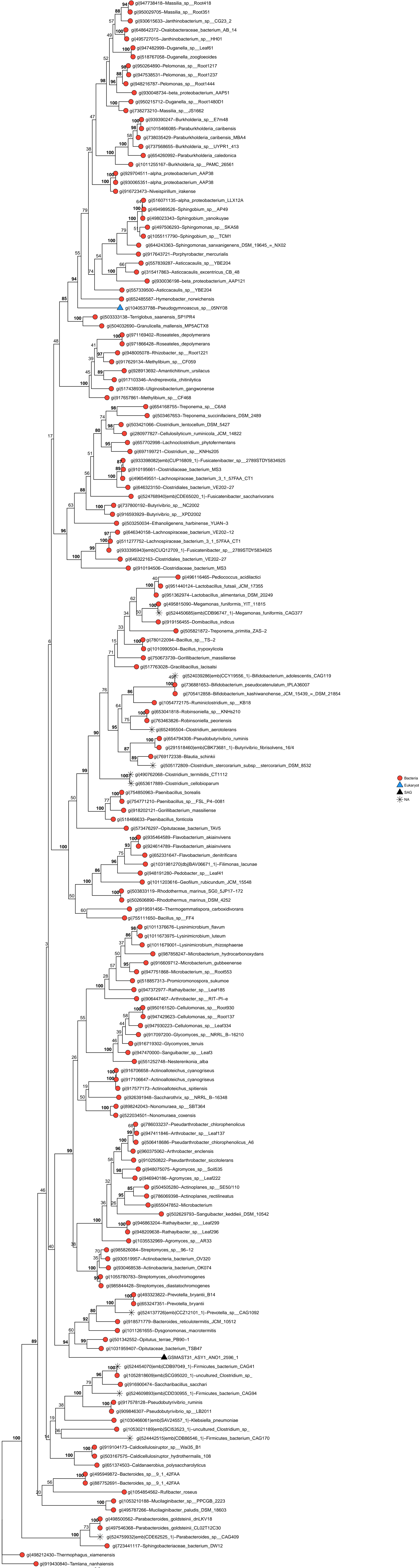


● gi545670825-alpha_proteobacterium_RS24
 ● gi495793767-alpha_proteobacterium_IMCC14465
 0.2









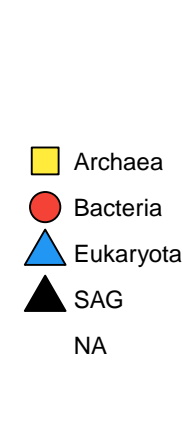
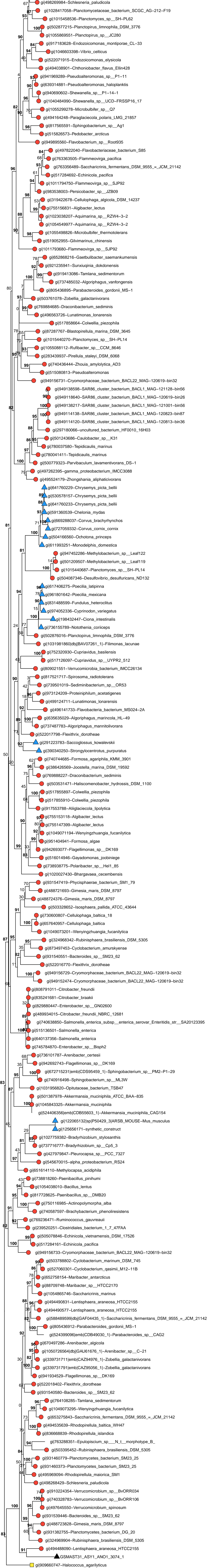
- Bacteria
- ▲ Eukaryota
- ▲ SAG
- * NA

gij19430840-Tamlana_xianhaiensis
 0.2

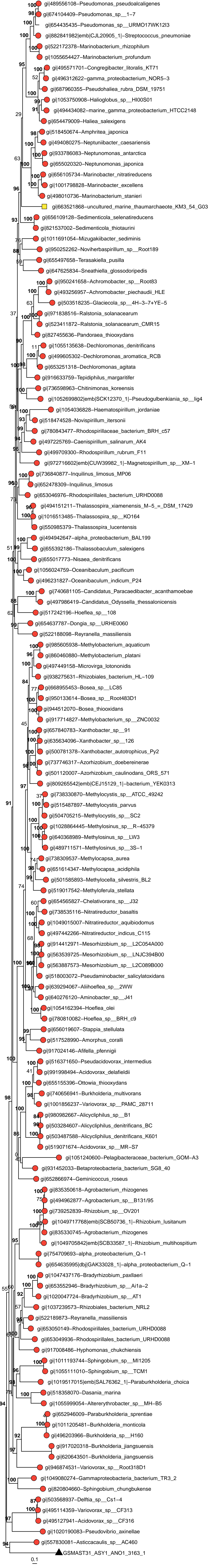


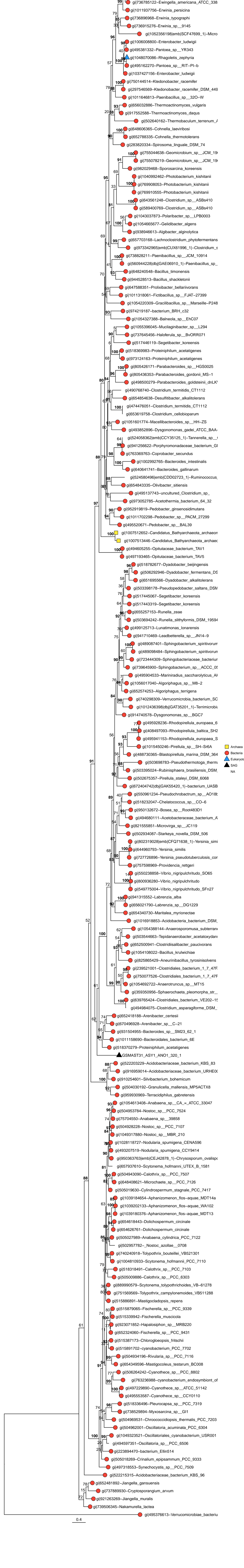
■ Archaea
● Bacteria
▲ SAG
★ NA

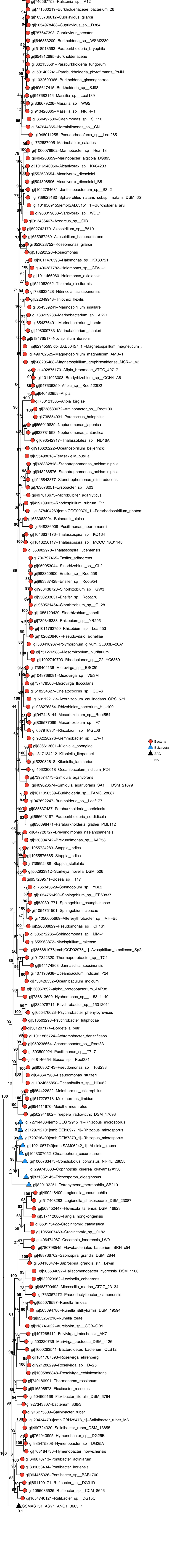
gij850511076-Methanosarcina_mazei
0.3

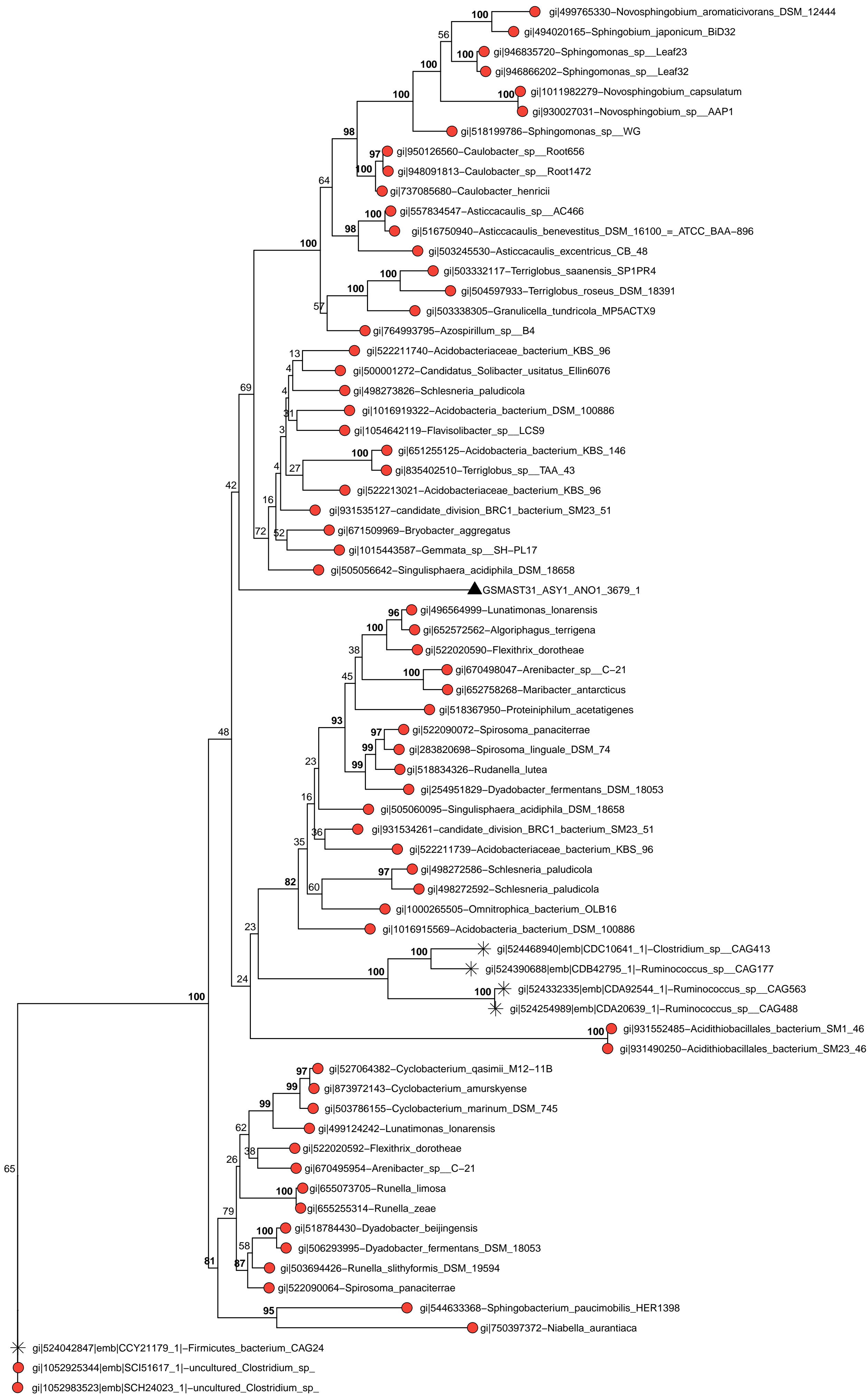


18 ● gij521961104-Zavarzinella_formosa
 67 ● gij498269984-Schlesneria_paludicola
 87 ● gij1028417058-Planctomycetaceae_bacterium_SCGC_AG-212-F19
 93 ● gij1015458536-Planctomyces_sp_SH-PL62
 100 ● gij502877215-Planctopirus_limnophila_DSM_3776
 90 ● gij1055869551-Planctopirus_sp_JC280
 84 ● gij917183628-Endozoicomonas_montiporae_CL-33
 94 ● gij1046603398-Vibrio_celticus
 49 ● gij522071915-Endozoicomonas_elysicola
 ● gij494038901-Chthoniobacter_flavus_Ellin428
 4 ● gij941969289-Pseudoalteromonas_sp_P1-11
 90 ● gij639314881-Pseudoalteromonas_haloplanktis
 ● gij940690602-Shewanella_sp_P1-14-1
 96 ● gij1040484990-Shewanella_sp_UCD-FRSSP16_17
 82 ● gij1055299278-Microbulbifer_sp_Q7
 90 ● gij494164248-Paraglaciecola_polaris_LMG_21857
 98 ● gij817565591-Sphingobacterium_sp_Ag1
 ● gij515826573-Pedobacter_arcticus
 ● gij949895560-Flavobacterium_sp_Root935
 84 ● gij497822040-Flavobacteriaceae_bacterium_S85
 68 ● gij763363505-Flammeovirga_pacifica
 93 ● gij763356489-Saccharicrinis_fermentans_DSM_9555_JCM_21142
 96 ● gij517284692-Echinicola_pacifica
 61 ● gij1011794750-Flammeovirga_sp_SJP92
 94 ● gij983538003-Persicobacter_sp_JZB09
 98 ● gij319422678-Cellulophaga_algicola_DSM_14237
 99 ● gij755156831-Algibacter_lectus
 88 ● gij1023038207-Aquimarina_sp_RZW4-3-2
 ● gij1054549977-Aquimarina_sp_RZW4-3-2
 98 ● gij1055498826-Microbulbifer_thermotolerans
 ● gij519052955-Gilvimirinus_chinensis
 ● gij1011793680-Flammeovirga_sp_SJP92
 78 ● gij652868216-Gaetbulbacter_saemankumensis
 92 ● gij921235941-Sunxiuqinia_dokdonensis
 100 ● gij919413086-Tamlana_sedimentorum
 ● gij737485032-Algoriphagus_vanfongensis
 ● gij805436895-Parabacteroides_gordonii_MS-1
 93 ● gij503761078-Zobellia_galactanivorans
 99 ● gij769884685-Draconibacterium_sediminis
 ● gij496563726-Lunatimonas_lonarensis
 ● gij517858664-Colwellia_piezophila
 78 ● gij87287767-Blastopirellula_marina_DSM_3645
 70 ● gij1015440270-Planctomyces_sp_SH-PL14
 93 ● gij1055088112-Rufibacter_sp_CCM_8646
 97 ● gij283439937-Pirellula_staleyii_DSM_6068
 99 ● gij740436444-Zhouia_amylyolytica_AD3
 ● gij515080813-Pseudoalteromonas
 ● gij949156731-Cryomorpaceae_bacterium_BACL22_MAG-120619-bin32
 ● gij949138598-SAR86_cluster_bacterium_BACL1_MAG-121128-bin56
 90 ● gij949118640-SAR86_cluster_bacterium_BACL1_MAG-120619-bin26
 100 ● gij949138217-SAR86_cluster_bacterium_BACL1_MAG-121001-bin56
 95 ● gij949114138-SAR86_cluster_bacterium_BACL1_MAG-120823-bin87
 ● gij949114120-SAR86_cluster_bacterium_BACL1_MAG-120813-bin36
 88 ● gij297180066-uncultured_bacterium_HF0010_16H03
 86 ● gij501243686-Caulobacter_sp_K31
 95 ● gij780037580-Tepidicaulis_marinus
 ● gij780041411-Tepidicaulis_marinus
 100 ● gij500779323-Parvibaculum_lavamentivorans_DS-1
 79 ● gij497262395-gamma_proteobacterium_IMCC3088
 ● gij495524179-Zhongshania_aliphaticivorans
 ● gij641760229-Chrysemys_picta_bellii
 89 ● gij530578157-Chrysemys_picta_bellii
 88 ● gij641760233-Chrysemys_picta_bellii
 93 ● gij591360539-Chelonina_mydas
 100 ● gij669288037-Corvus_brachyrhynchos
 ● gij727059332-Corvus_cornix_cornix
 ● gij504166560-Ochotona_princeps
 ● gij611993251-Monodelphis_domestica
 81 ● gij947452286-Methylobacterium_sp_Leaf122
 14 ● gij501209507-Methylobacterium_sp_Leaf119
 100 ● gij1015440687-Planctomyces_sp_SH-PL14
 ● gij504087346-Desulfovibrio_desulfuricans_ND132
 96 ● gij617406275-Poecilia_latipinna
 78 ● gij961801642-Poecilia_mexicana
 90 ● gij831488599-Fundulus_heteroclitus
 15 ● gij974052336-Cyprinodon_variegatus
 100 ● gij198432447-Ciona_intestinalis
 24 ● gij736155789-Notothenia_coriiceps
 77 ● gij502876016-Planctopirus_limnophila_DSM_3776
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 100 ● gij752320930-Cupriavidus_basilensis
 4 ● gij517126097-Cupriavidus_sp_UYPR2_512
 ● gij809021551-Verrucomicrobia_bacterium_IMCC26134
 90 ● gij817521717-Spirosoma_radiotolerans
 42 ● gij739501019-Sediminibacterium_sp_OR53
 18 ● gij973124209-Proteiniphilum_acetatifgenes
 96 ● gij499124711-Lunatimonas_lonarensis
 92 ● gij496141733-Flavobacteria_bacterium_MS024-2A
 89 ● gij635635029-Algoriphagus_marincola_HL-49
 73 ● gij737487783-Algoriphagus_mannitolivorans
 ● gij522017798-Flexithrix_dorotheae
 92 ● gij291223783-Saccoglossus_kowalevskii
 ● gij390340250-Strongylocentrotus_purpuratus
 62 ● gij740744685-Formosa_agariphila_KMM_3901
 96 ● gij386426569-Joostella_marina_DSM_19592
 75 ● gij769882227-Draconibacterium_sediminis
 45 ● gij503531471-Haliscomenobacter_hydrossis_DSM_1100
 100 ● gij517855897-Colwellia_piezophila
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 97 ● gij755153118-Algibacter_lectus
 80 ● gij755147399-Algibacter_lectus
 92 ● gij1049071194-Wenyngzhuangia_fucanilytica
 99 ● gij951404941-Formosa_algae
 33 ● gij942693077-Flagellimonas_sp_DK169
 88 ● gij516014946-Gayadomonas_joobiniege
 ● gij738938775-Polaribacter_sp_Hel1_85
 ● gij1020027430-Bhargavaea_cecembensis
 70 ● gij931547419-Phycisphaerae_bacterium_SM1_79
 10 ● gij488721693-Gimesia_maris_DSM_8797
 30 ● gij488724376-Gimesia_maris_DSM_8797
 91 ● gij503328652-Isosphaera_pallida_ATCC_43644
 100 ● gij730600807-Cellulophaga_baltica_18
 ● gij657640957-Cellulophaga_baltica
 ● gij1049073201-Wenyngzhuangia_fucanilytica
 95 ● gij324968342-Rubinisphaera_brasiliensis_DSM_5305
 35 ● gij873497453-Cyclobacterium_amurskyense
 82 ● gij931540551-Bacteroides_sp_SM23_62
 1 ● gij522016770-Flexithrix_dorotheae
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 ● gij949152474-Cryomorpaceae_bacterium_BACL22_MAG-120619-bin32
 ● gij808791011-Citrobacter_freundii
 ● gij835241681-Citrobacter_braakii
 32 ● gij829880447-Enterobacter_sp_GN02600
 89 ● gij489934015-Citrobacter_freundii_NBRC_12681
 ● gij740638850-Salmonella_enterica_subsp_enterica_serovar_Enteritidis_str_SA20123395
 100 ● gij515136501-Salmonella_enterica
 ● gij640137356-Salmonella_enterica
 ● gij745784870-Enterobacter_sp_Bisph2
 68 ● gij736101787-Arenibacter_certesii
 35 ● gij942692743-Flagellimonas_sp_DK169
 95 ● gij672715231[emb]CDS95459_1-Sphingobacterium_sp_PM2-P1-29
 ● gij740916498-Sphingobacterium_sp_ML3W
 ● gij1031956820-Opitutaceae_bacterium_TSB47
 ● gij501387978-Akkermansia_muciniphila_ATCC_BAA-835
 100 ● gij1045843325-Akkermansia_muciniphila
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 ● gij122065132[sp]P50429_3[ARSB_MOUSE]-Mus_musculus
 ● gij125656171-synthetic_construct
 81 ● gij1027759382-Bradyrhizobium_stylosanthis
 43 ● gij737716777-Bradyrhizobium_sp_Cp5_3
 97 ● gij427979847-Pleurocapsa_sp_PCC_7327
 ● gij545670015-alpha_proteobacterium_RS24
 ● gij651614110-Methylocapsa_acidiphila
 43 ● gij738818260-Paenibacillus_pinihumi
 94 ● gij1054038010-Bacillus_lentus
 95 ● gij817728625-Paenibacillus_sp_DMB20
 100 ● gij750116985-Actinopolymorpha_alba
 ● gij740587597-Brachyobacterium_phenoliresistens
 92 ● gij769236471-Ruminococcus_gauvreauii
 ● gij239520251-Clostridiales_bacterium_1_7_7FAA
 99 ● gij505078846-Echinicola_vietnamensis_DSM_17526
 ● gij517284161-Echinicola_pacifica
 ● gij949156733-Cryomorpaceae_bacterium_BACL22_MAG-120619-bin32
 95 ● gij503788802-Cyclobacterium_marinum_DSM_745
 86 ● gij527060301-Cyclobacterium_qasimii_M12-11B
 78 ● gij652758154-Maribacter_antarcticus
 97 ● gij88709748-Maribacter_sp_HTCC2170
 ● gij1054865746-Saccharicrinis_marinus
 88 ● gij494490831-Lentisphaera_araneosa_HTCC2155
 38 ● gij494490577-Lentisphaera_araneosa_HTCC2155
 82 ● gij588489599[dbj]GAF04435_1-Saccharicrinis_fermentans_DSM_9555_JCM_21142
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 95 ● gij1050726564[dbj]GAU61676_1-Arenibacter_sp_C-21
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 67 ● gij522018402-Flexithrix_dorotheae
 ● gij931540580-Bacteroides_sp_SM23_62
 96 ● gij764108285-Tamlana_sedimentorum
 99 ● gij1049073295-Wenyngzhuangia_fucanilytica
 73 ● gij653275843-Saccharicrinis_fermentans_DSM_9555_JCM_21142
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 94 ● gij763288351-Epulopiscium_sp_N_t_morphotype_B_1
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 47 ● gij931460373-Planctomycetes_bacterium_SM23_25
 44 ● gij495969094-Rhodopirellula_maiorica_SM1
 ● gij498268429-Schlesneria_paludicola
 74 ● gij910224354-Verrucomicrobium_sp_BvORR034
 100 ● gij740328783-Verrucomicrobium_sp_BvORR106
 96 ● gij497645550-Verrucomicrobium_spinorum
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 ● GSMAST31_ASY1_ANO1_3074_1
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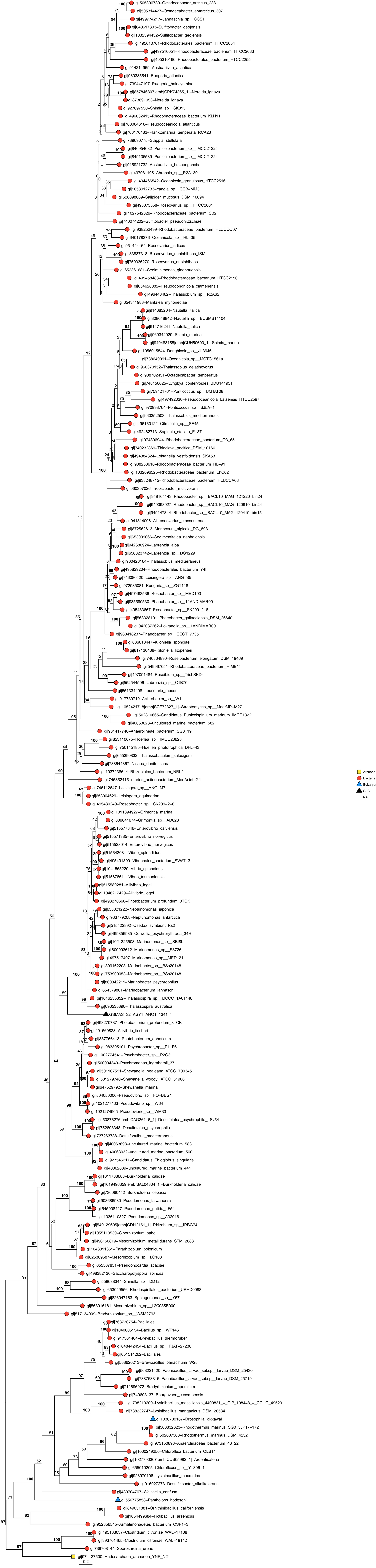
● Bacteria
▲ SAG
✱ NA

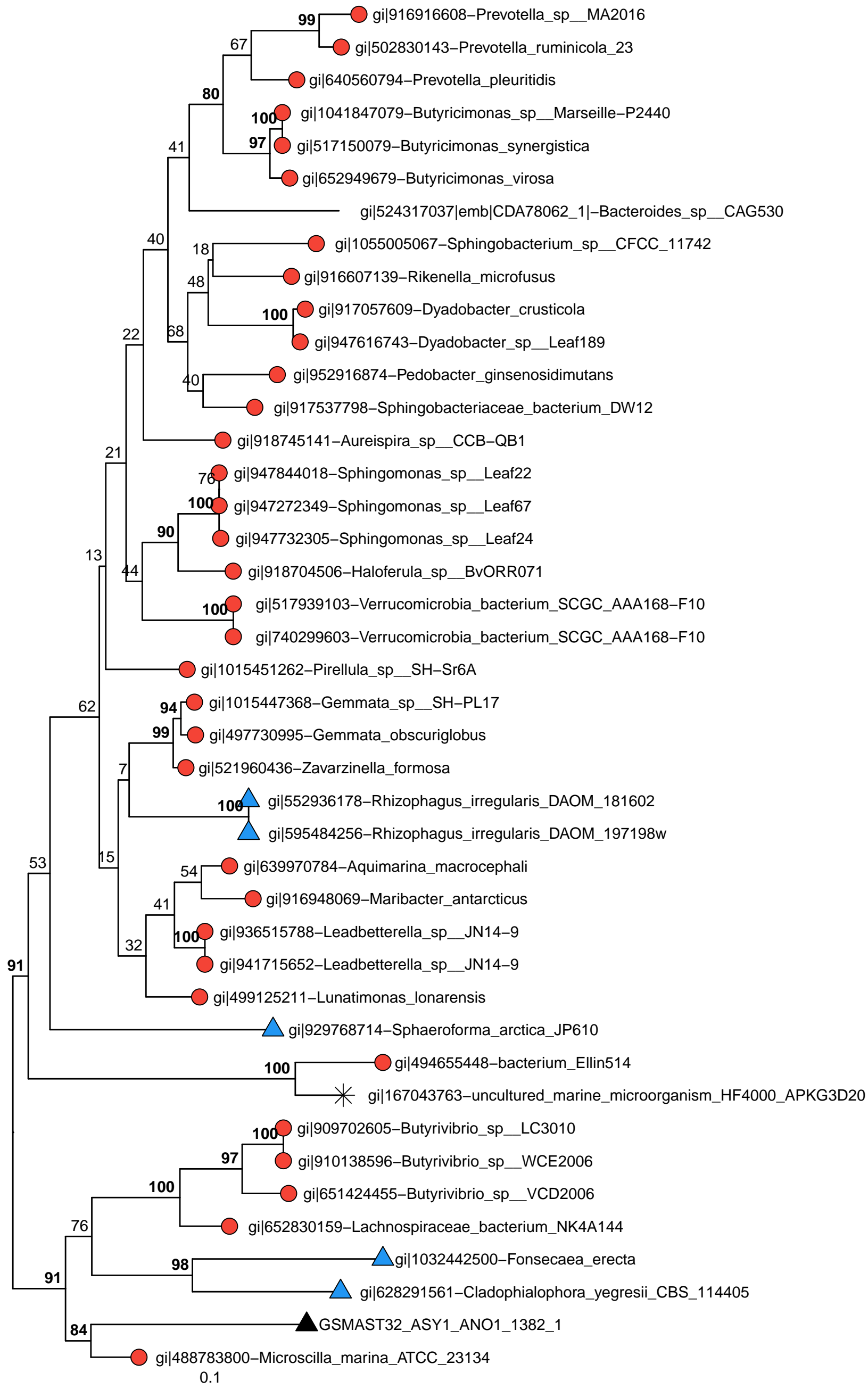
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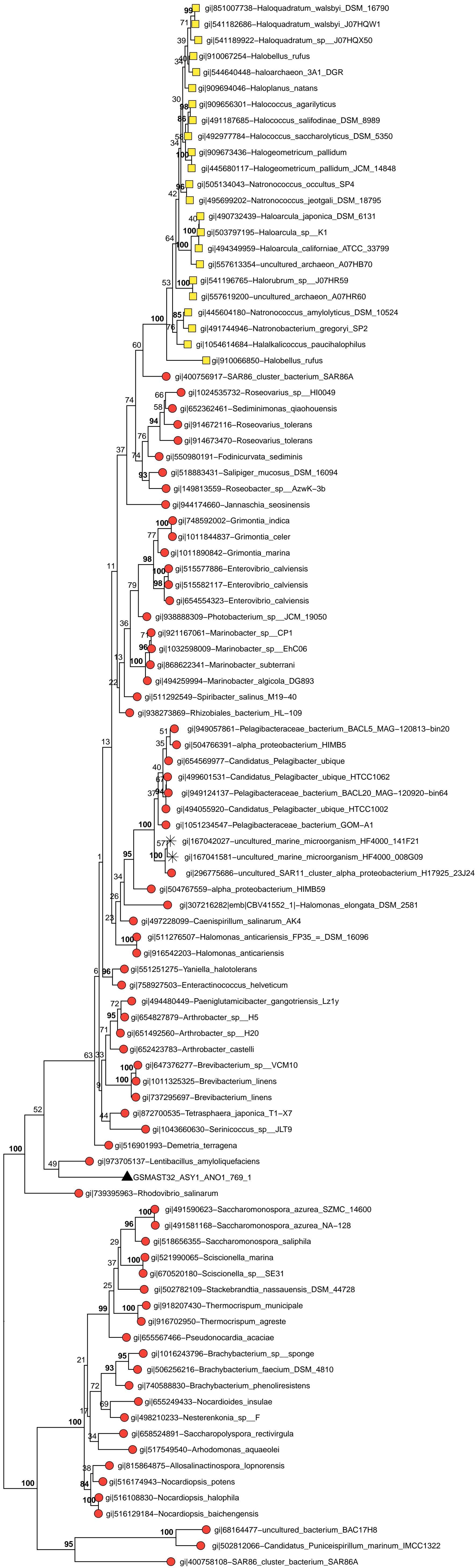


● Bacteria
▲ SAG
NA

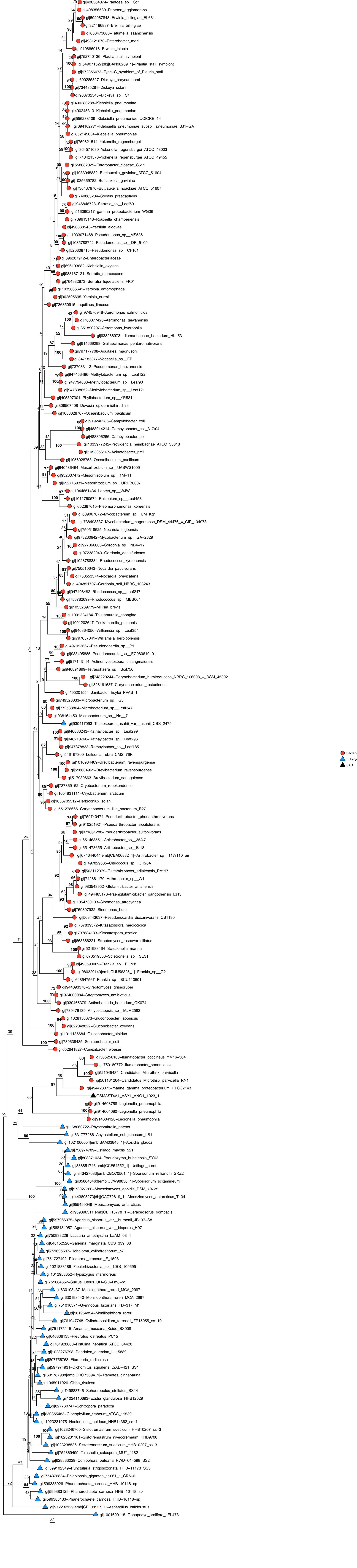
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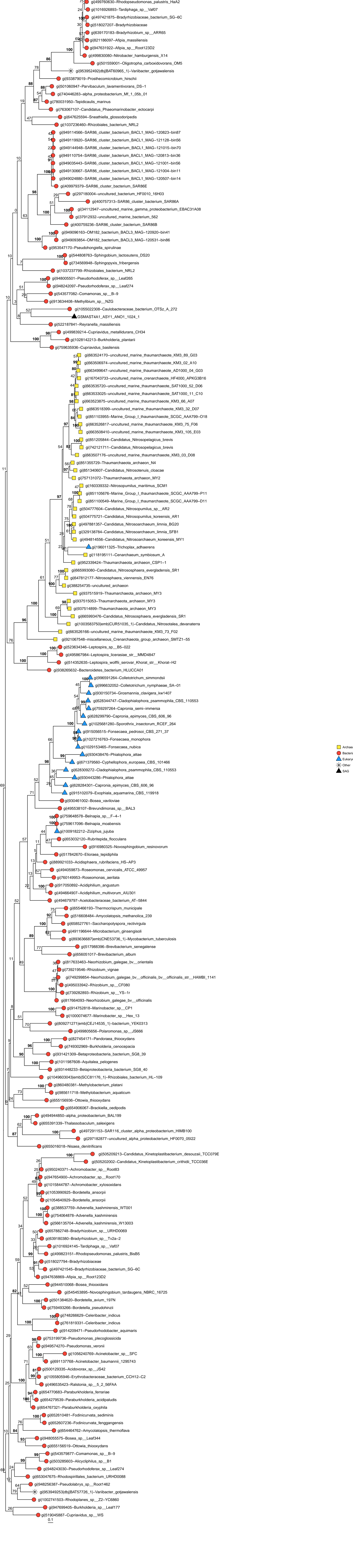


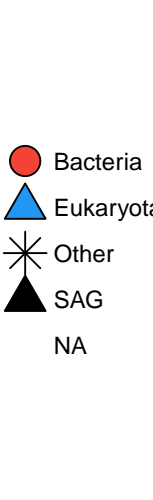
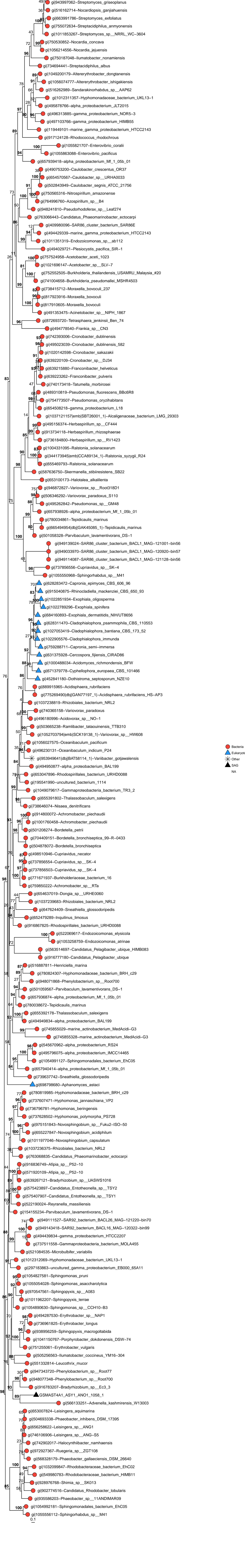


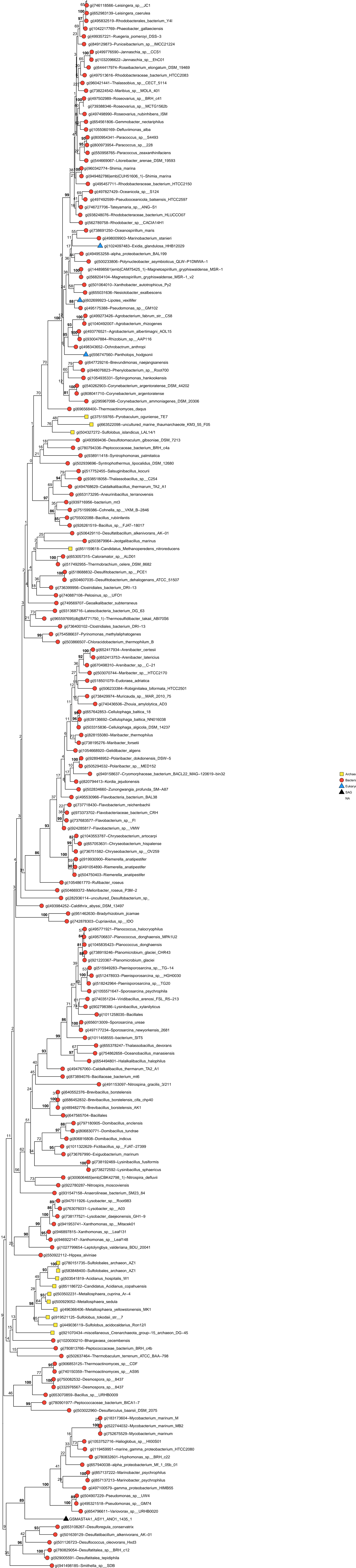


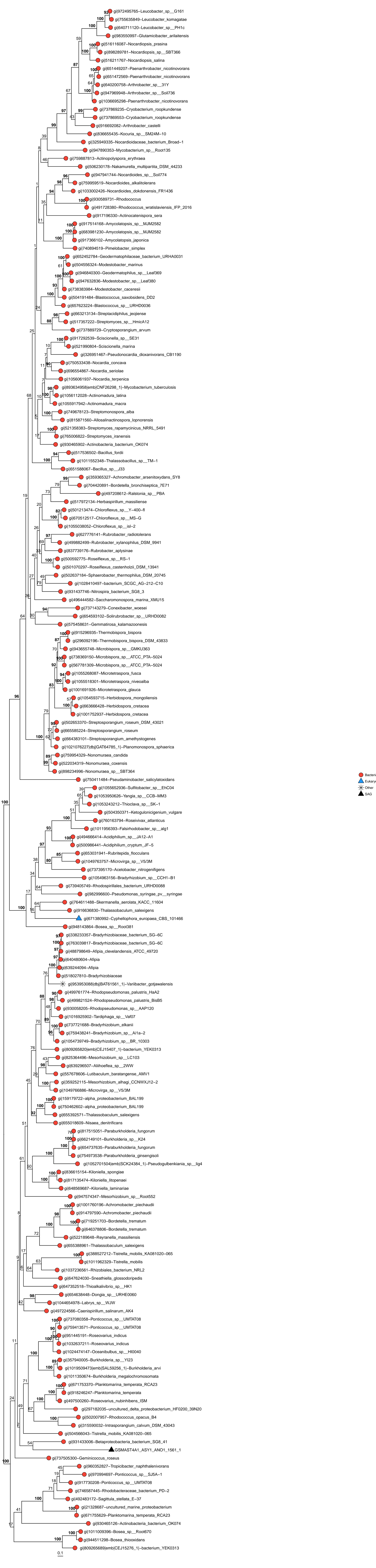
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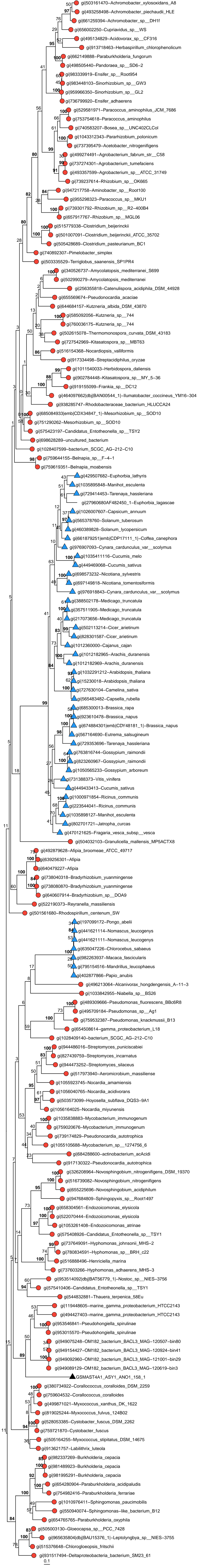


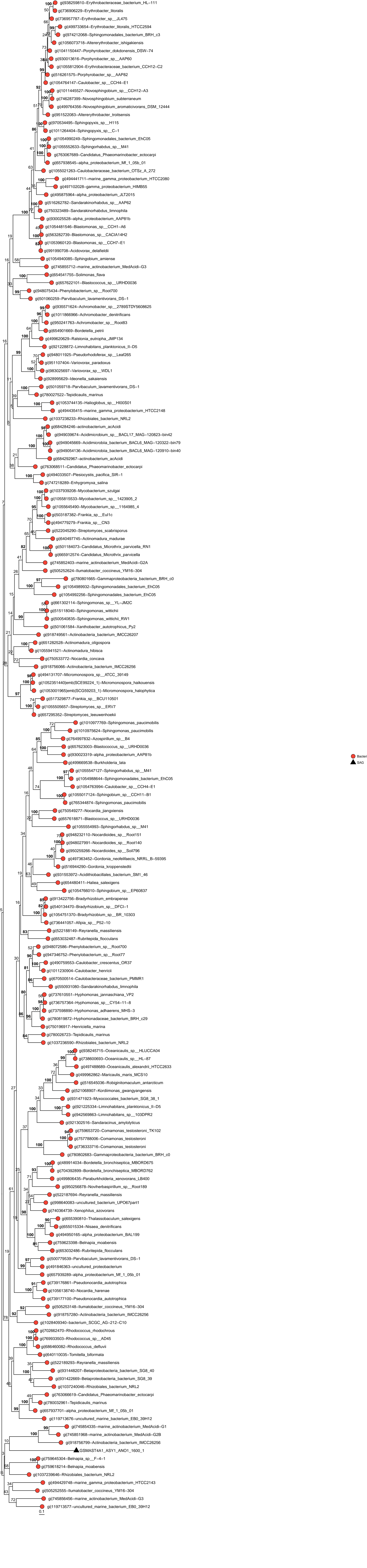


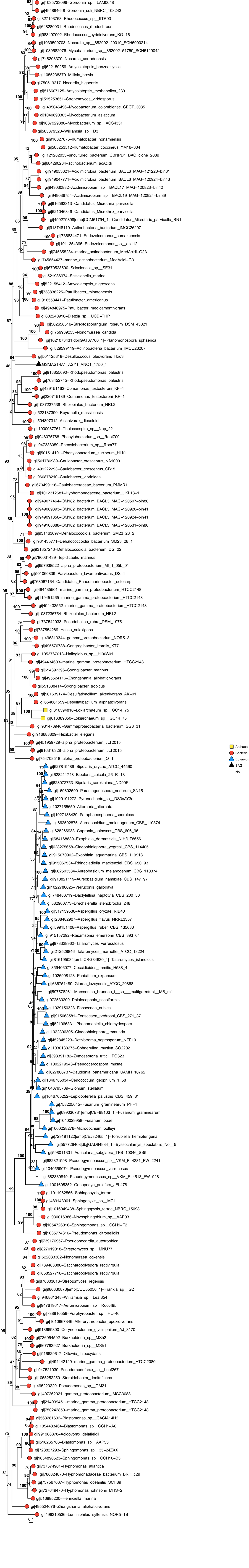


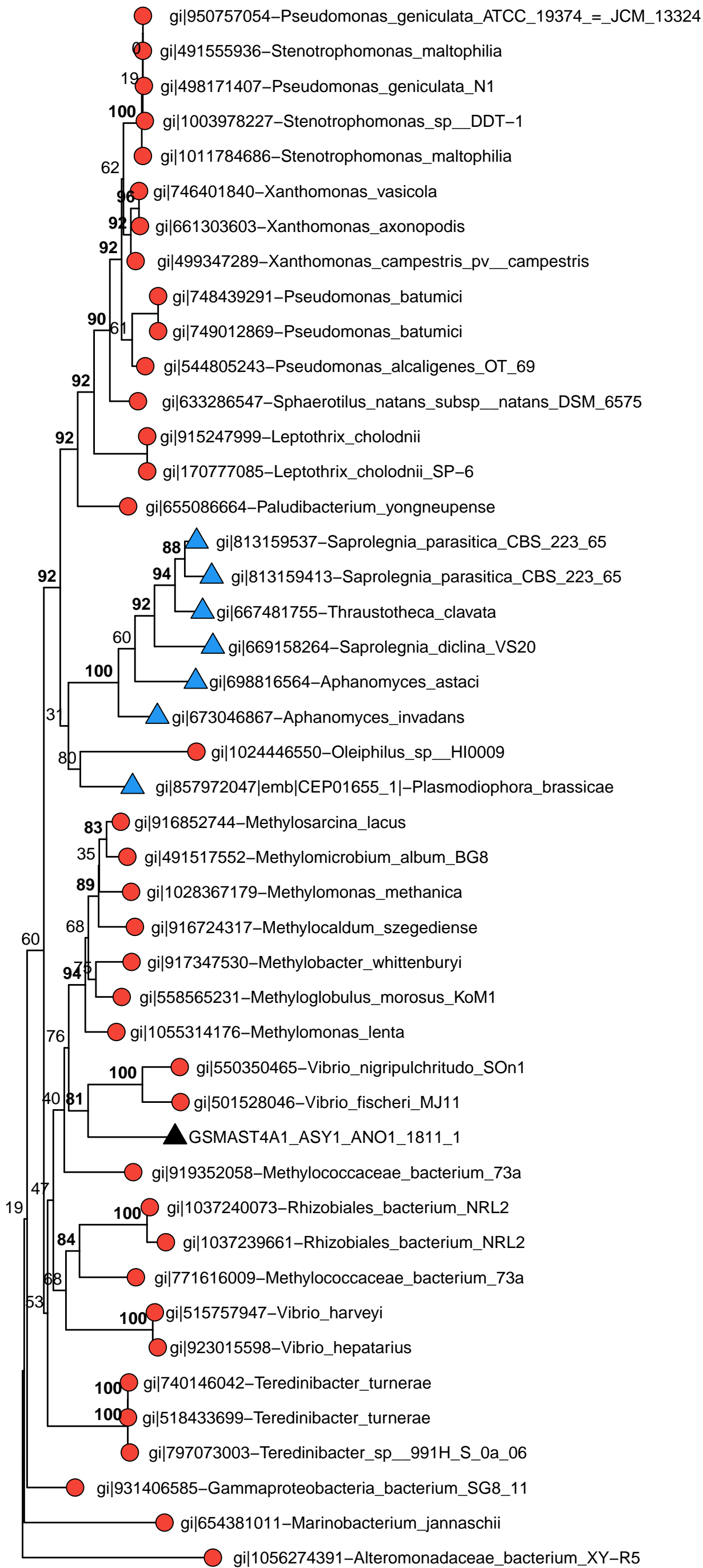
● Bacteria
▲ Eukaryota
★ Other
▲ SAG

0.1



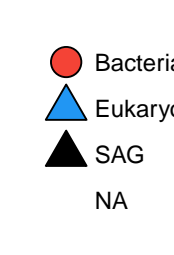
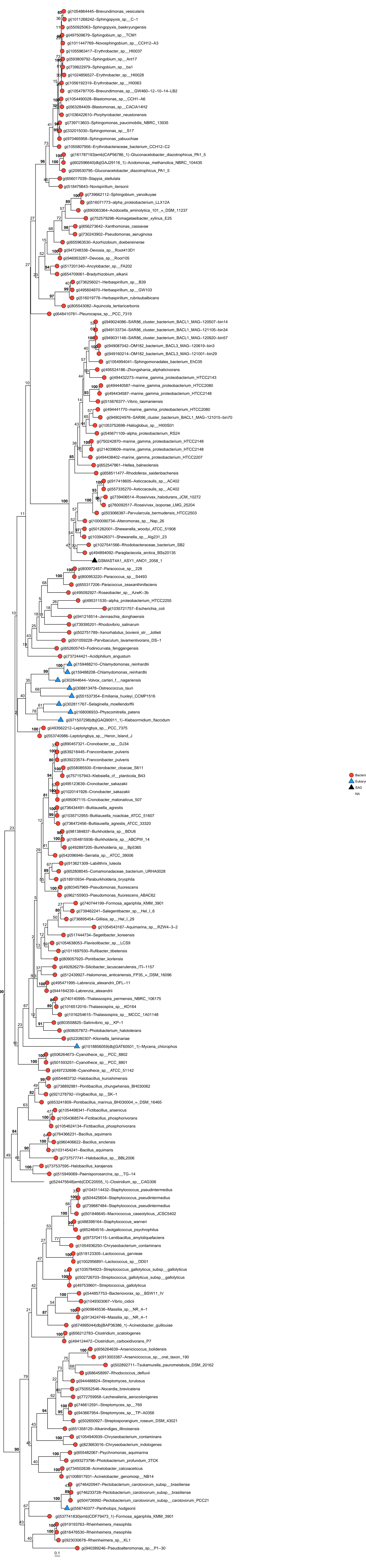


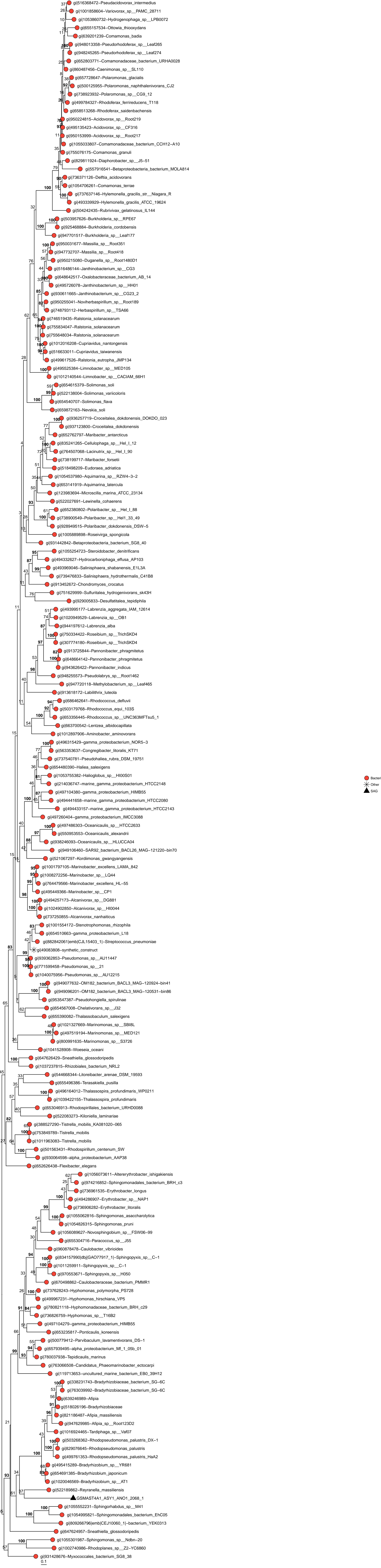




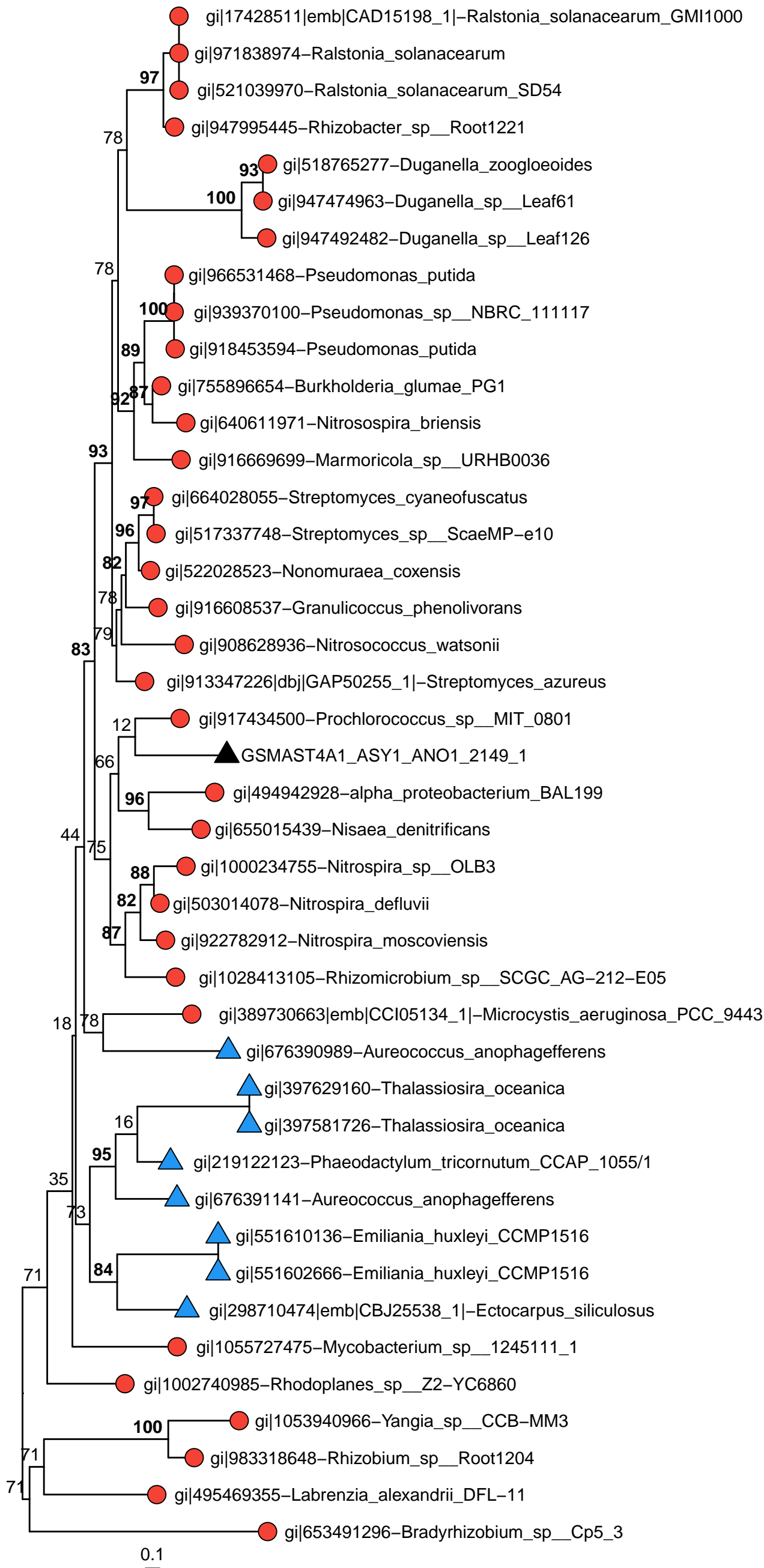
- Bacteria
- ▲ Eukaryota
- ▲ SAG

0.1

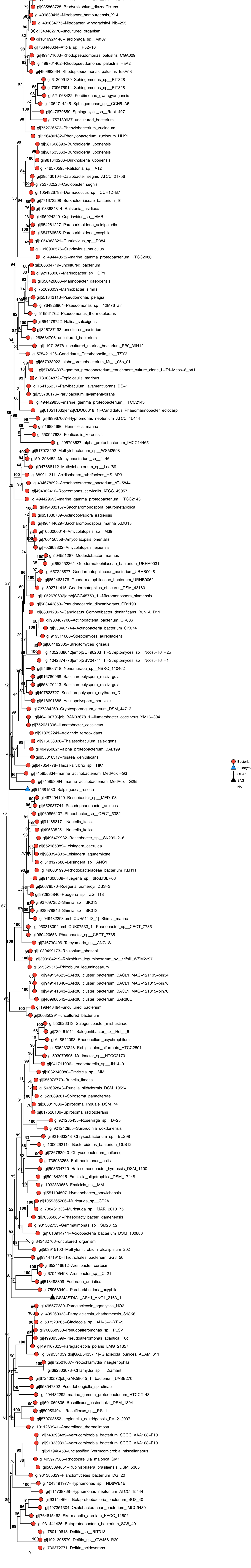


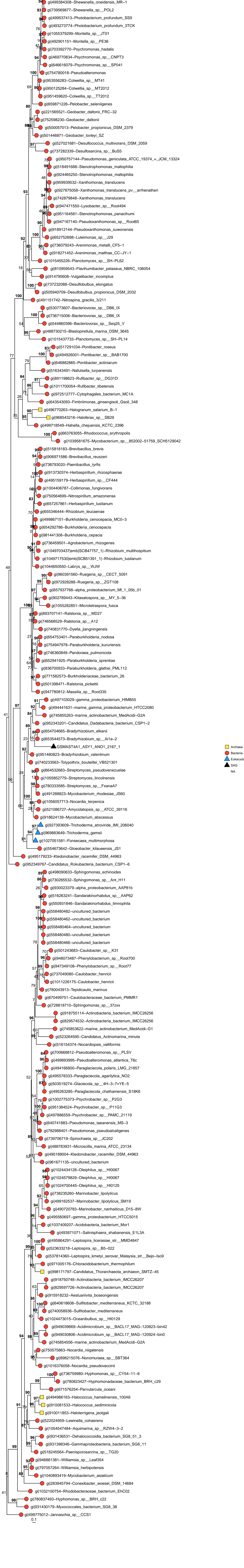


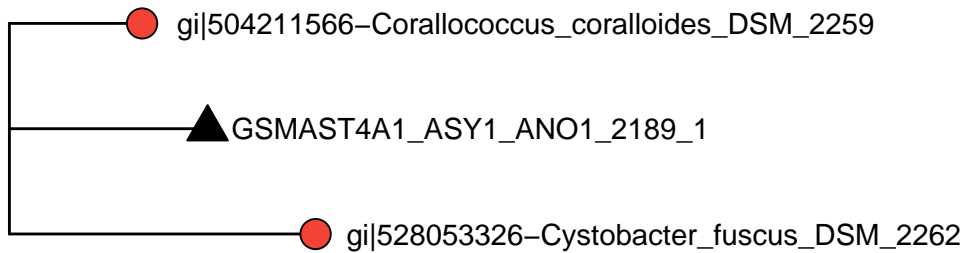
● Bacteria
★ Other
▲ SAG



- Bacteria
- ▲ Eukaryota
- ▲ SAG



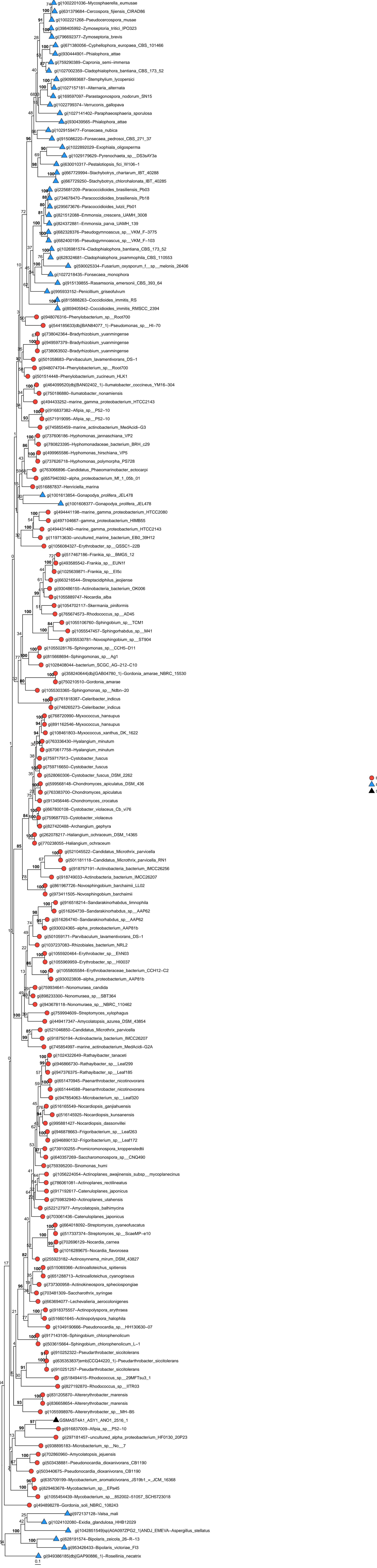


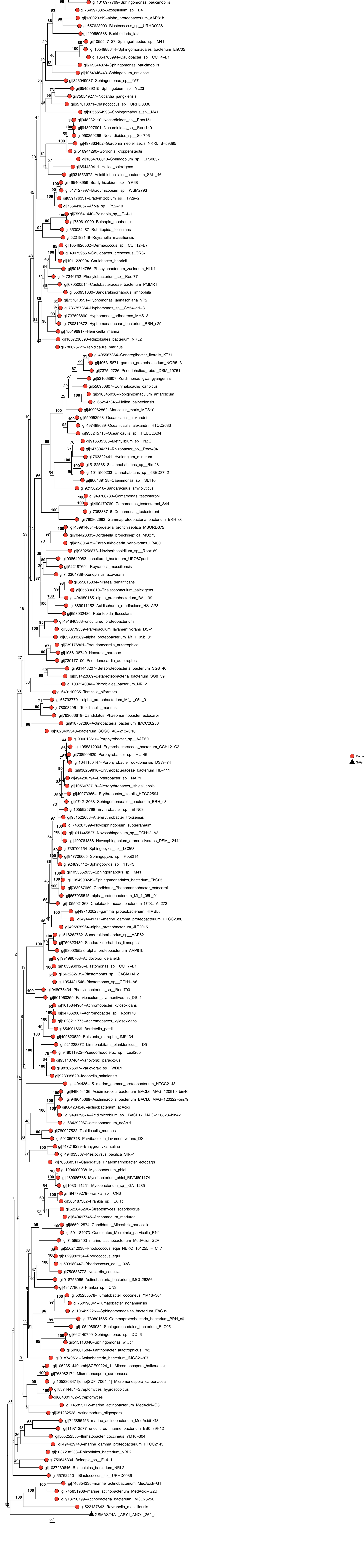


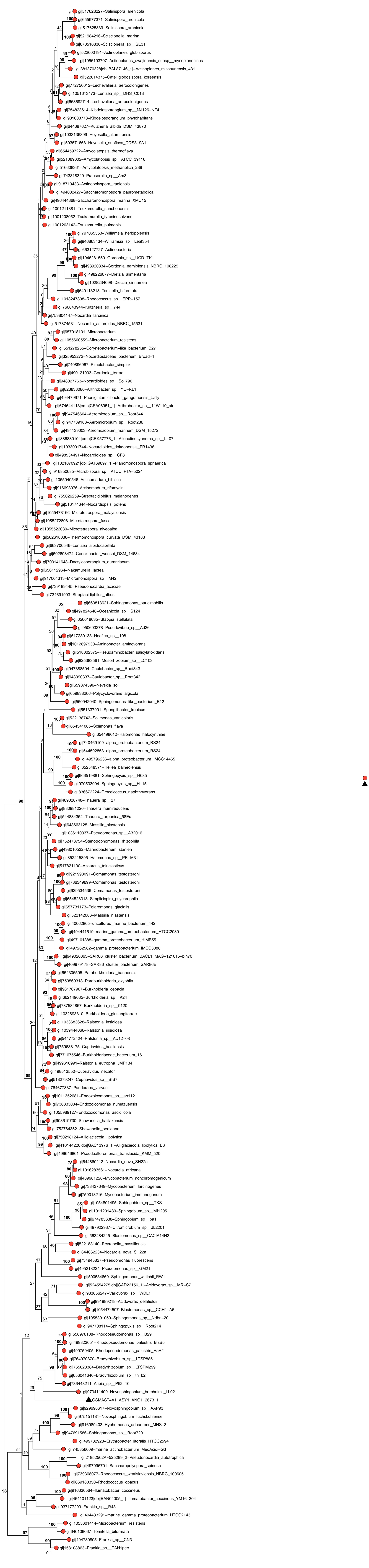
● Bacteria
▲ SAG

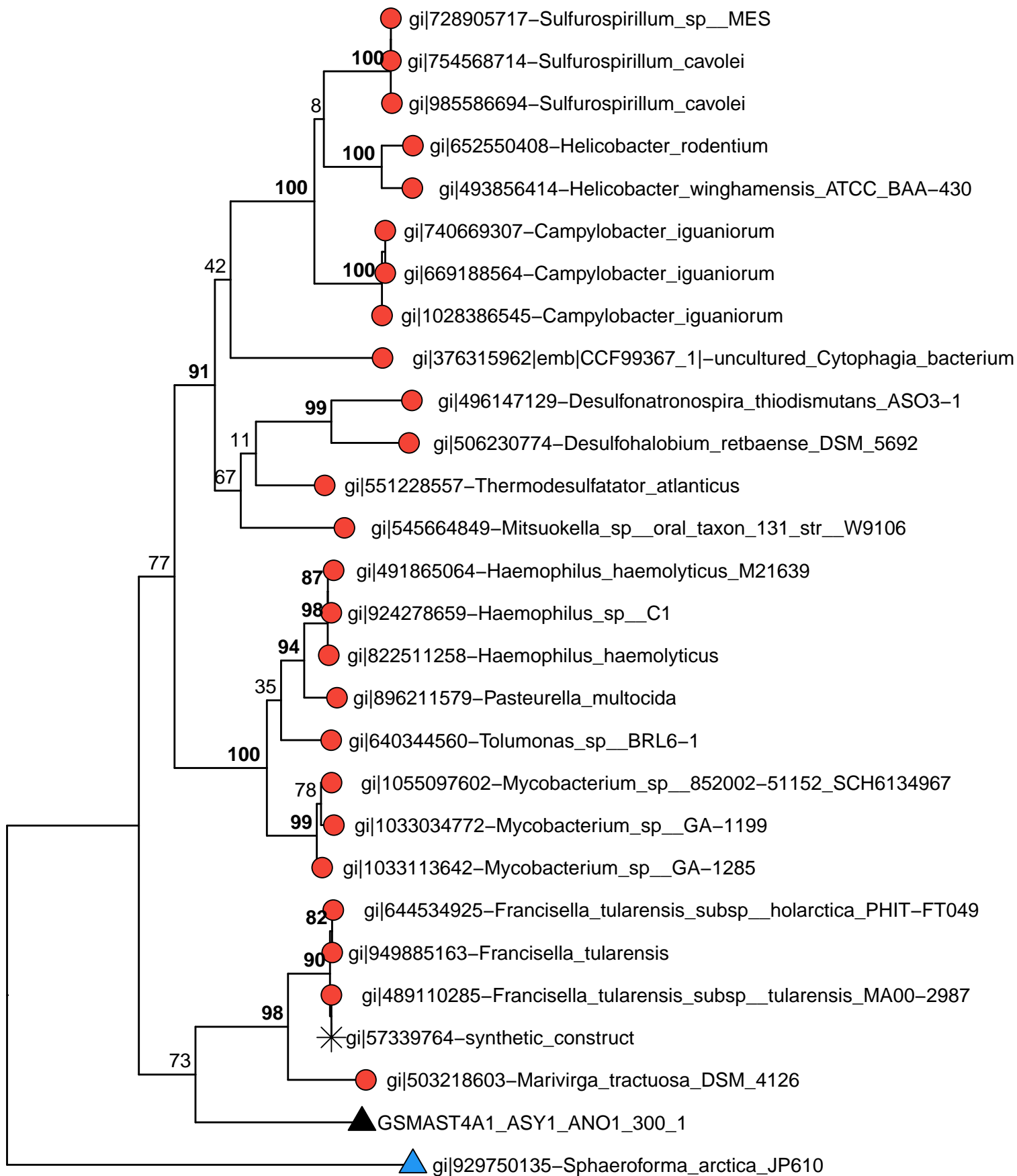
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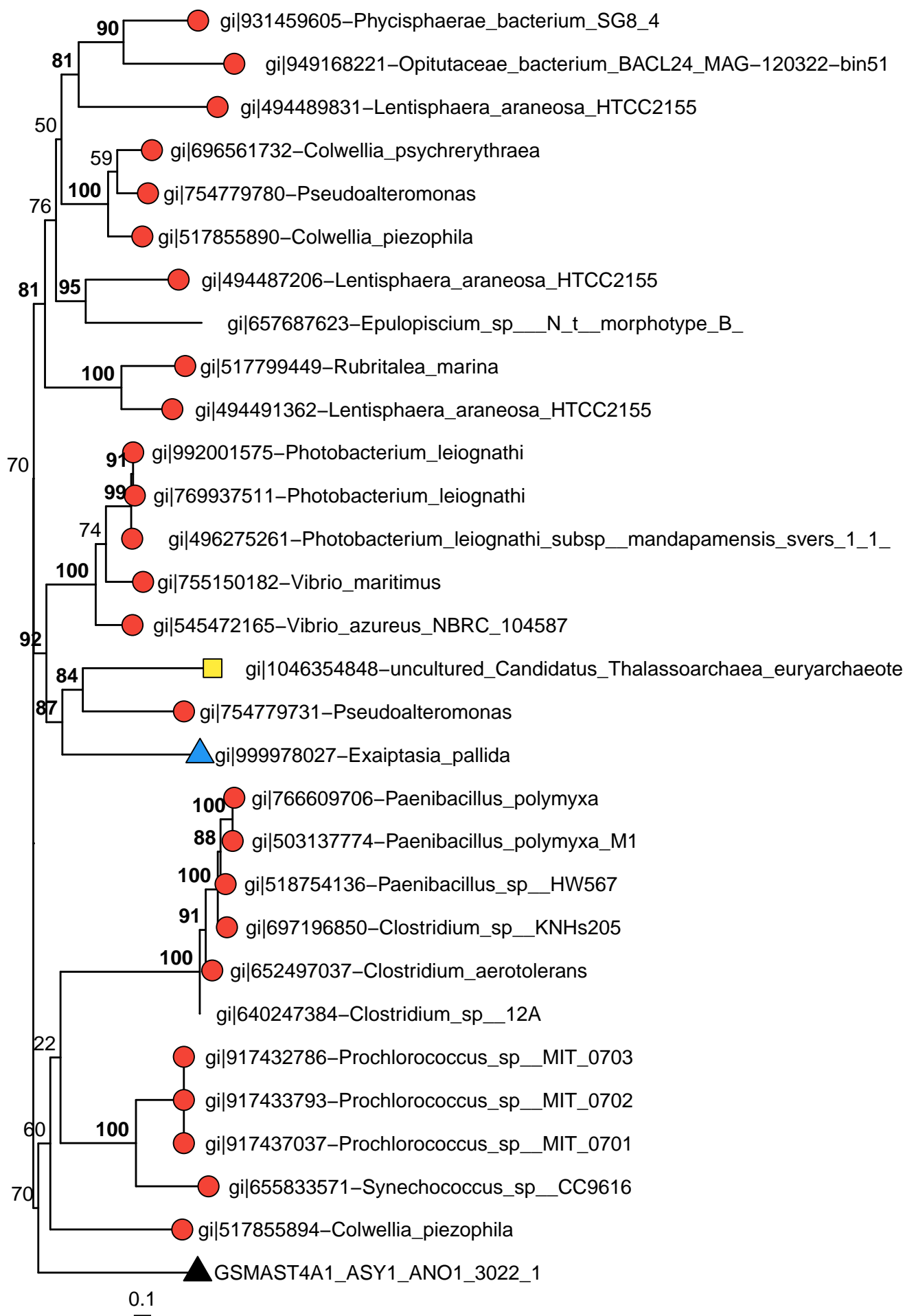


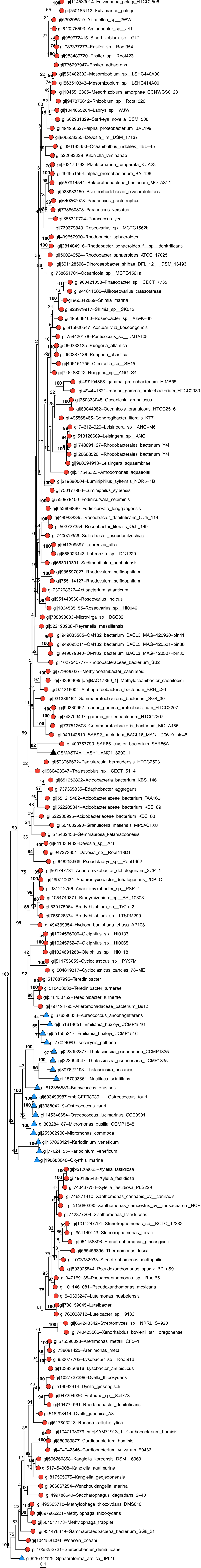


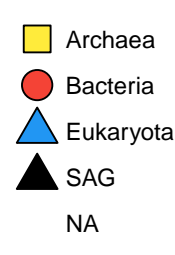
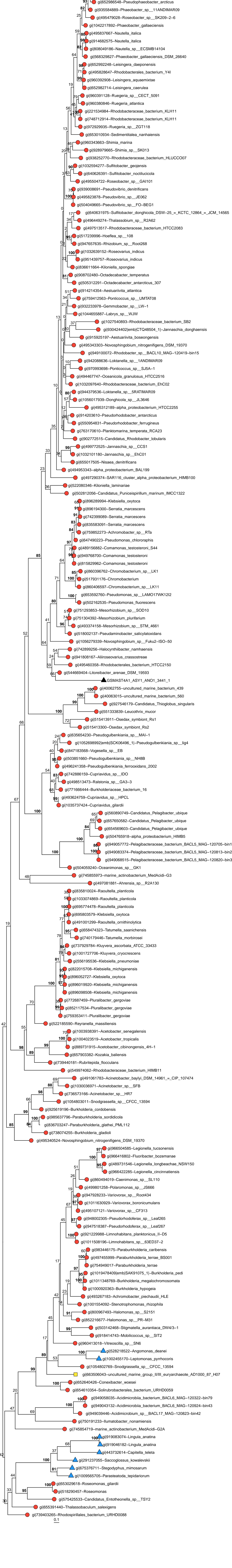


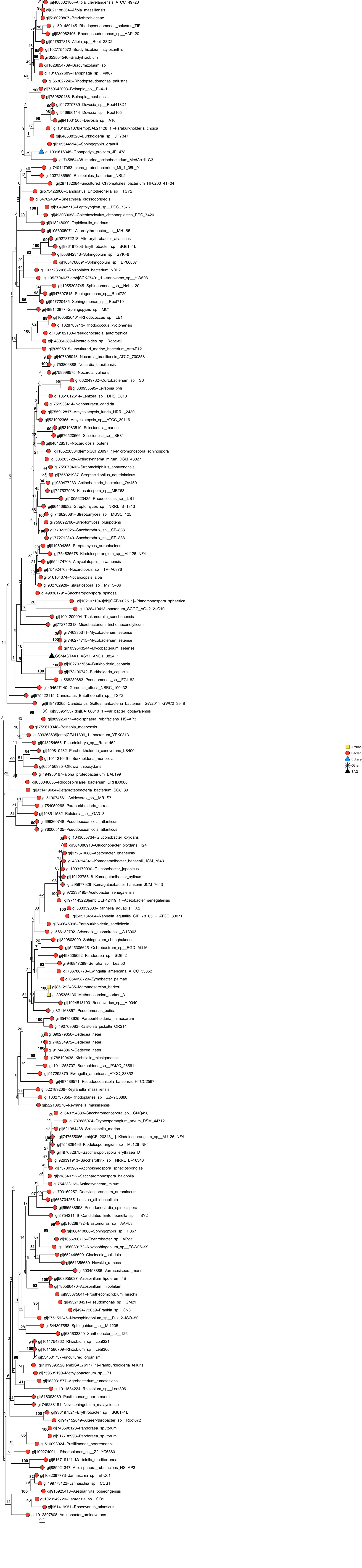


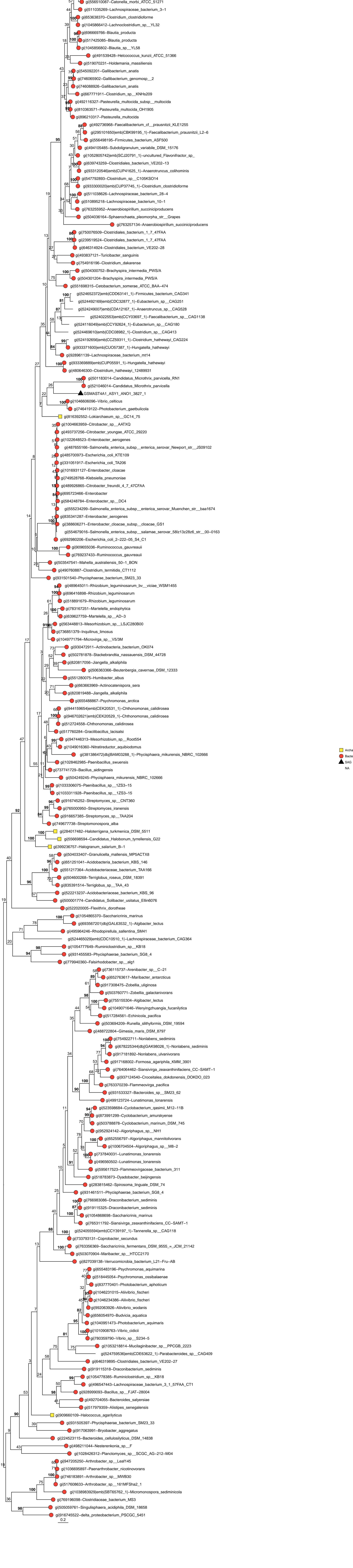
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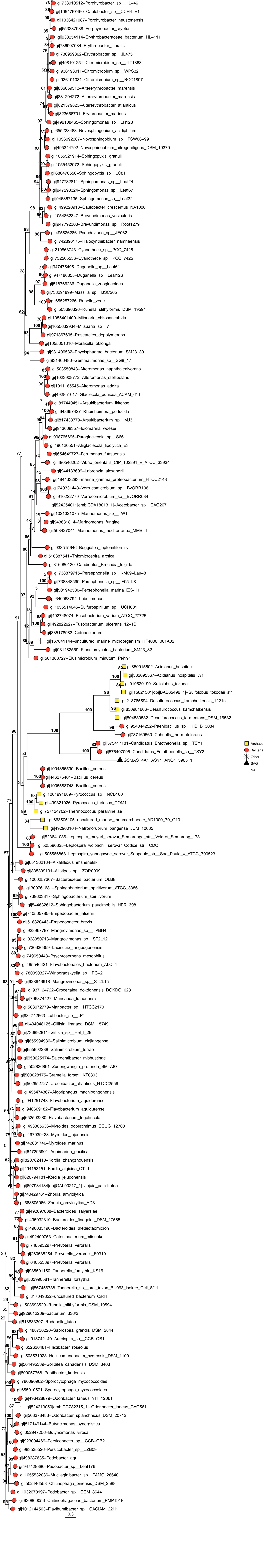


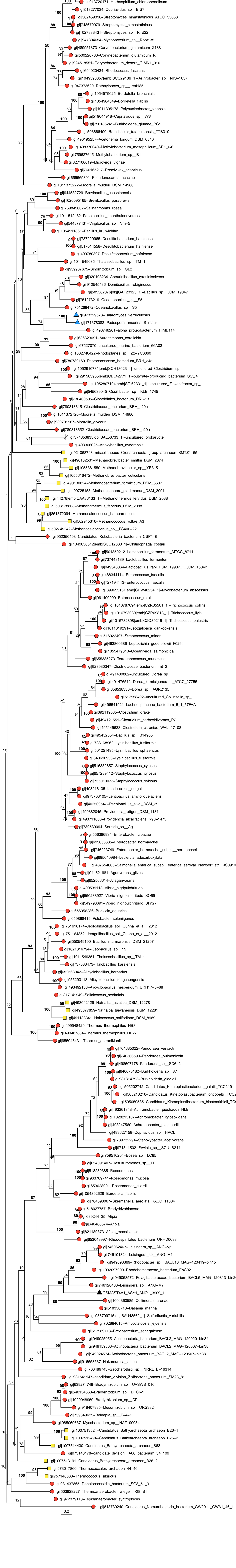


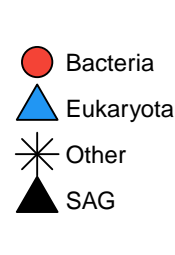
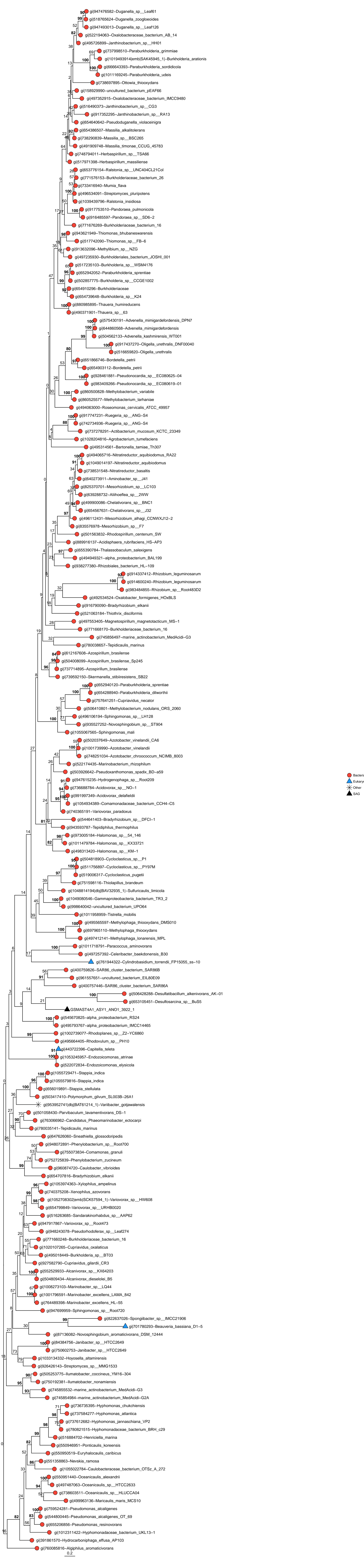




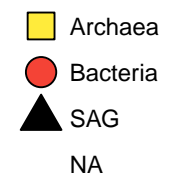
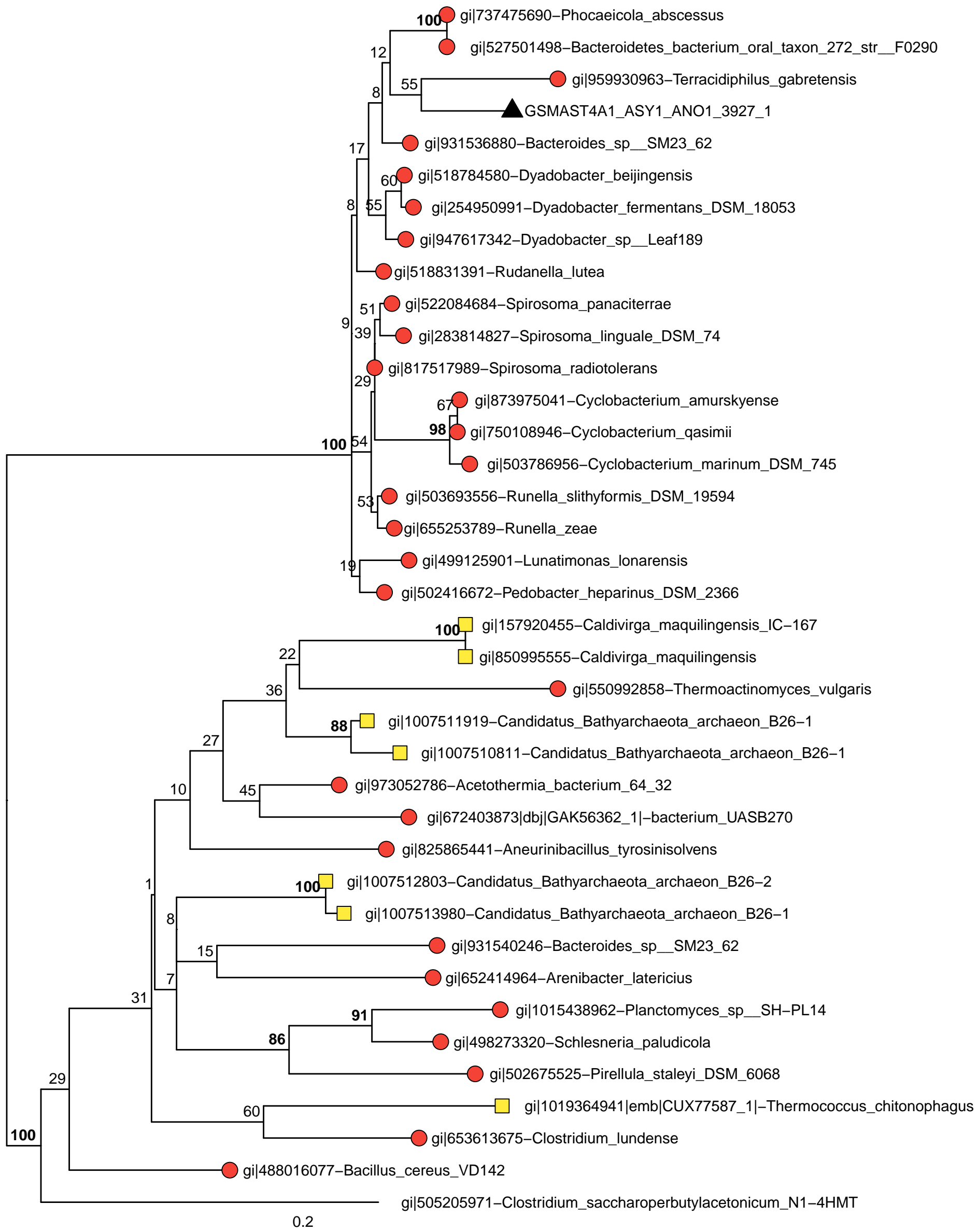




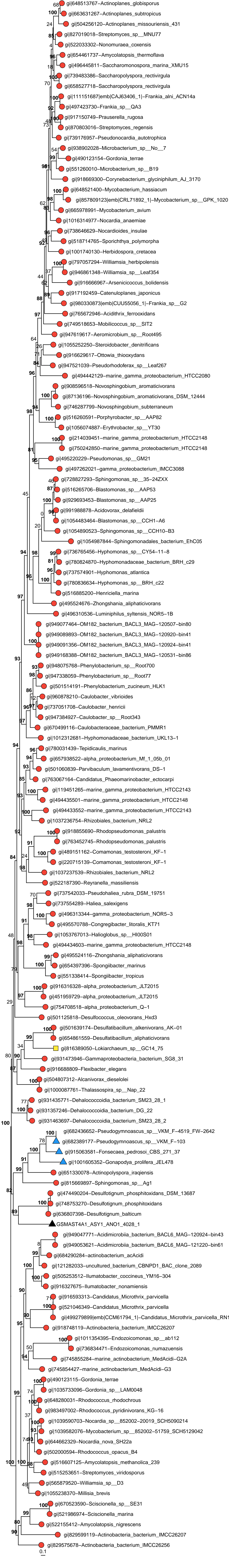


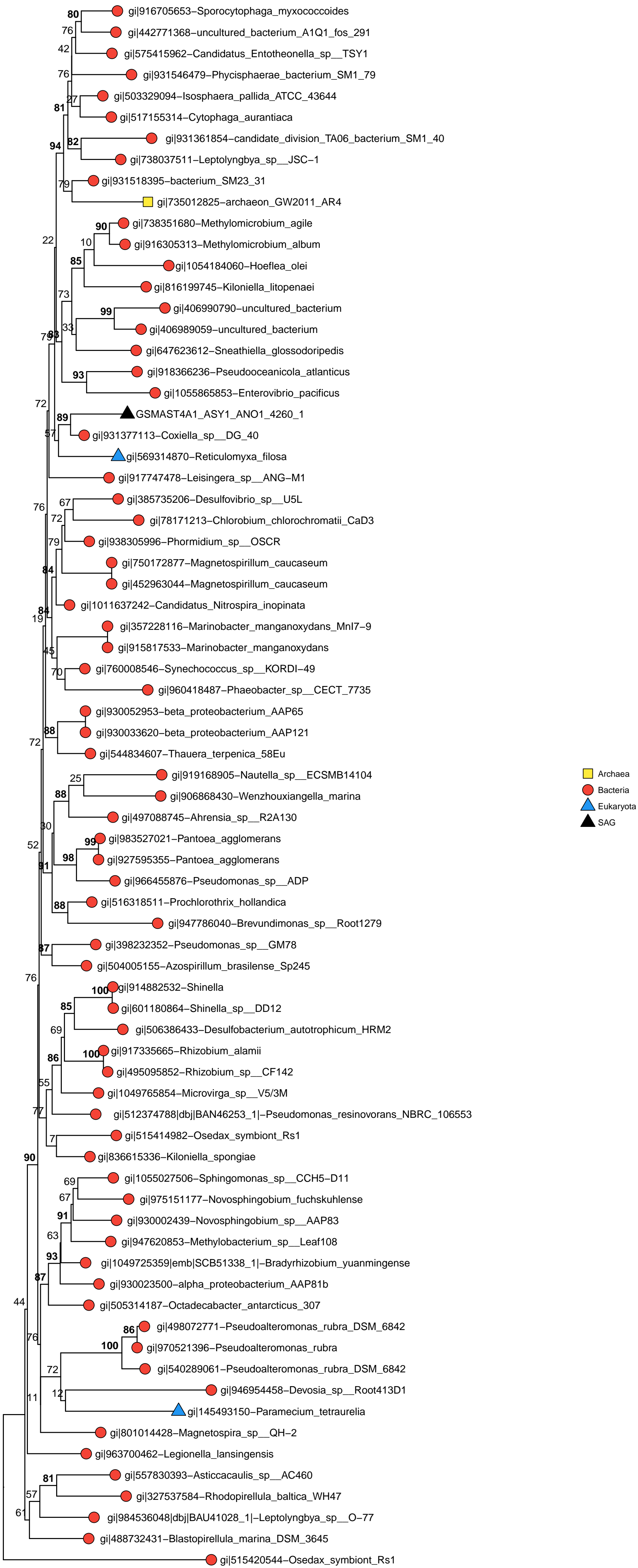


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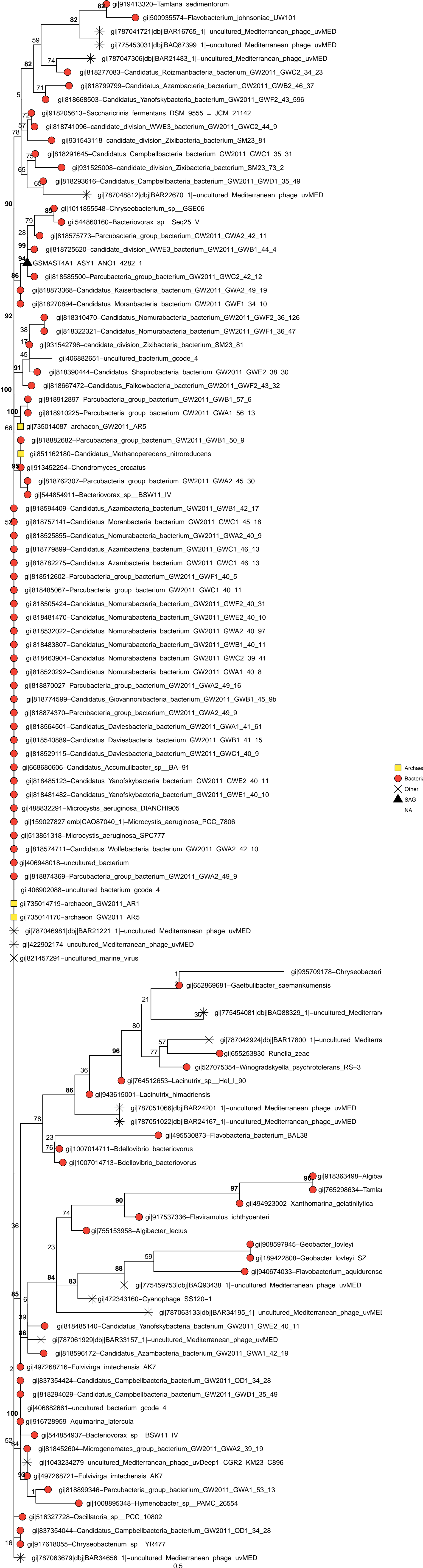


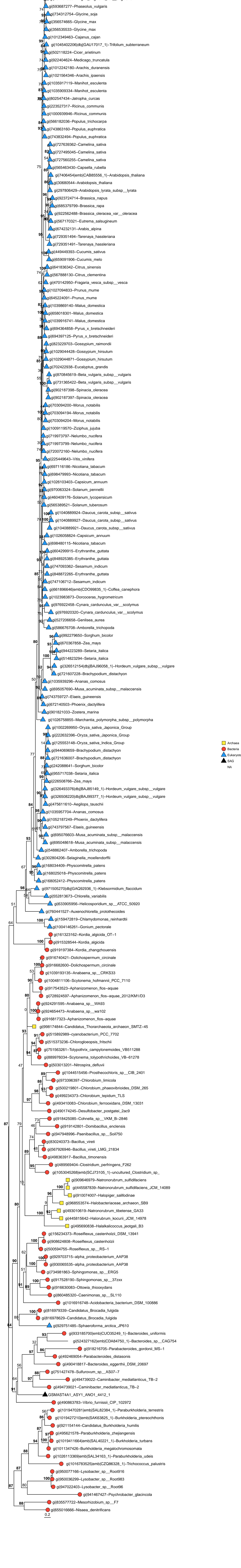
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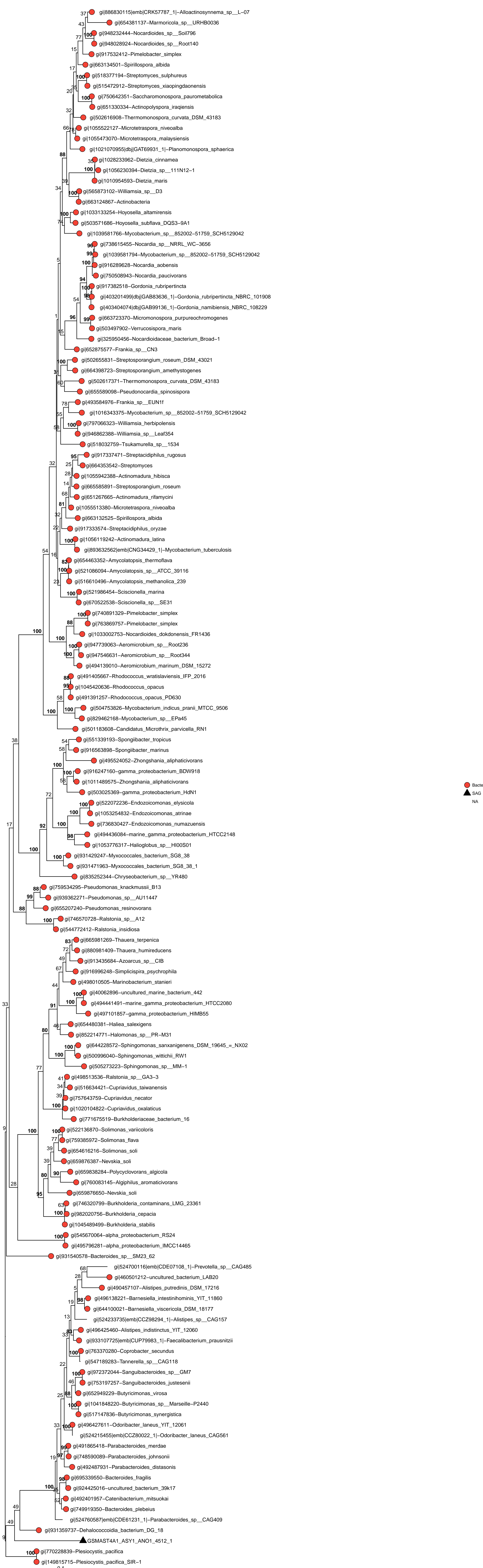


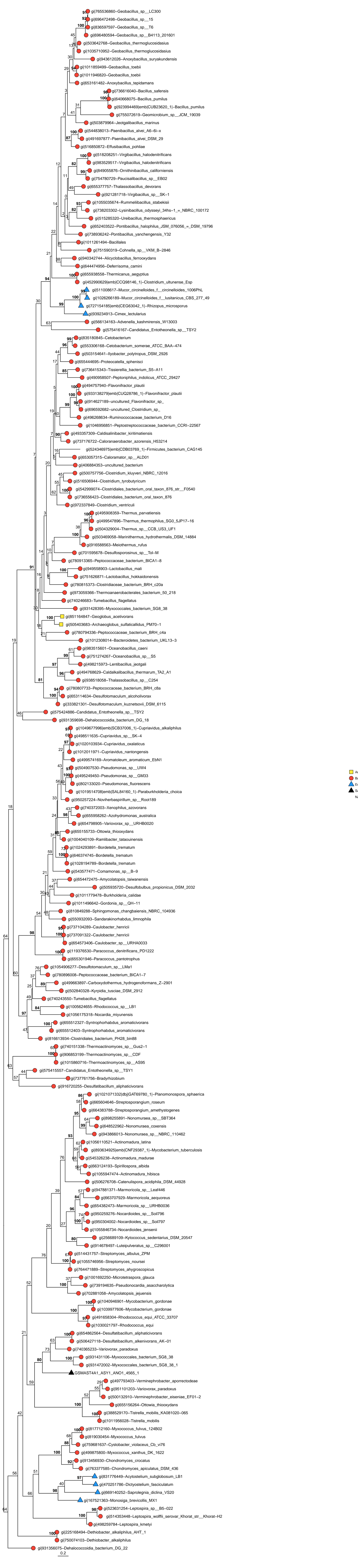


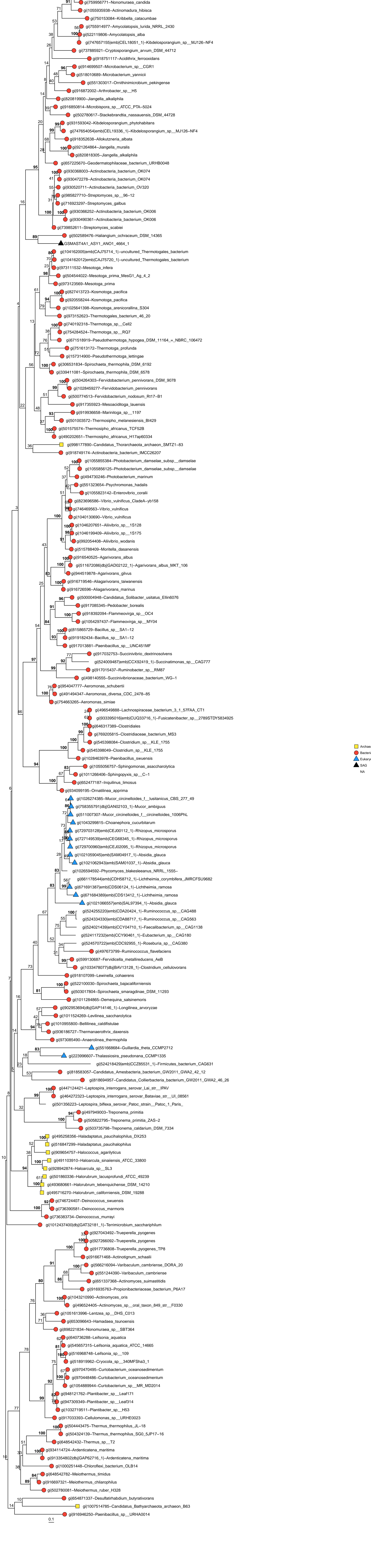
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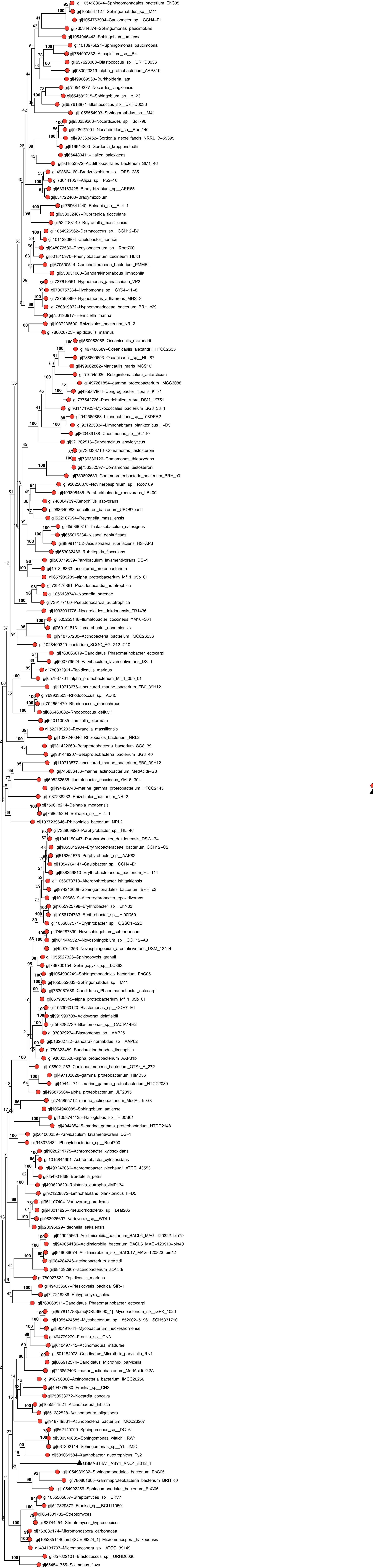




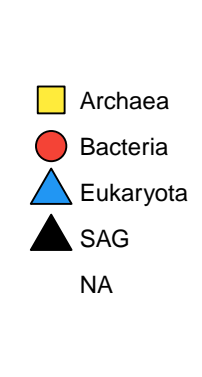
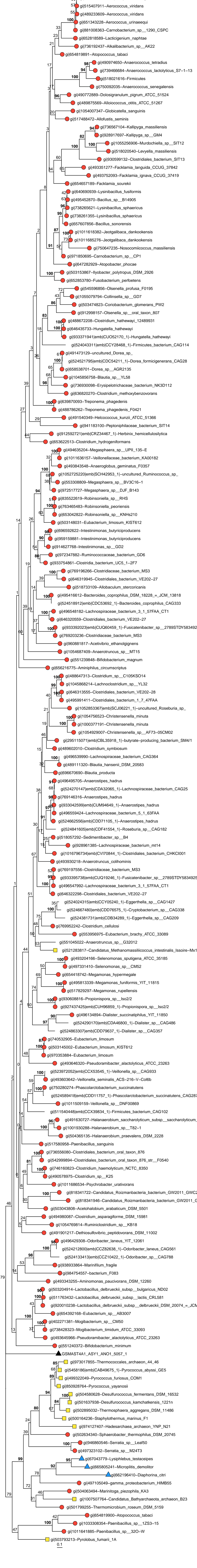


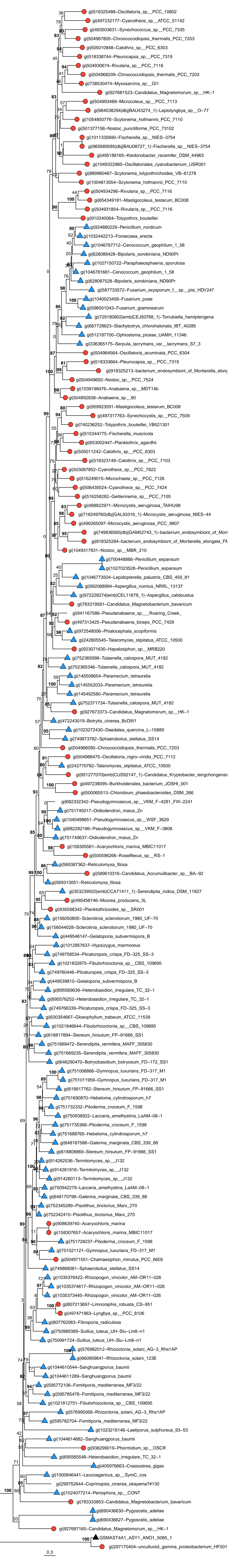


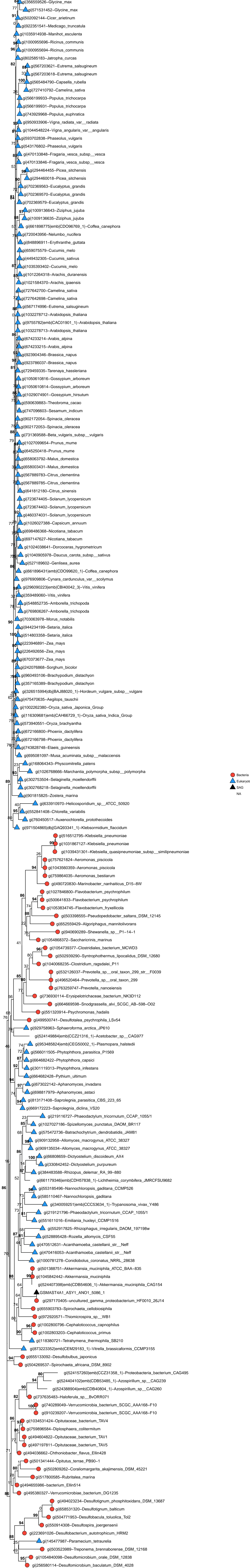


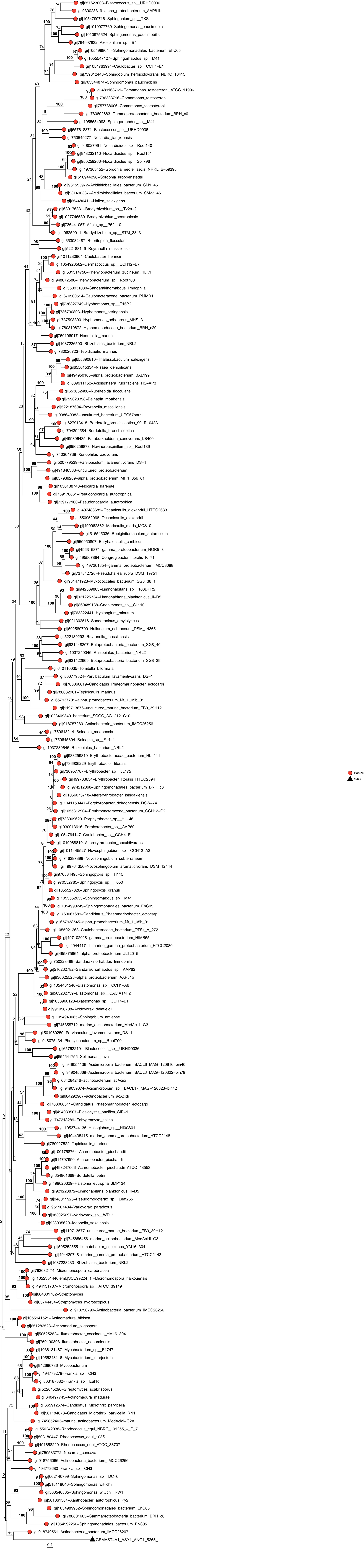


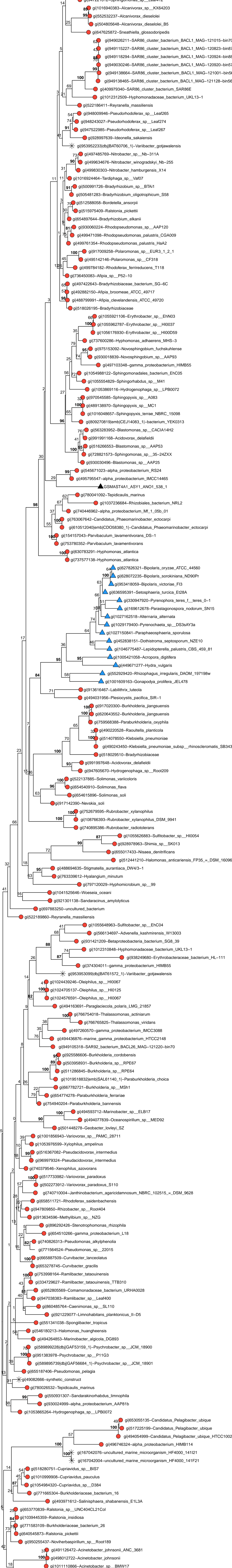
● Bacteria
▲ SAG



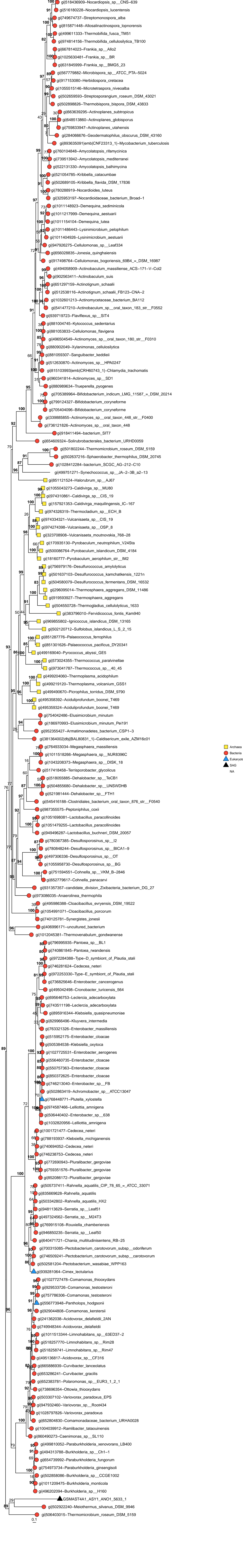


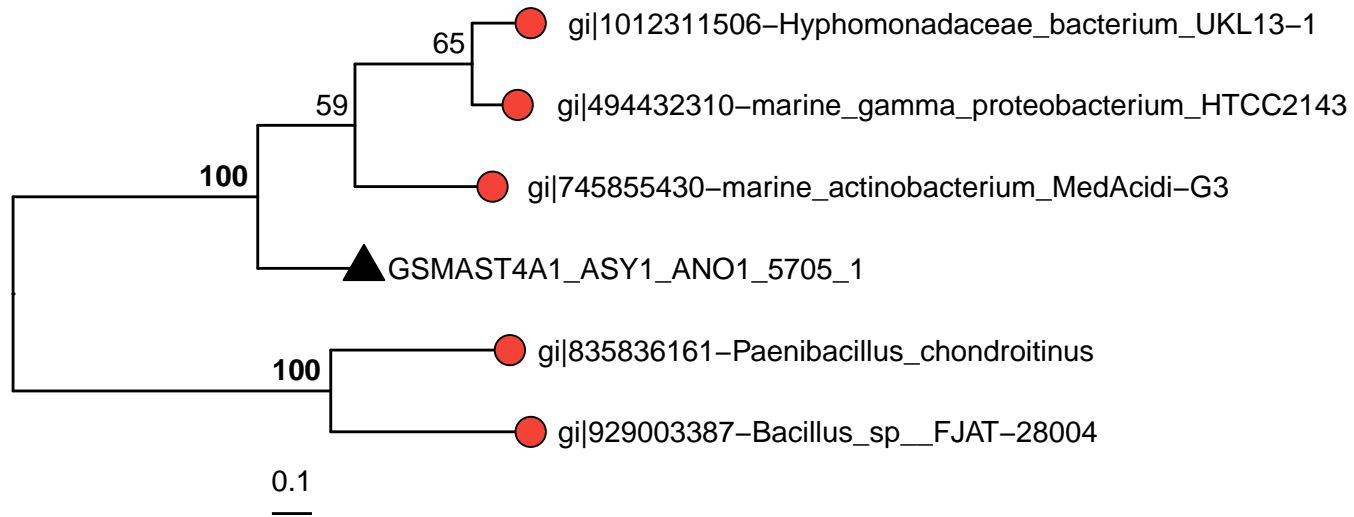




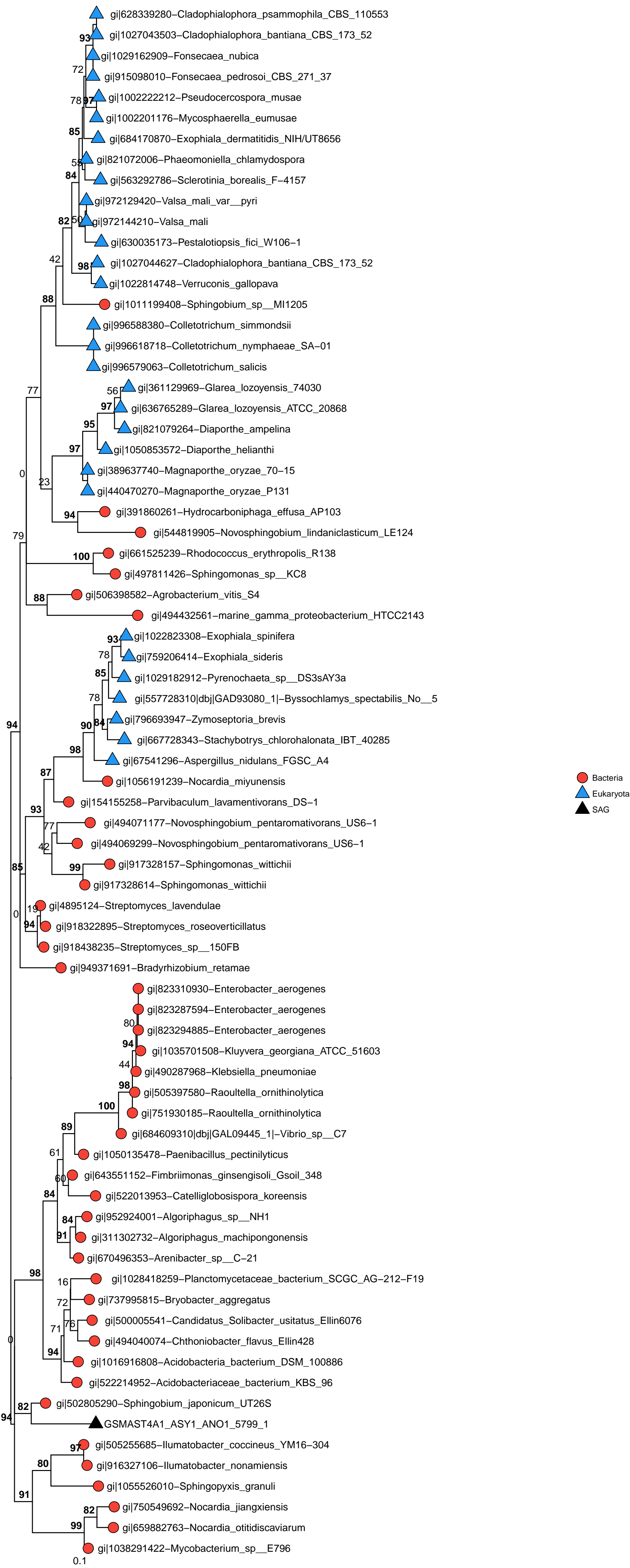


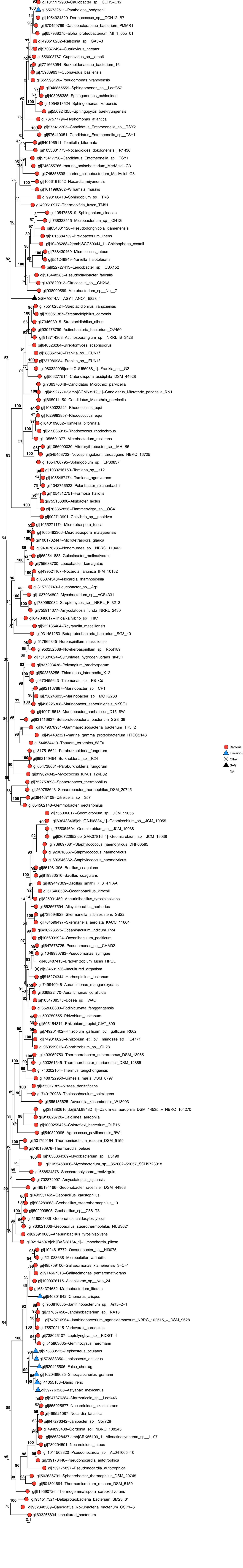
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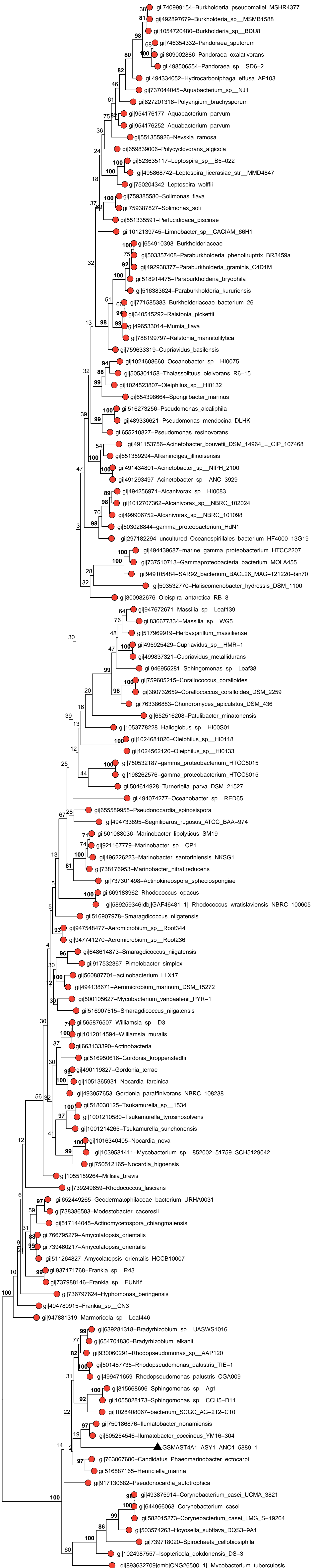


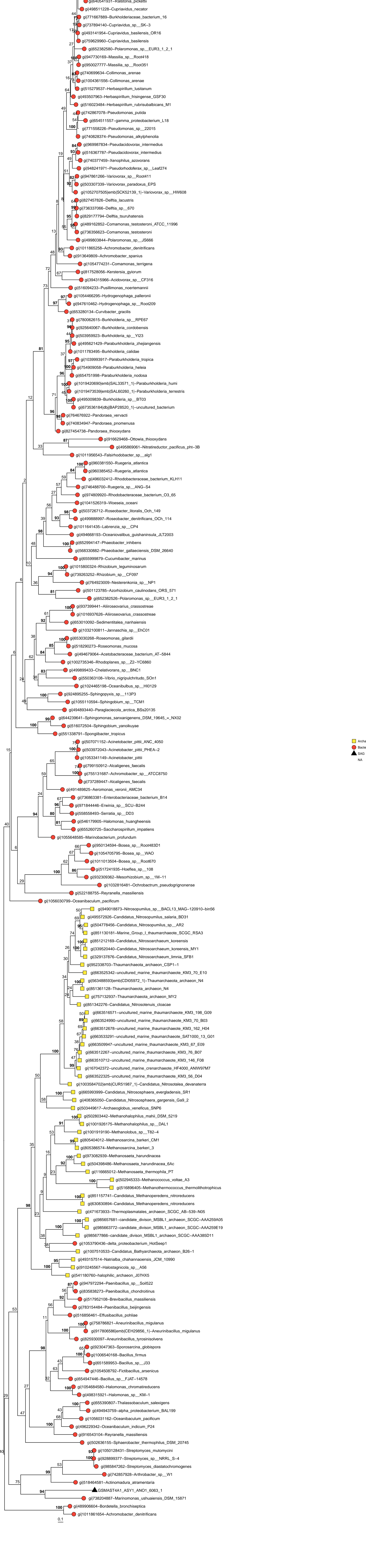


● Bacteria
▲ SAG



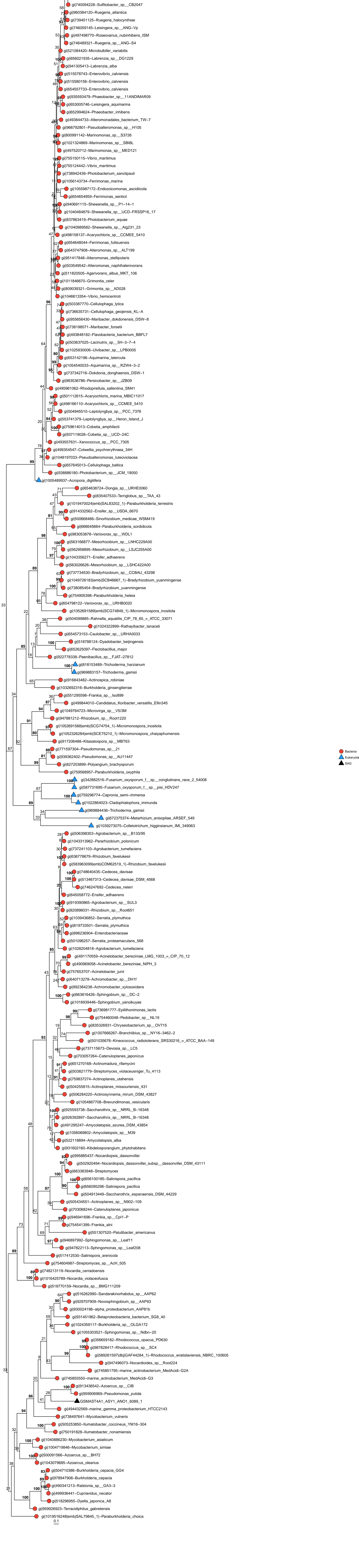


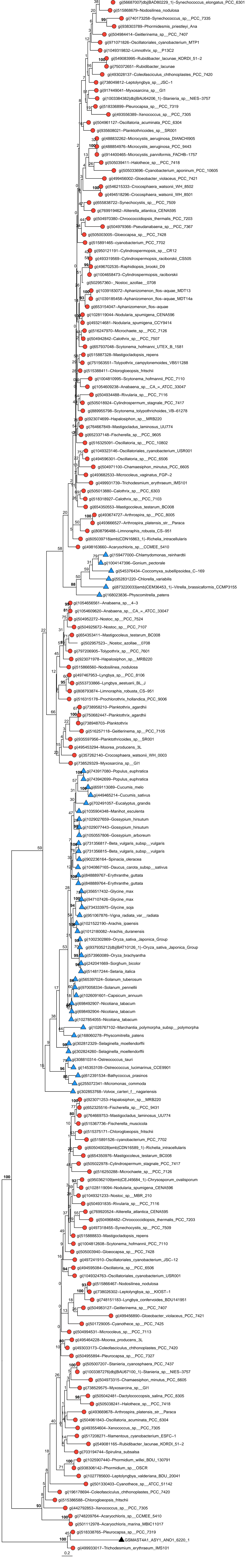




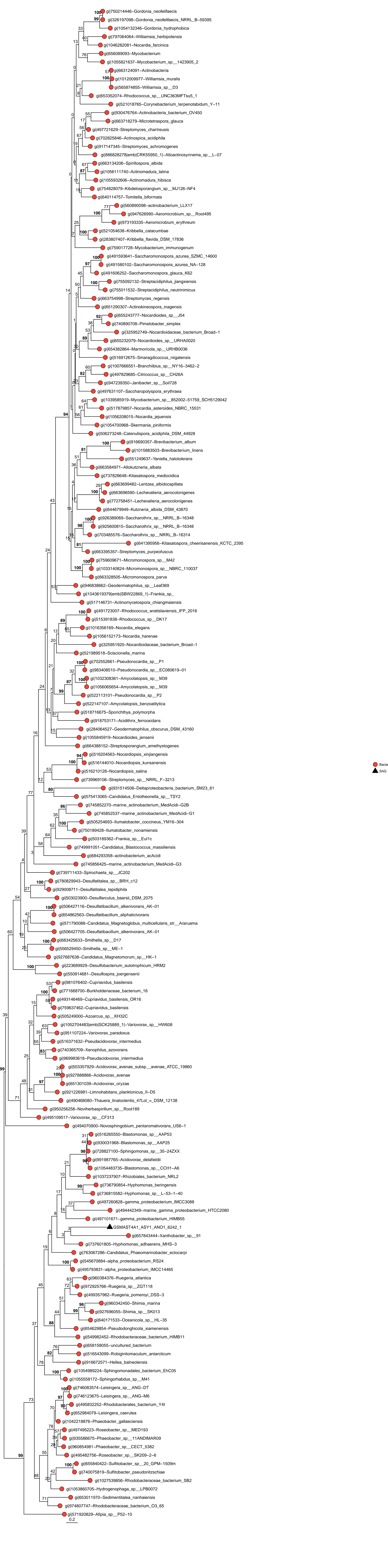
■ Archaea
● Bacteria
▲ SAG
■ NA

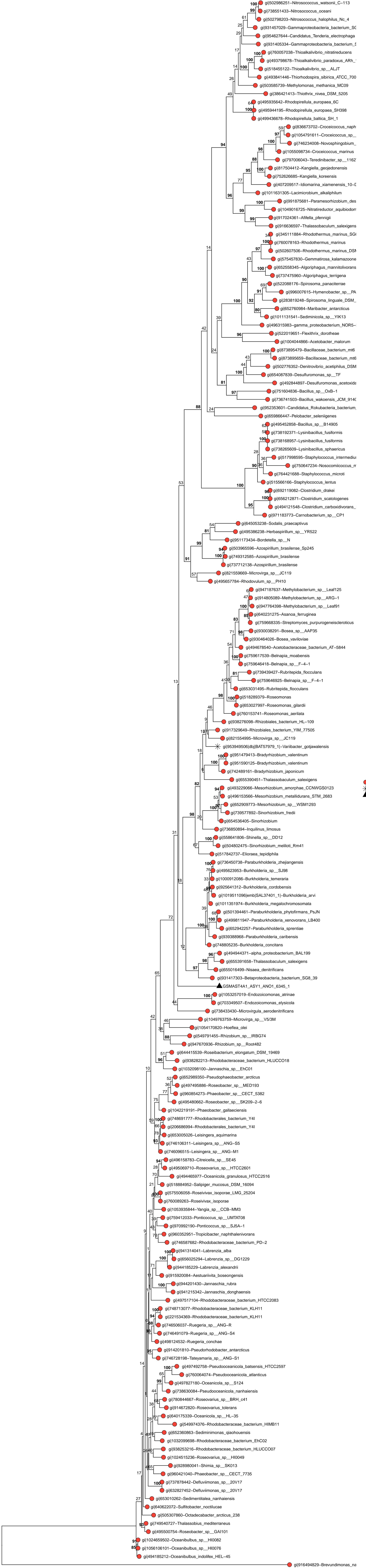
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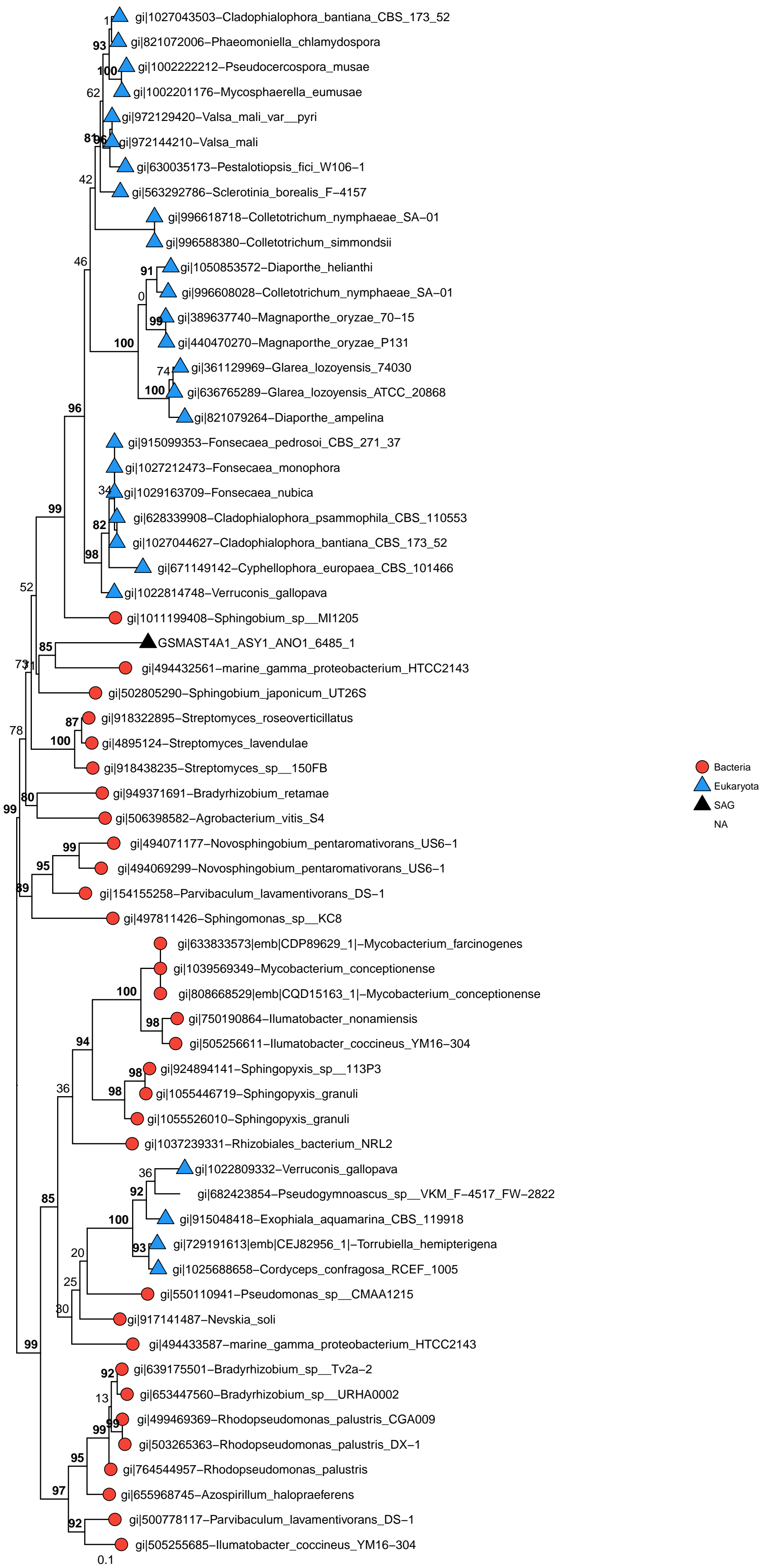


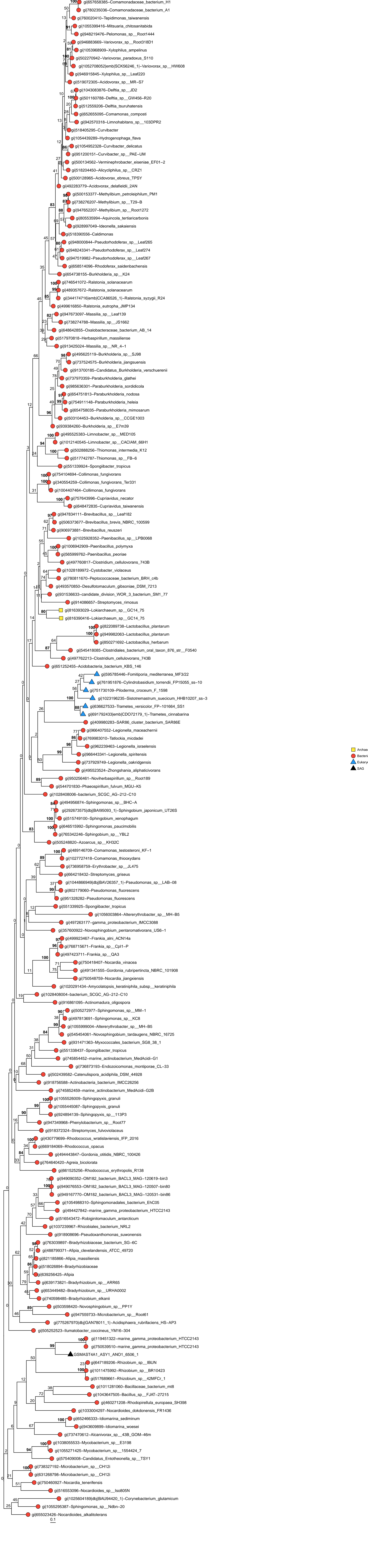
● Bacteria
 ▲ Eukaryota
 ▲ SAG
 NA



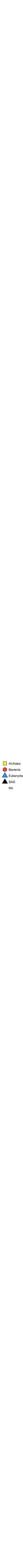
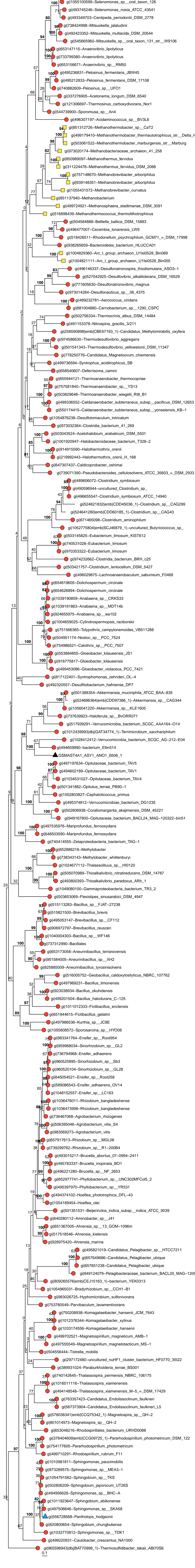


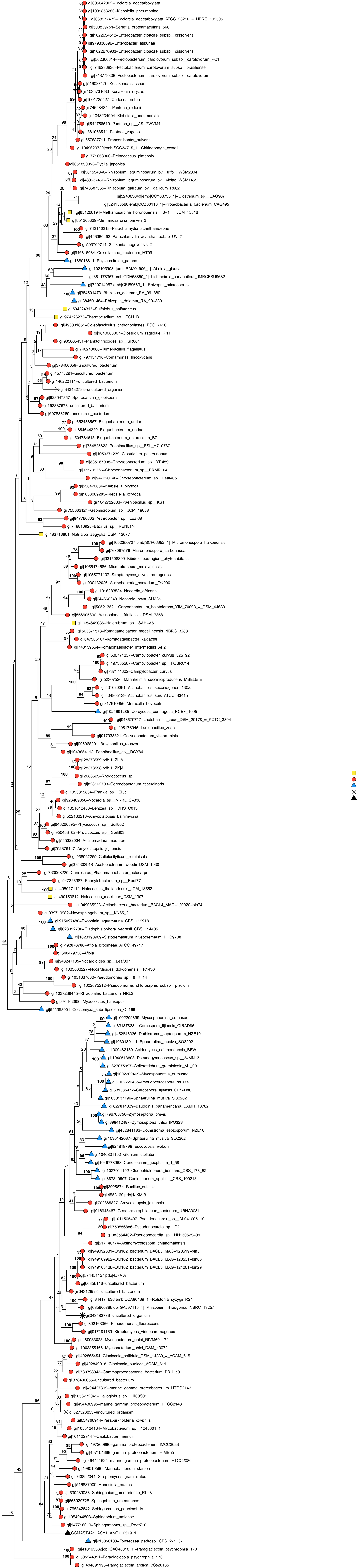
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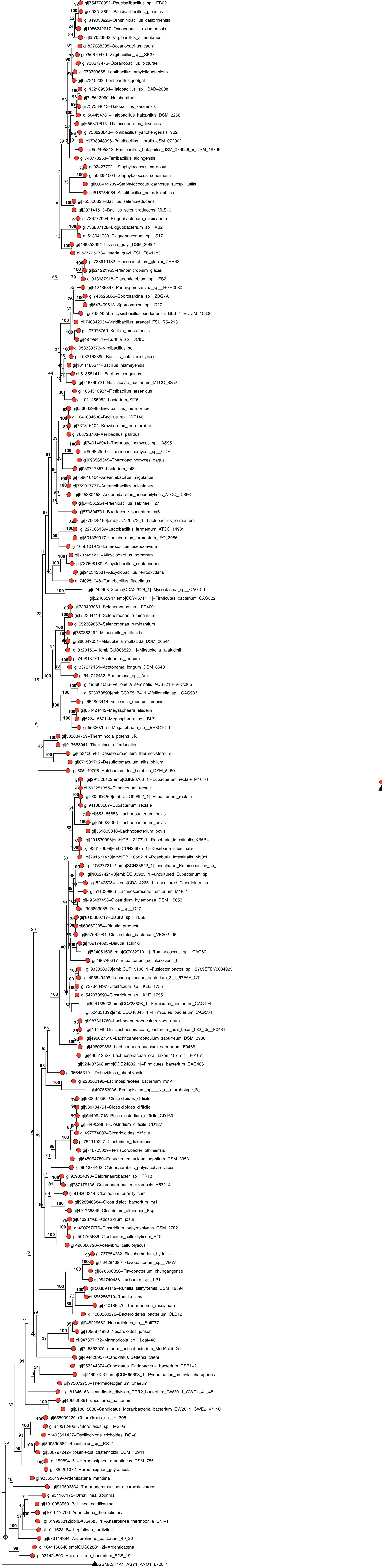


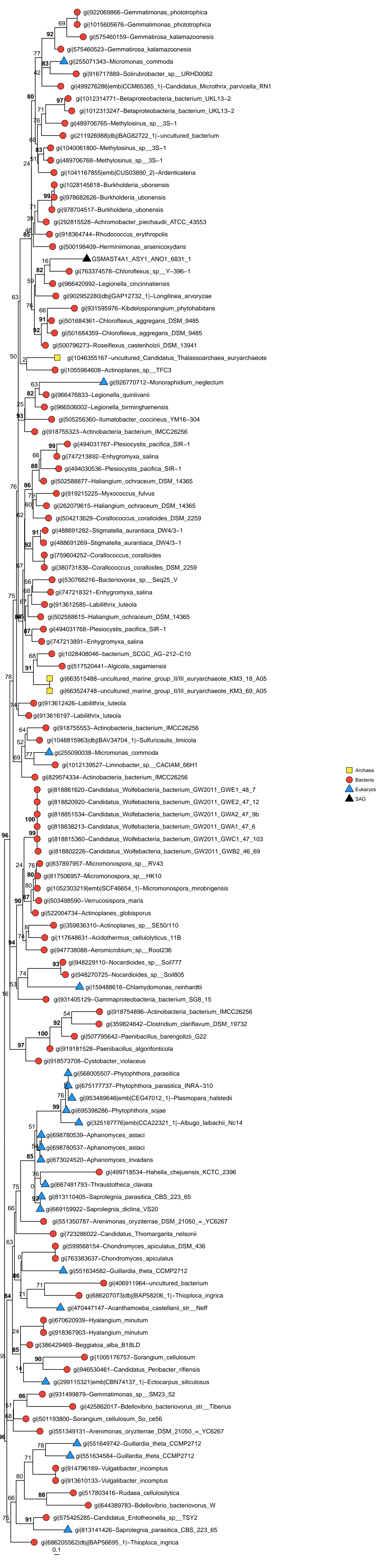


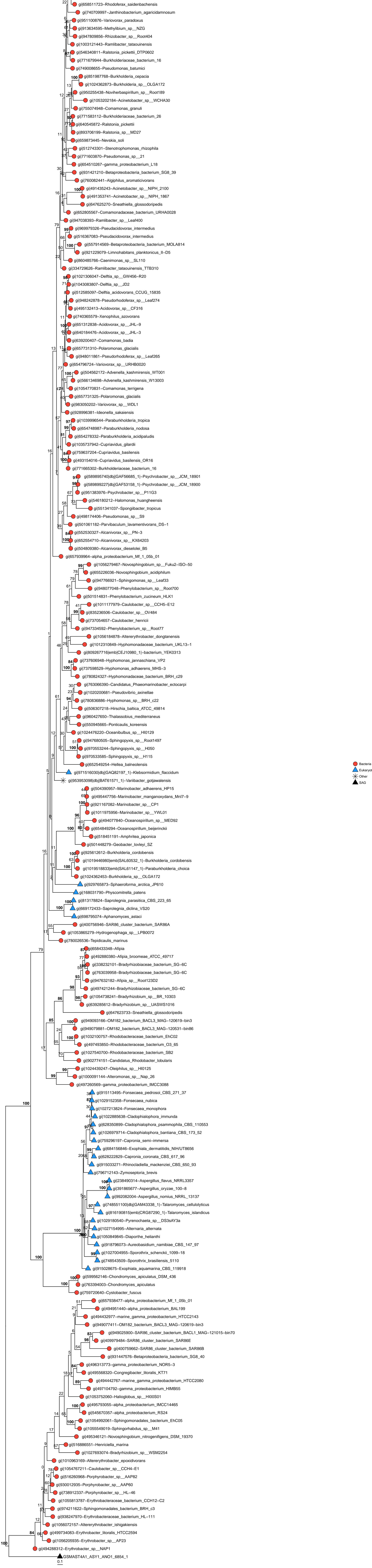
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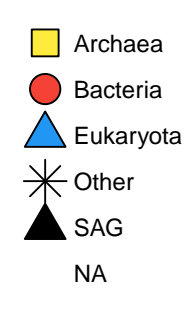
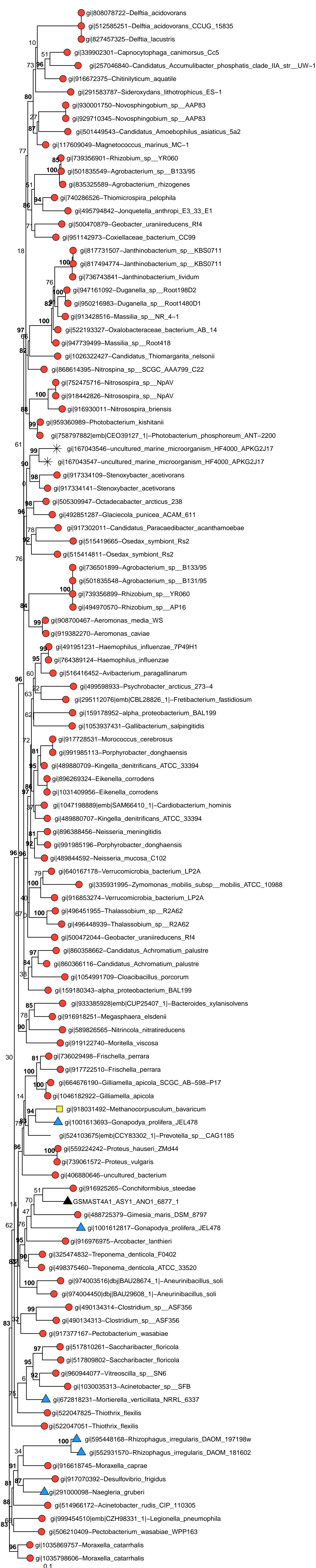


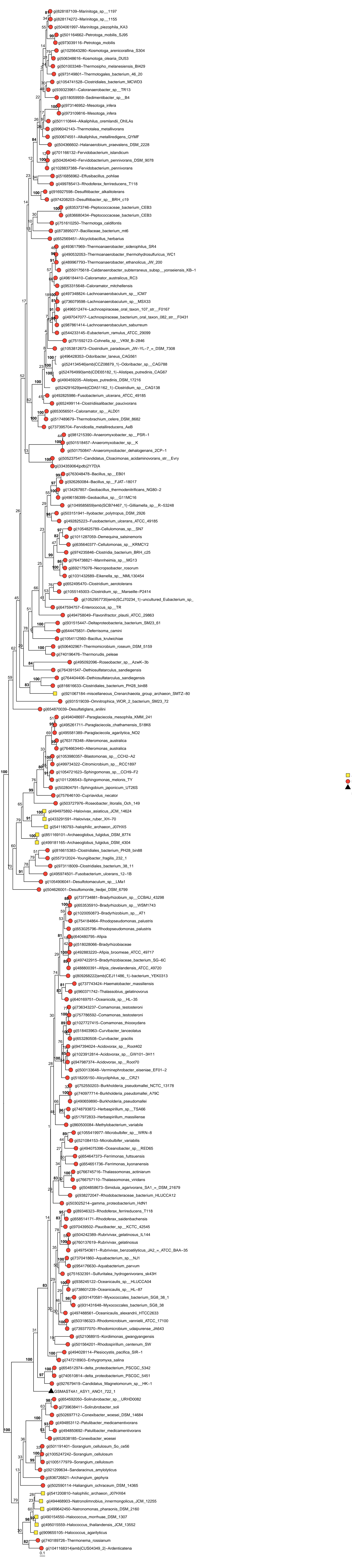


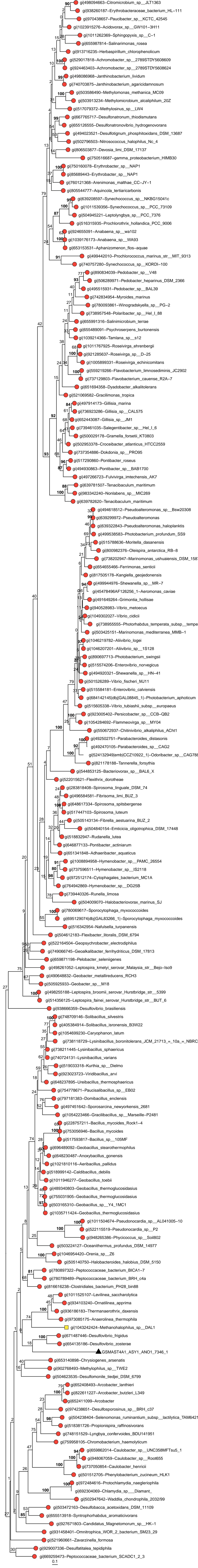


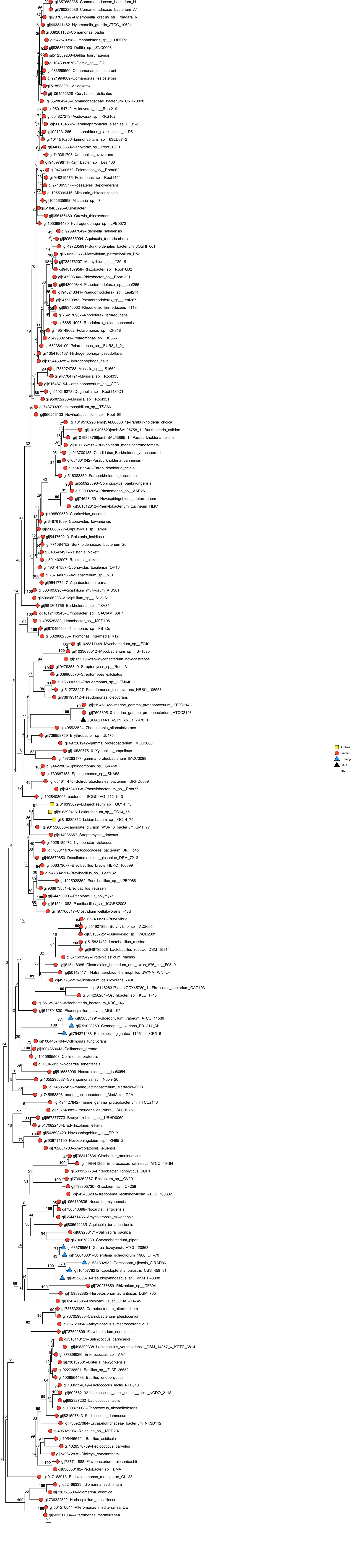


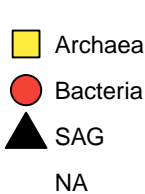
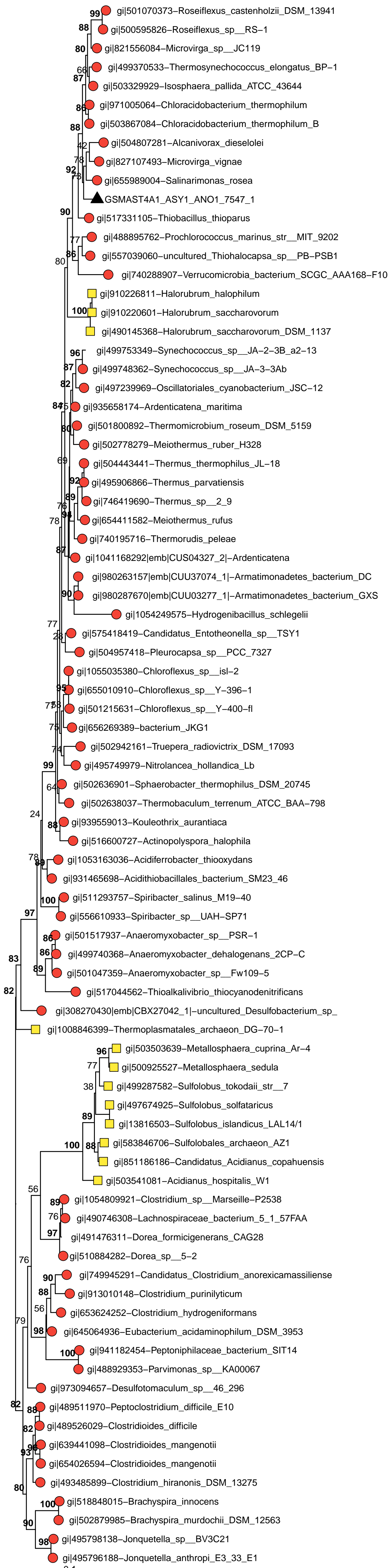


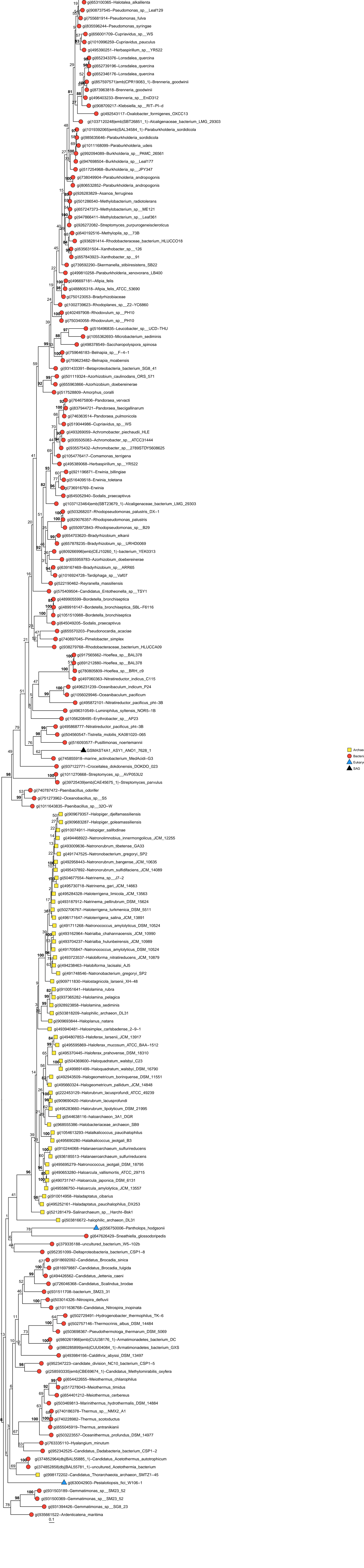


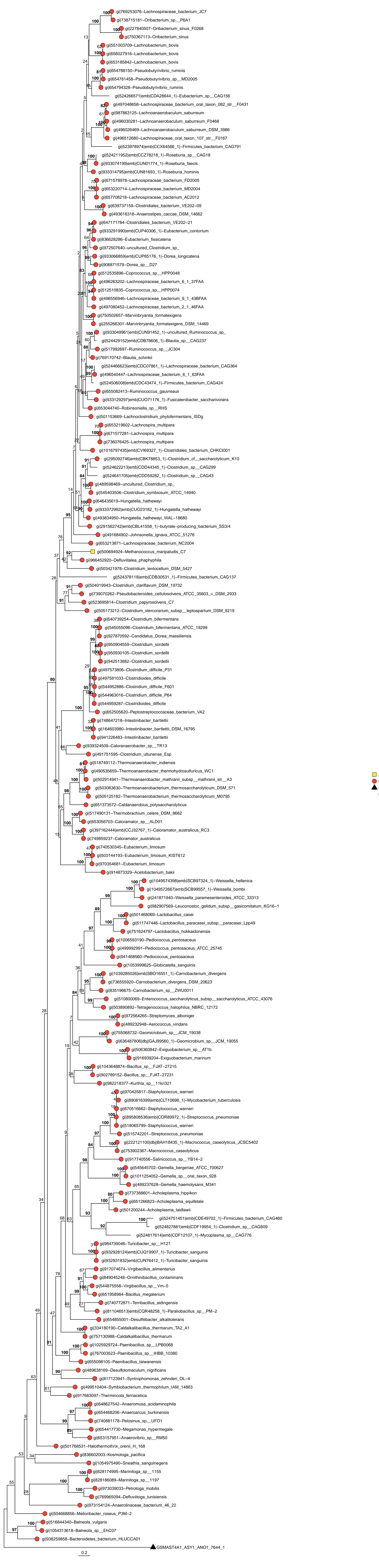




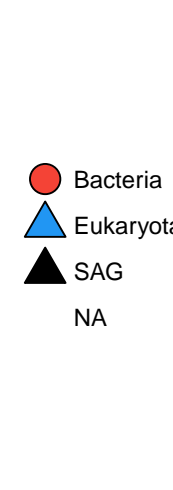




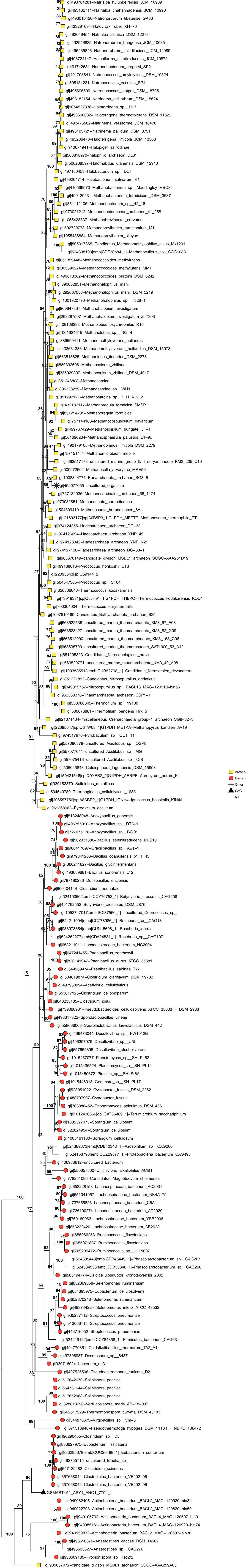


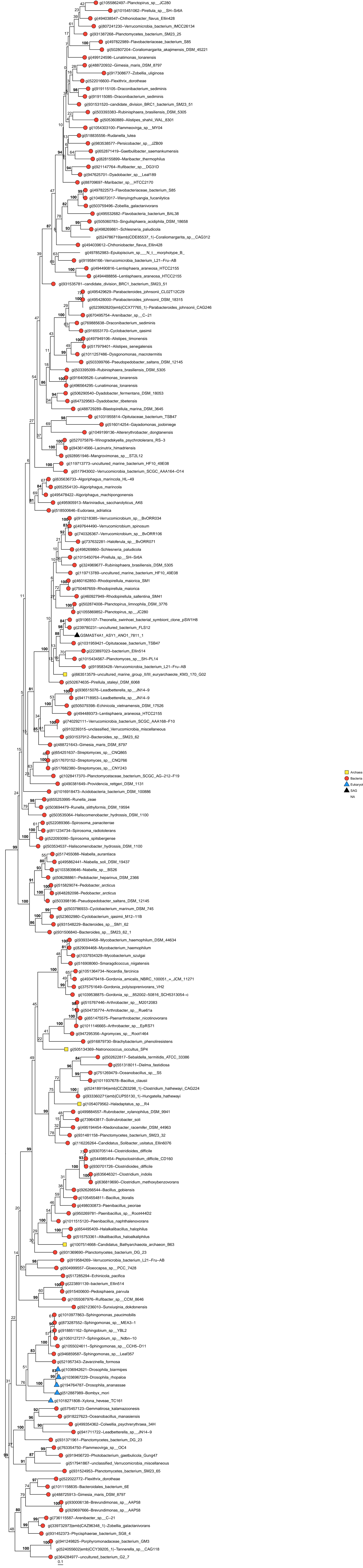


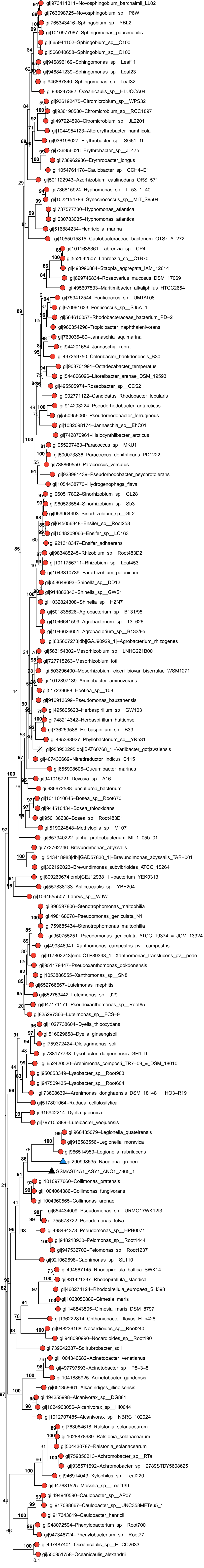
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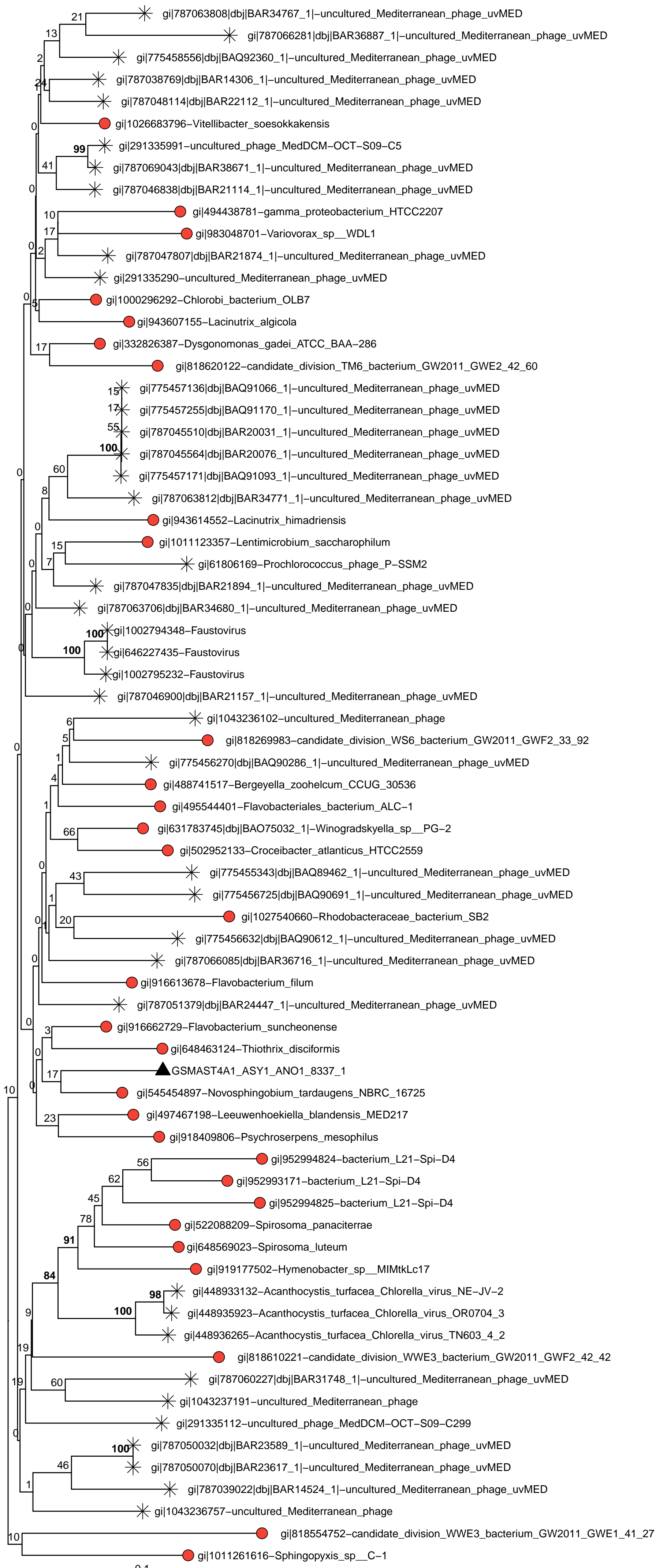
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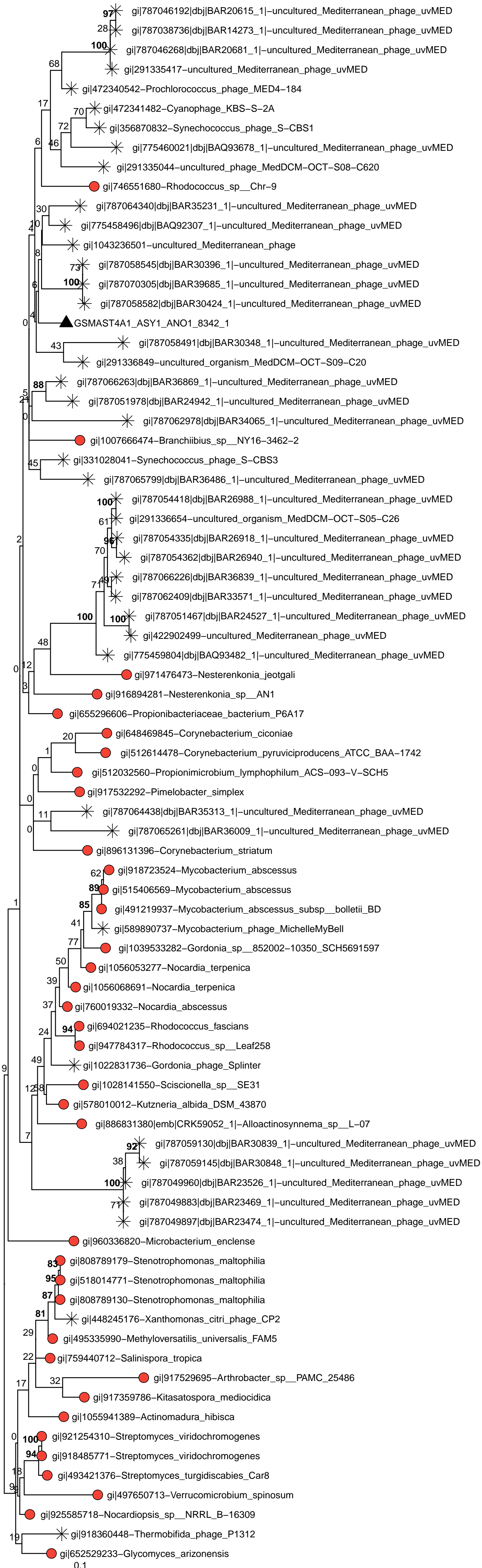


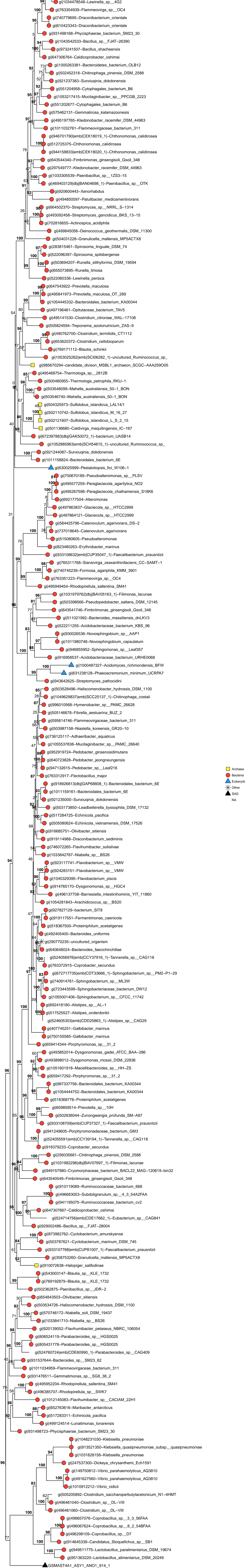


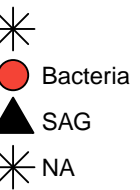
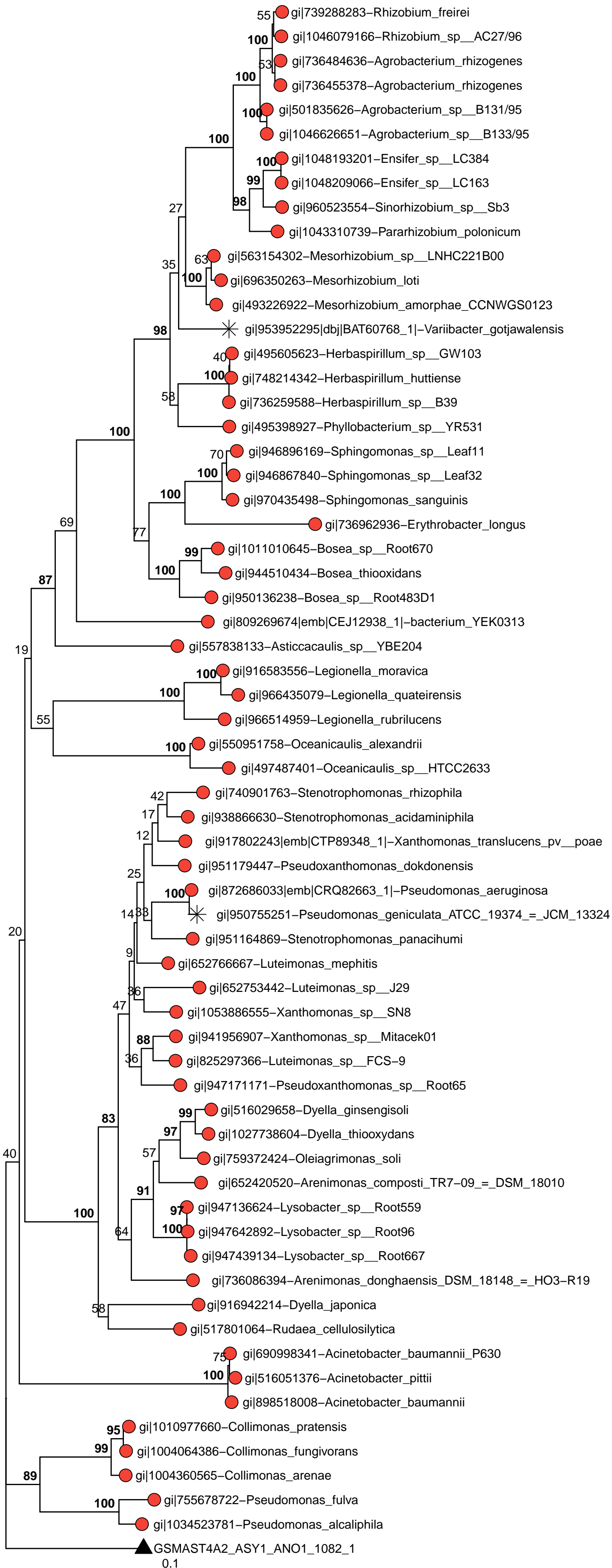


- Bacteria
- ▲ Eukaryota
- ✱ Other
- ▲ SAG
- NA

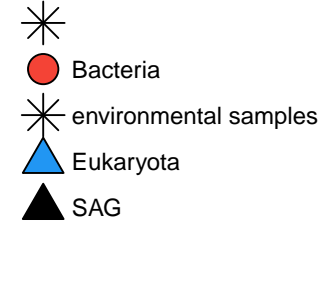
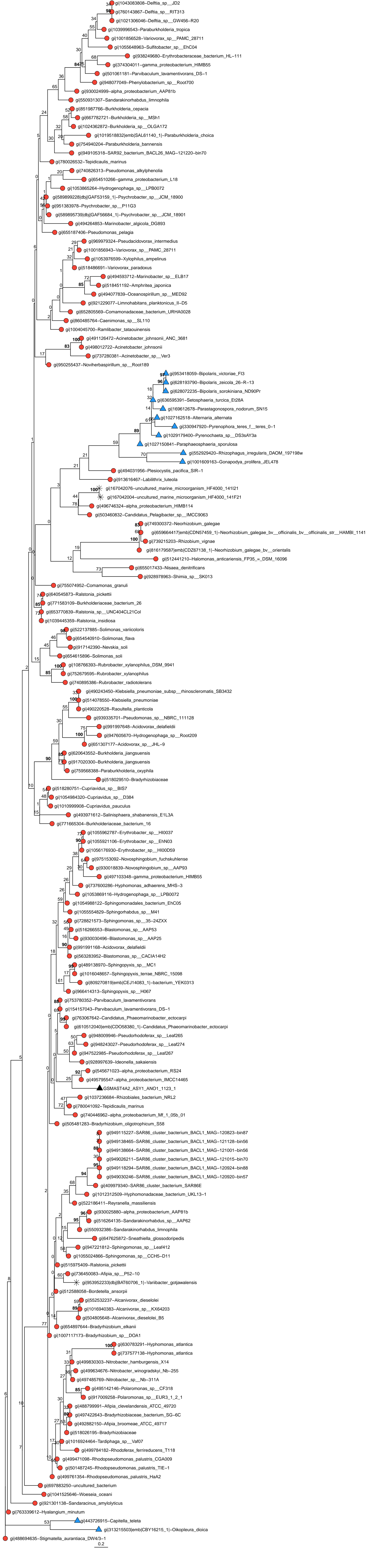




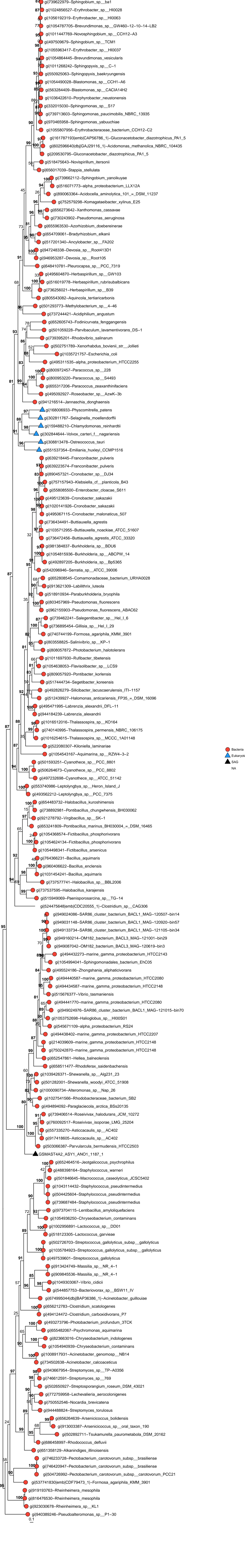


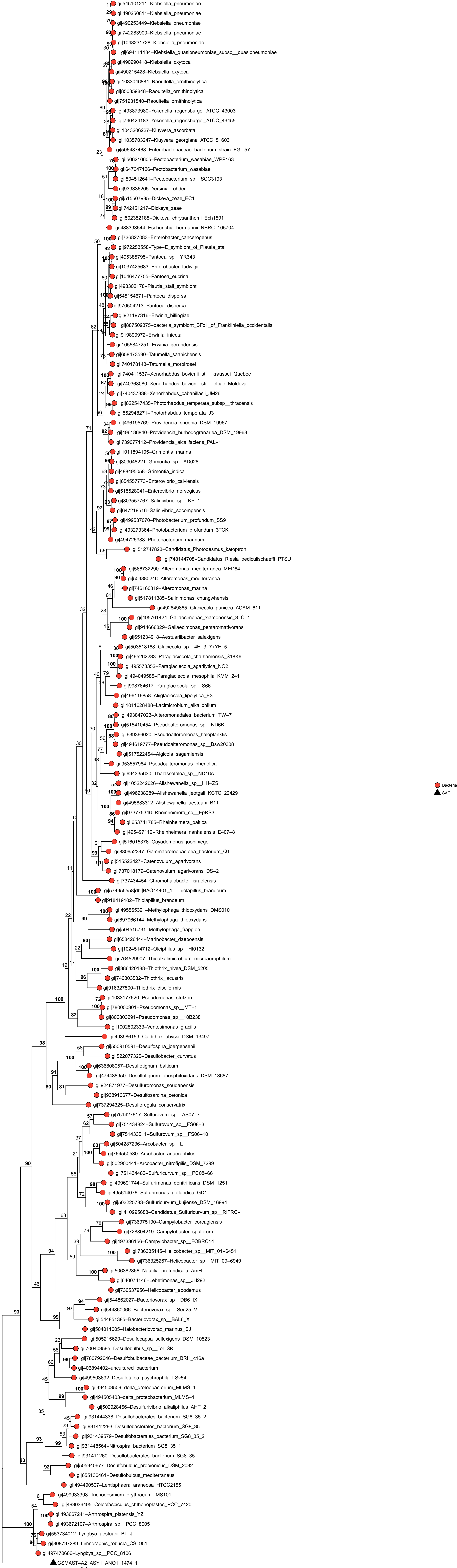


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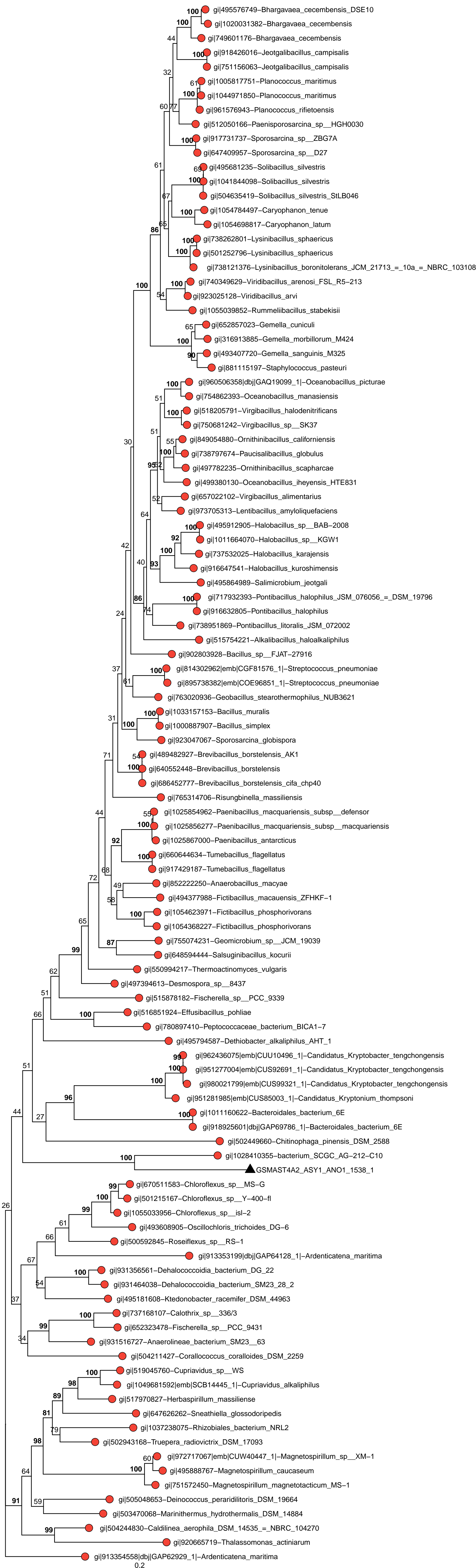


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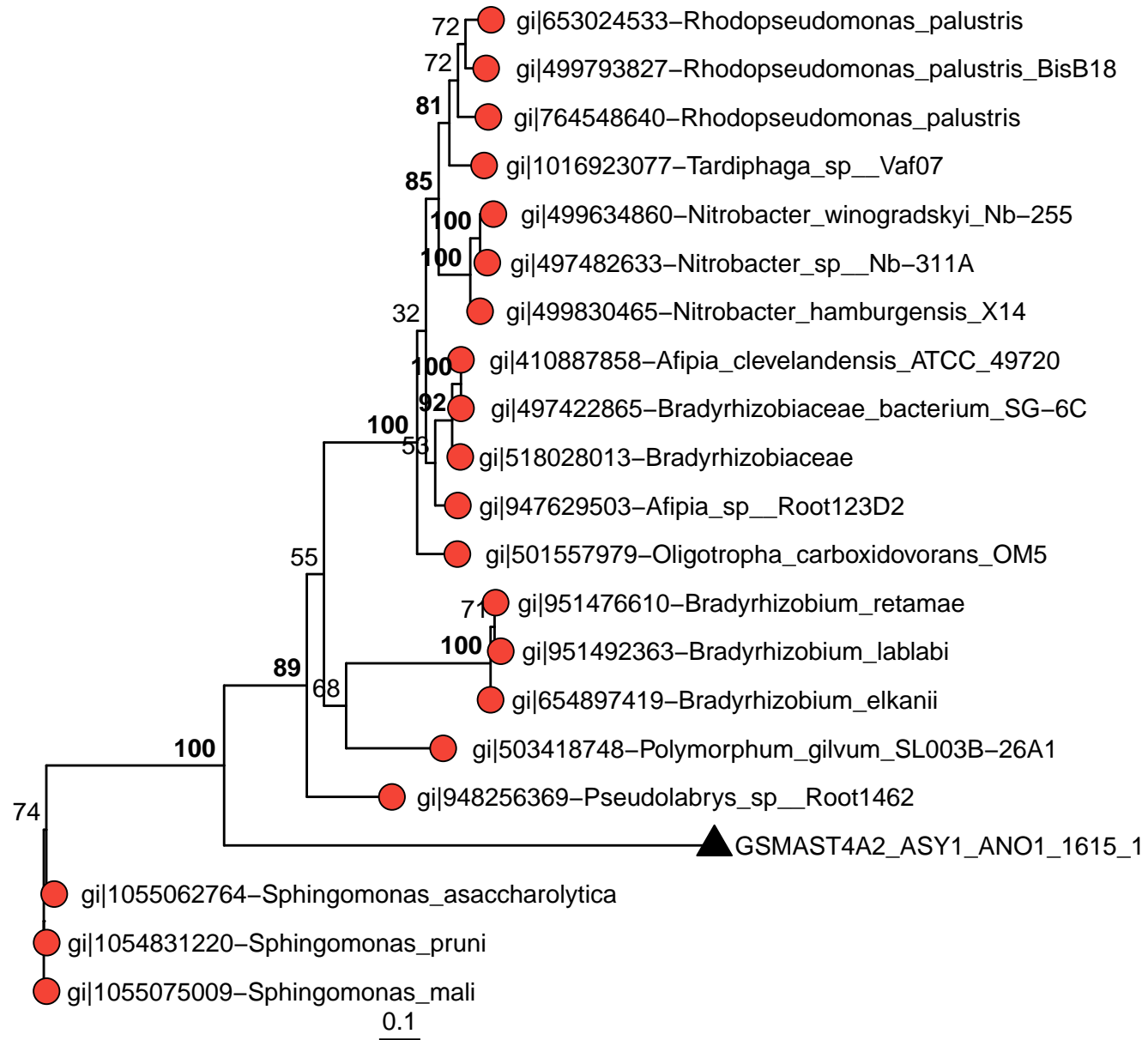


● Bacteria
 ▲ SAG

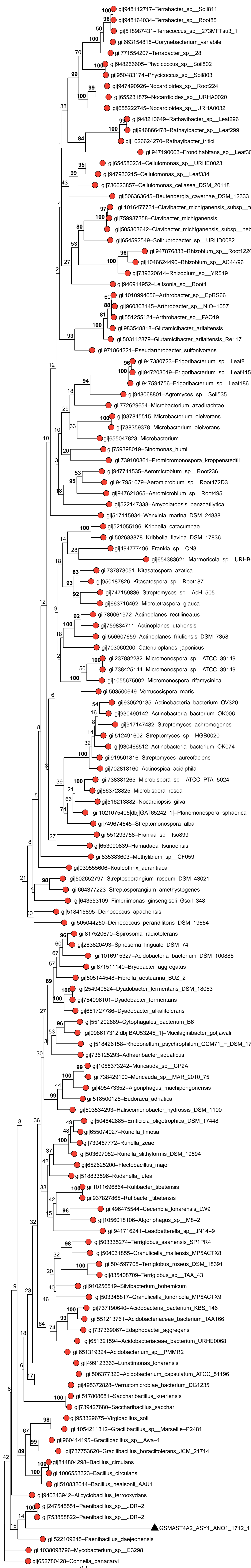


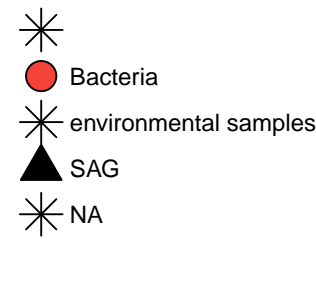
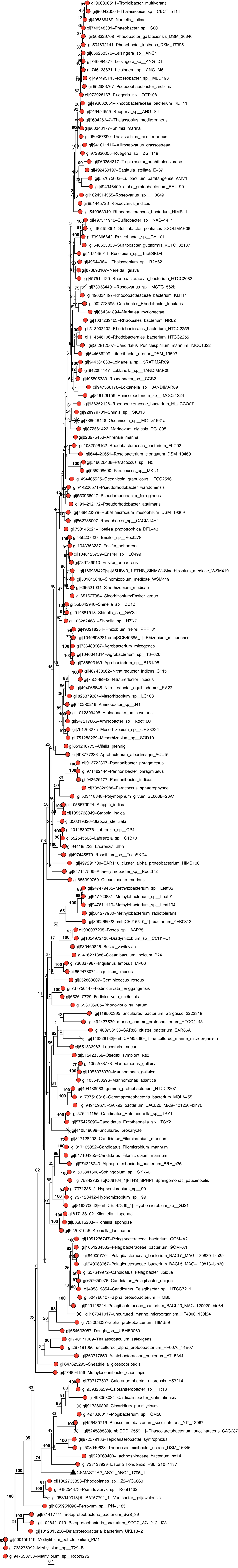
● Bacteria
▲ SAG

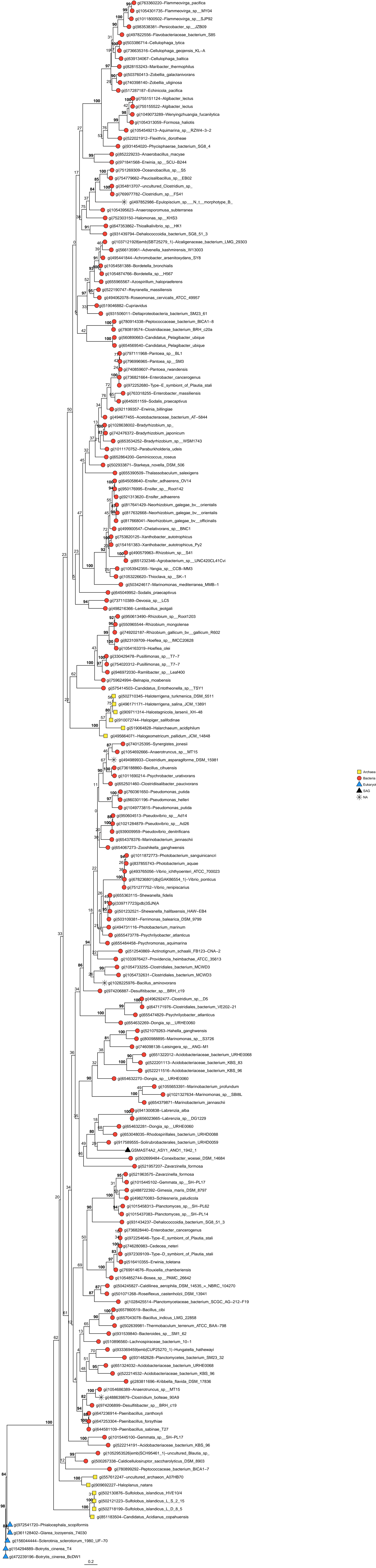
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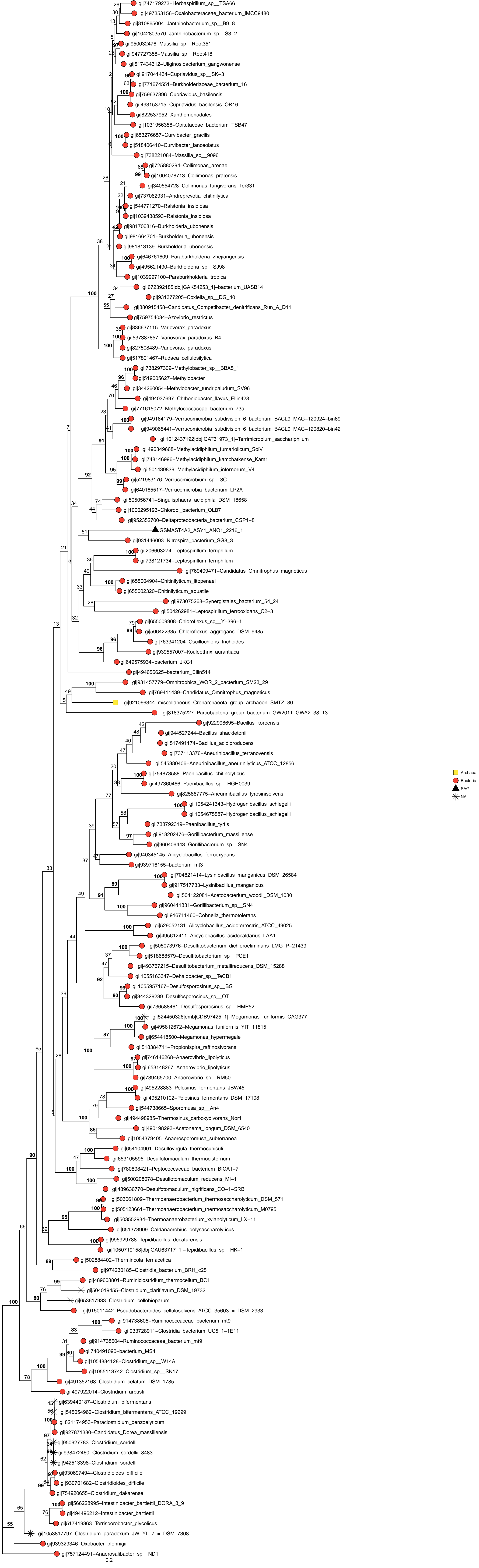


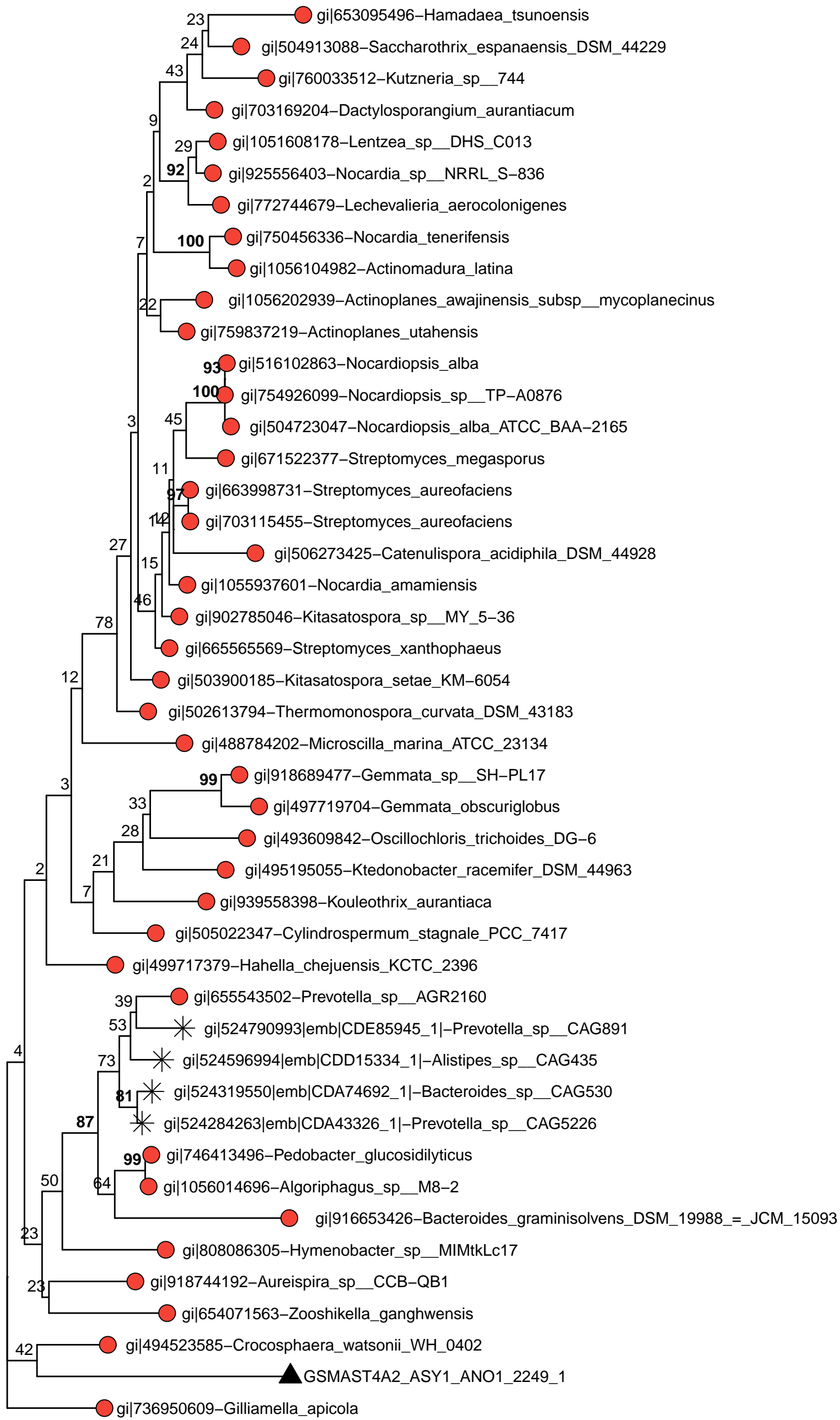
● Bacteria
▲ SAG





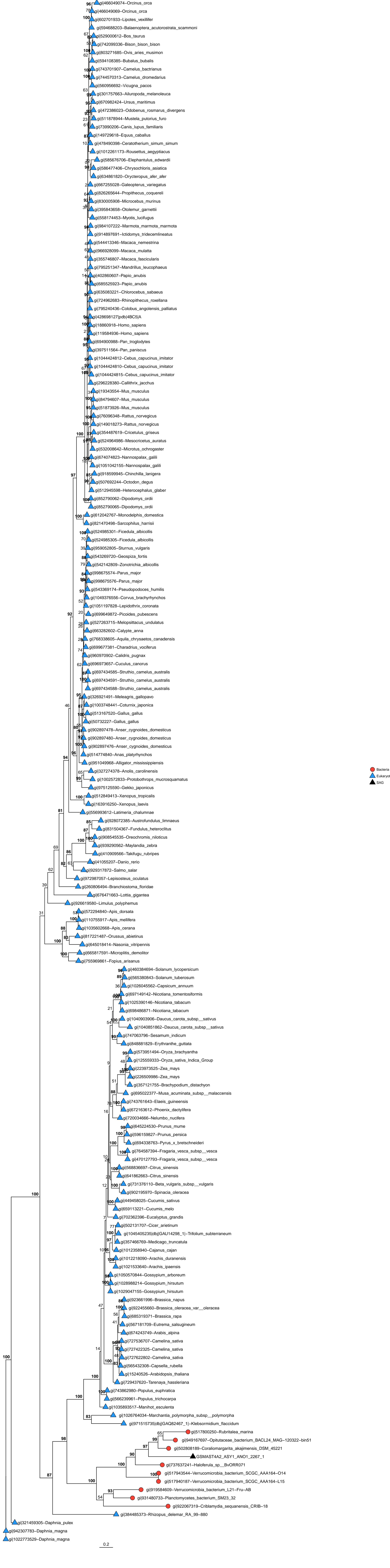




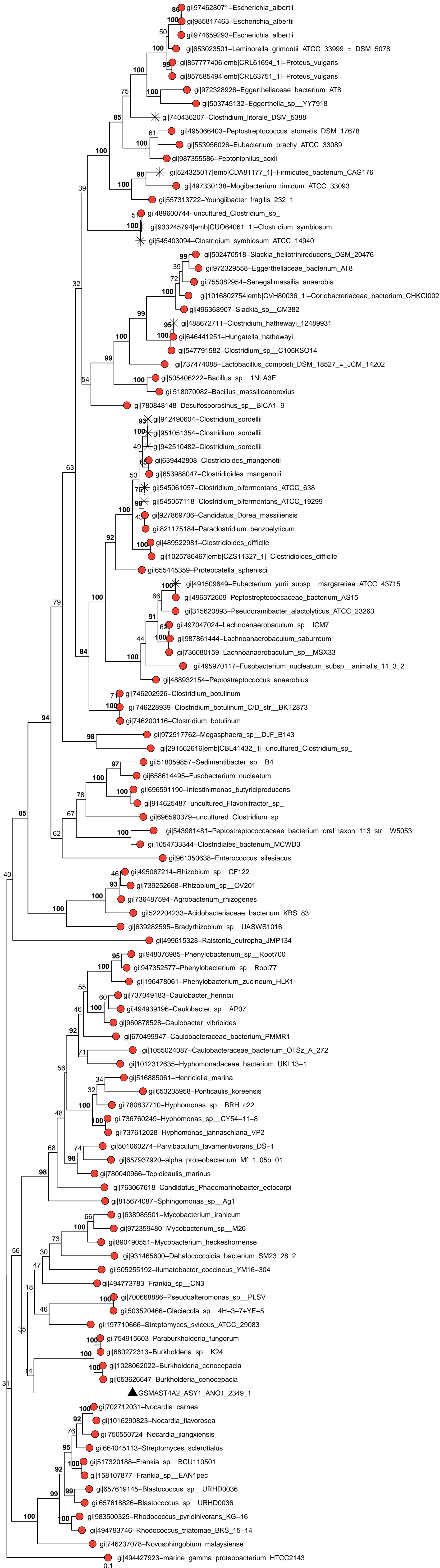


● Bacteria
▲ SAG
✱ NA

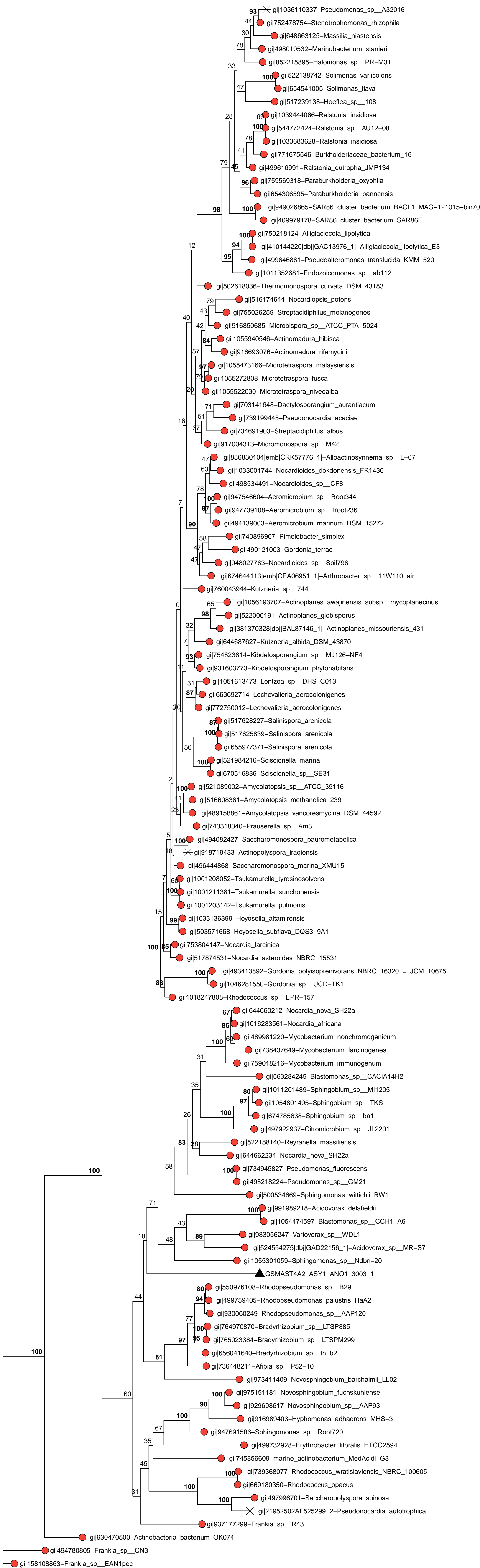
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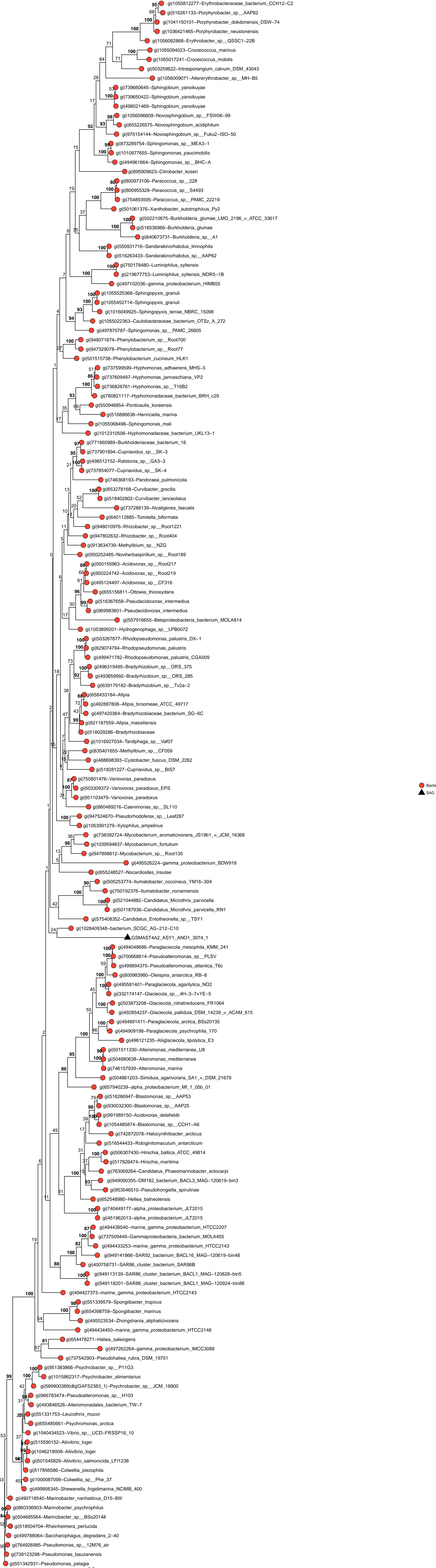


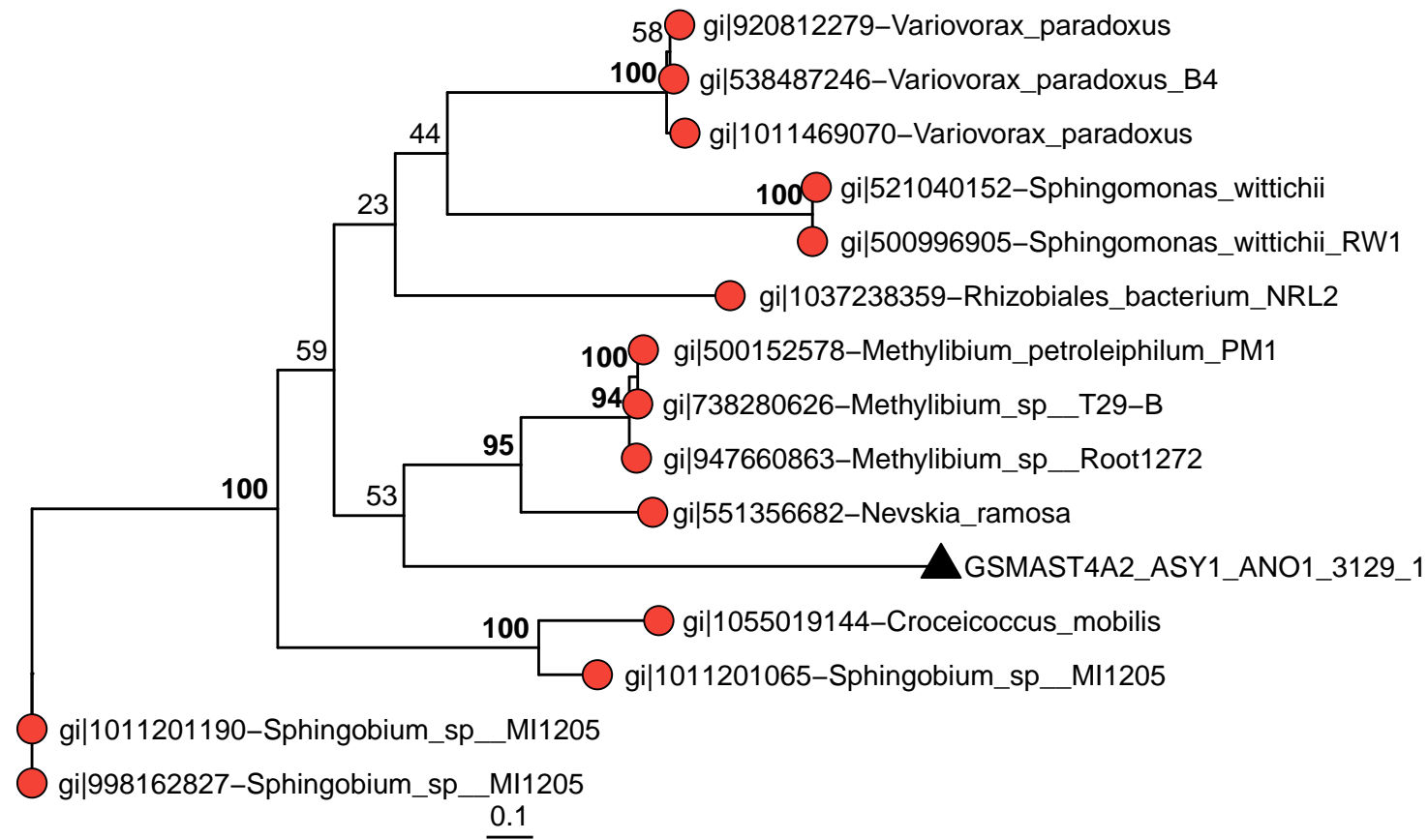


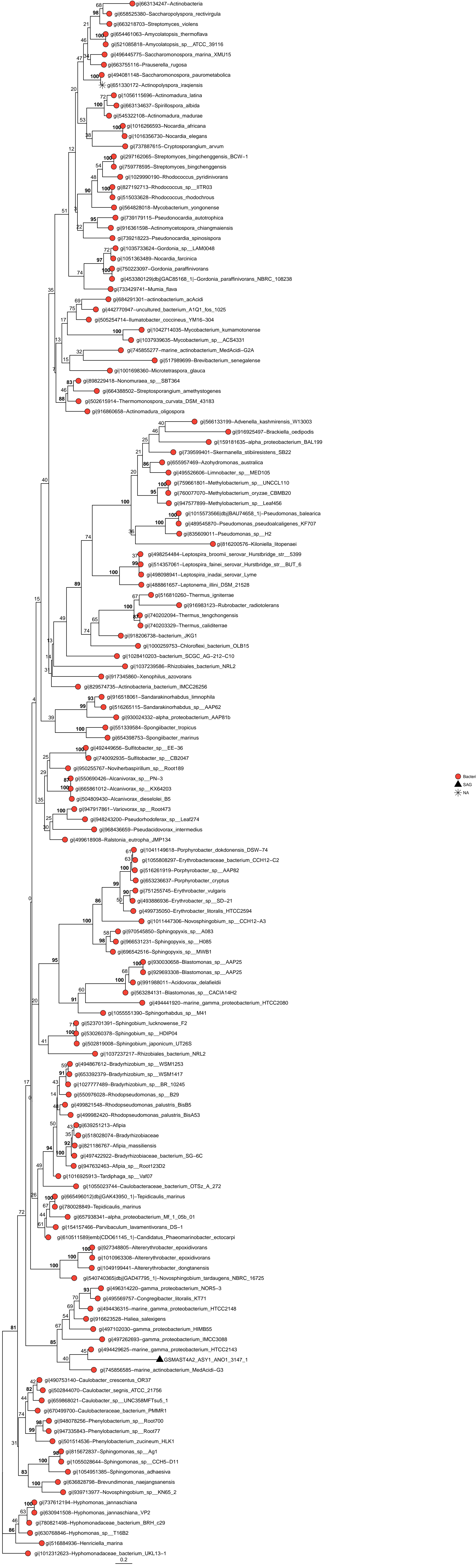


● Bacteria
▲ SAG
✱ NA

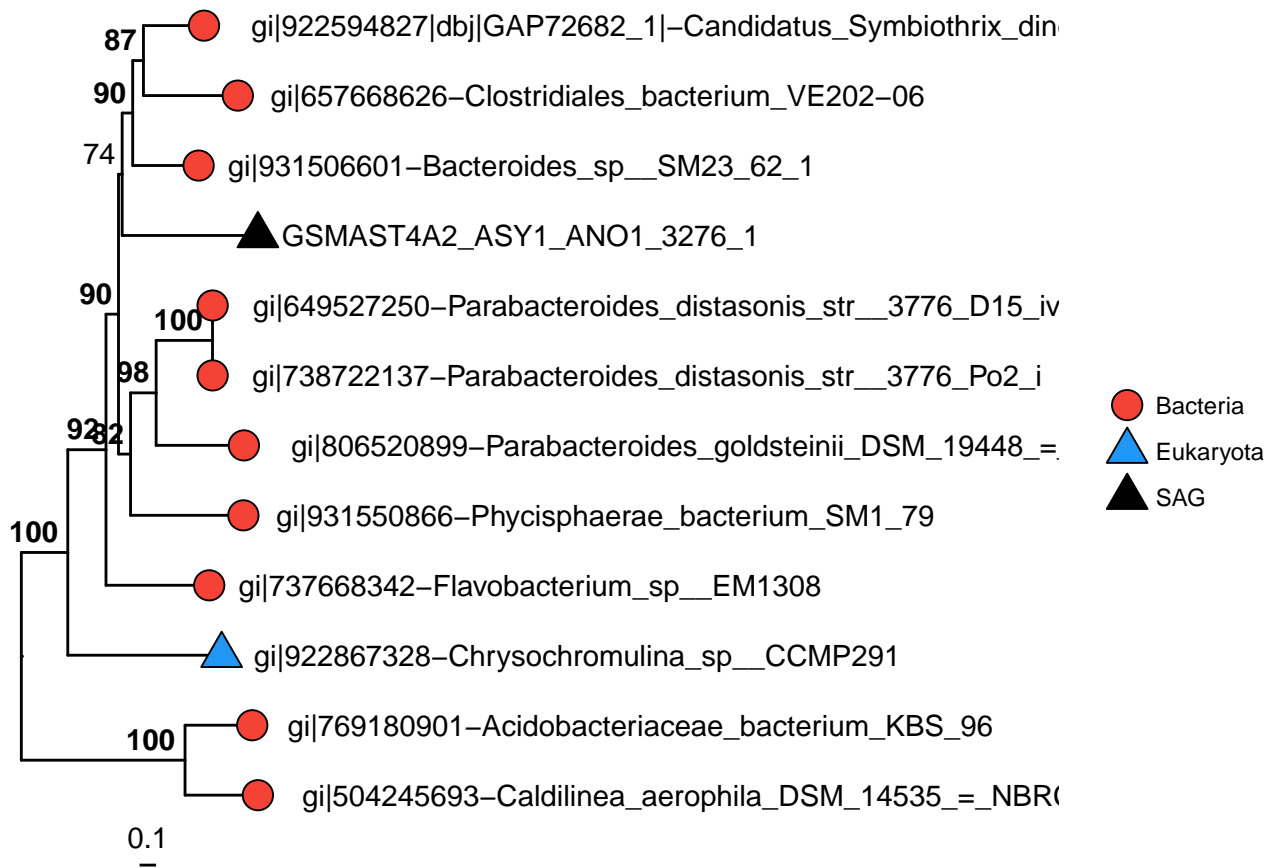
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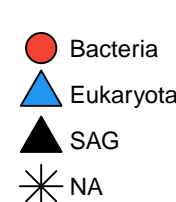
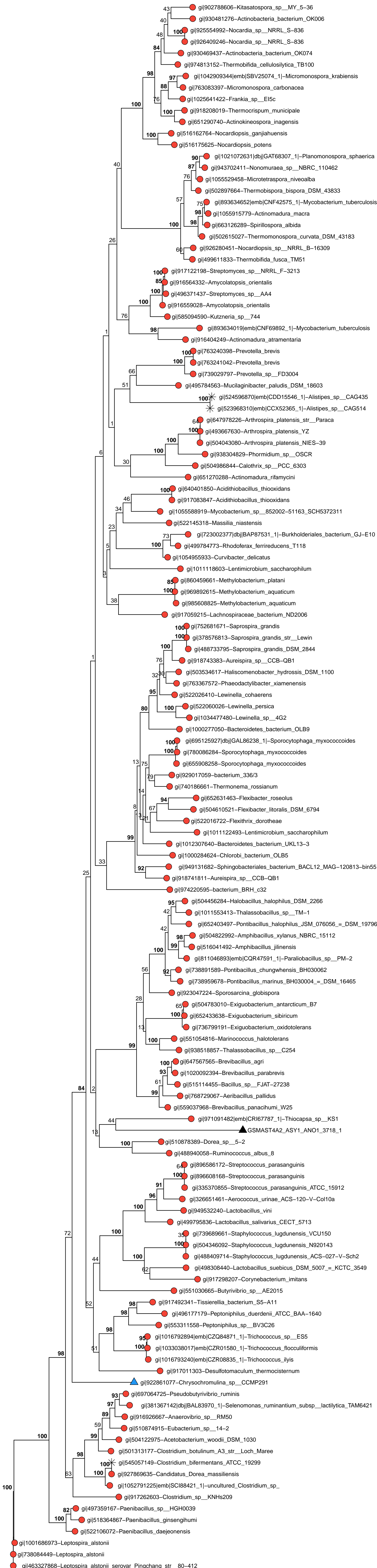


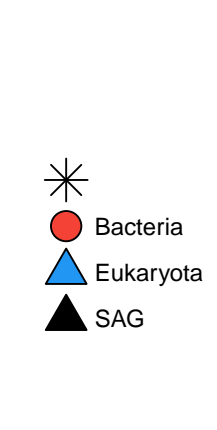
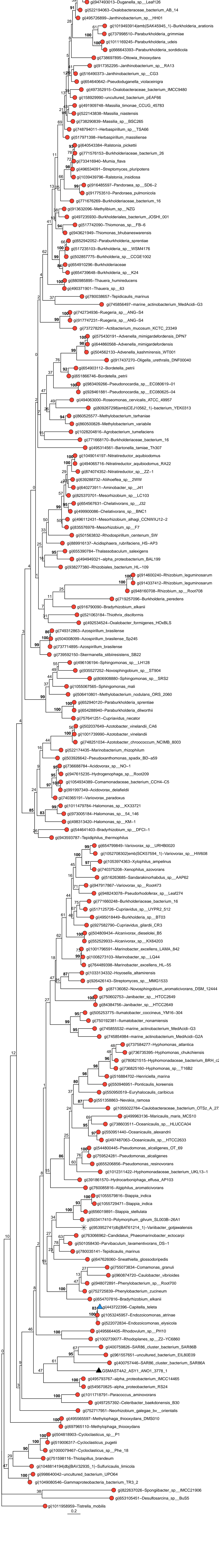




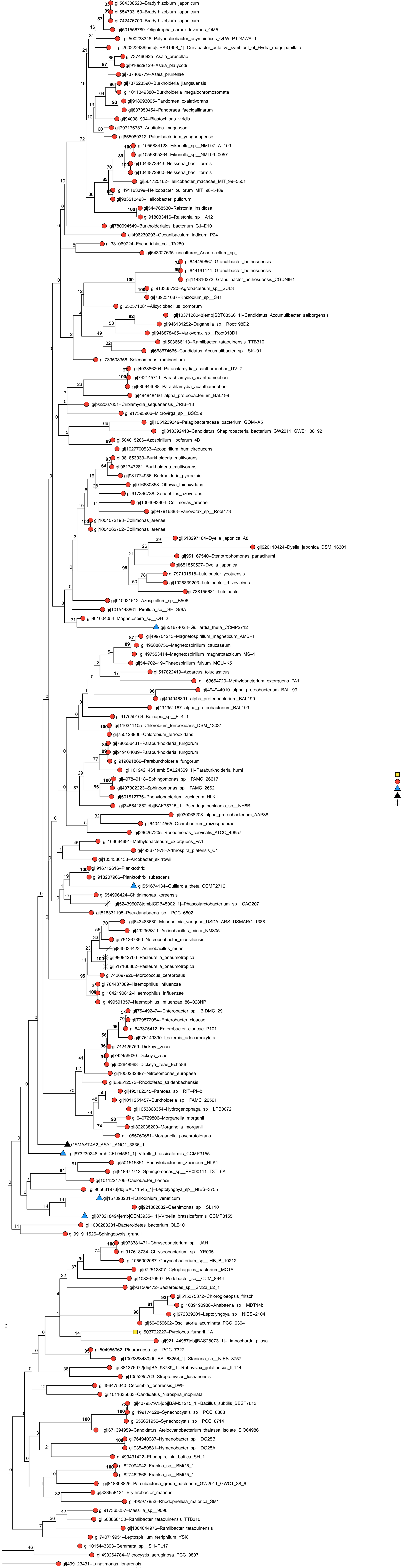
● Bacteria
▲ SAG
✱ NA

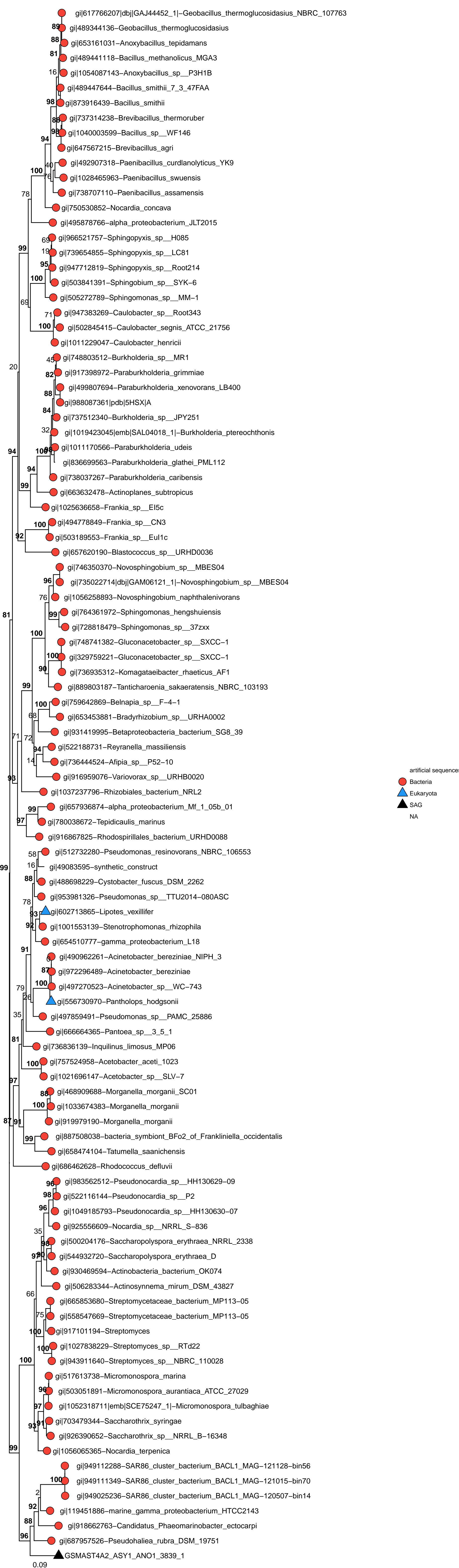


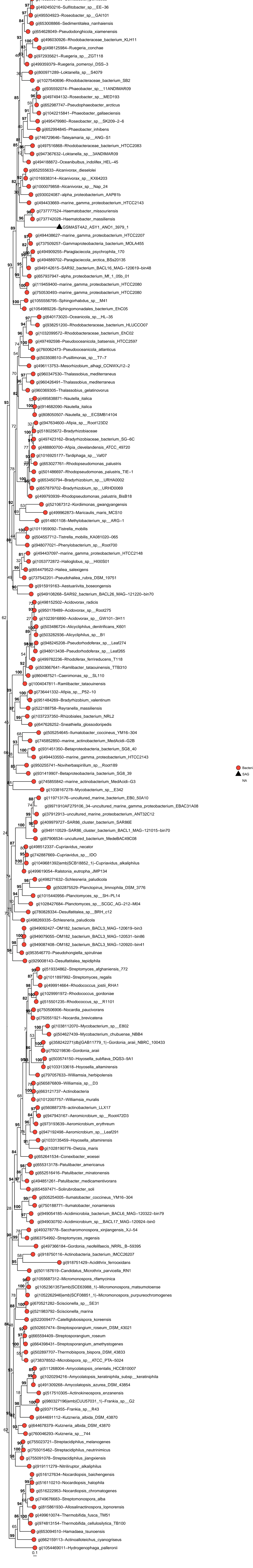


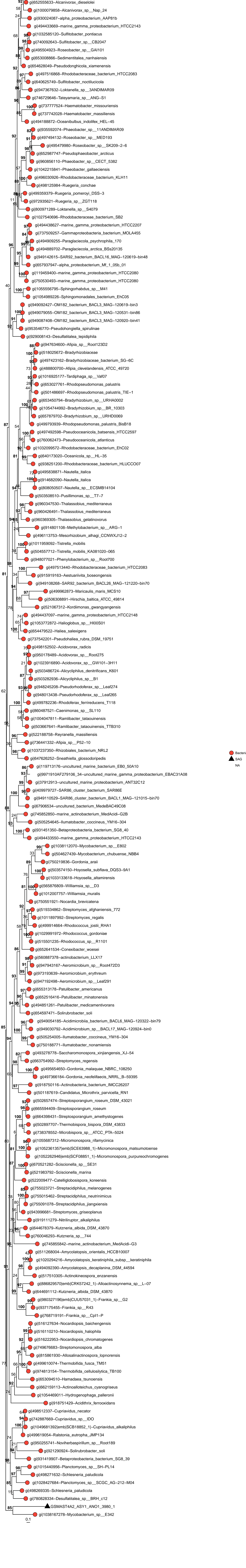


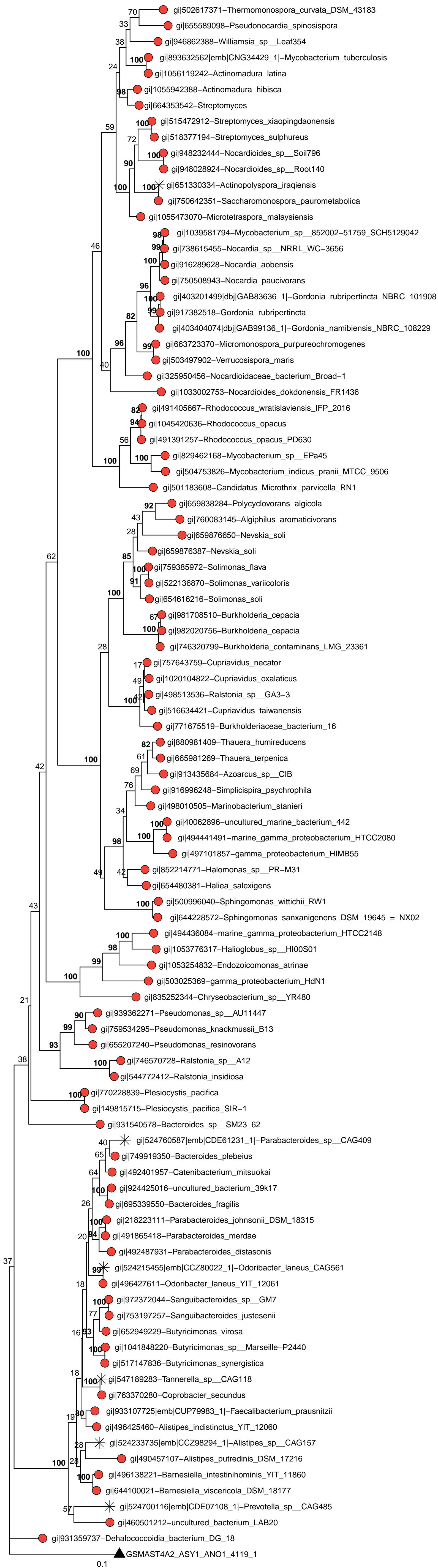
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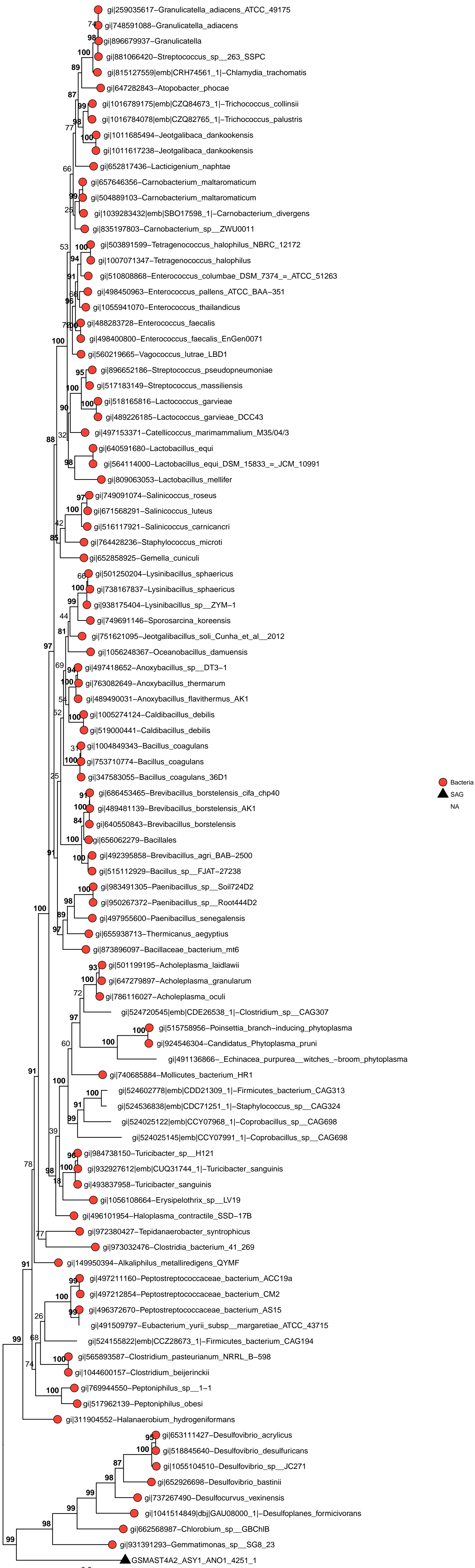


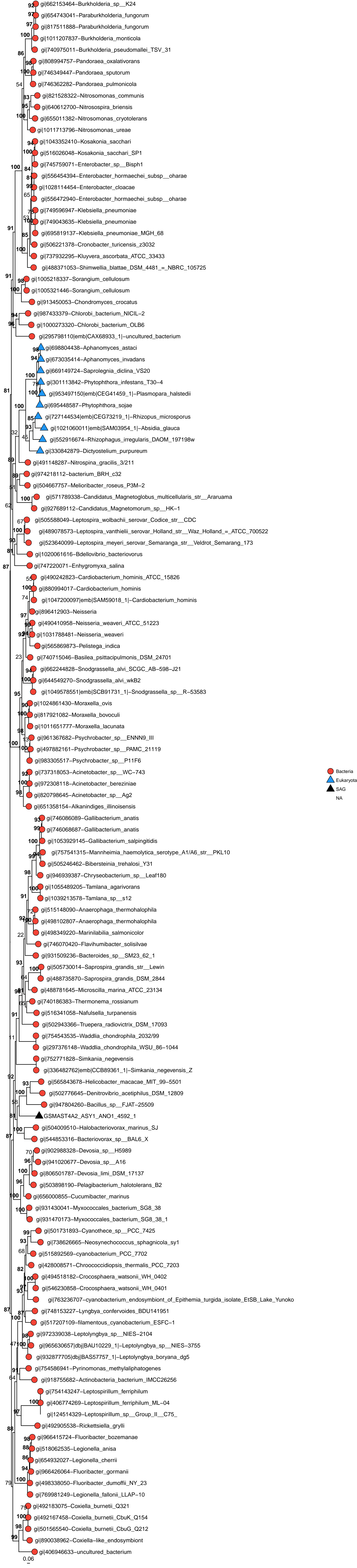


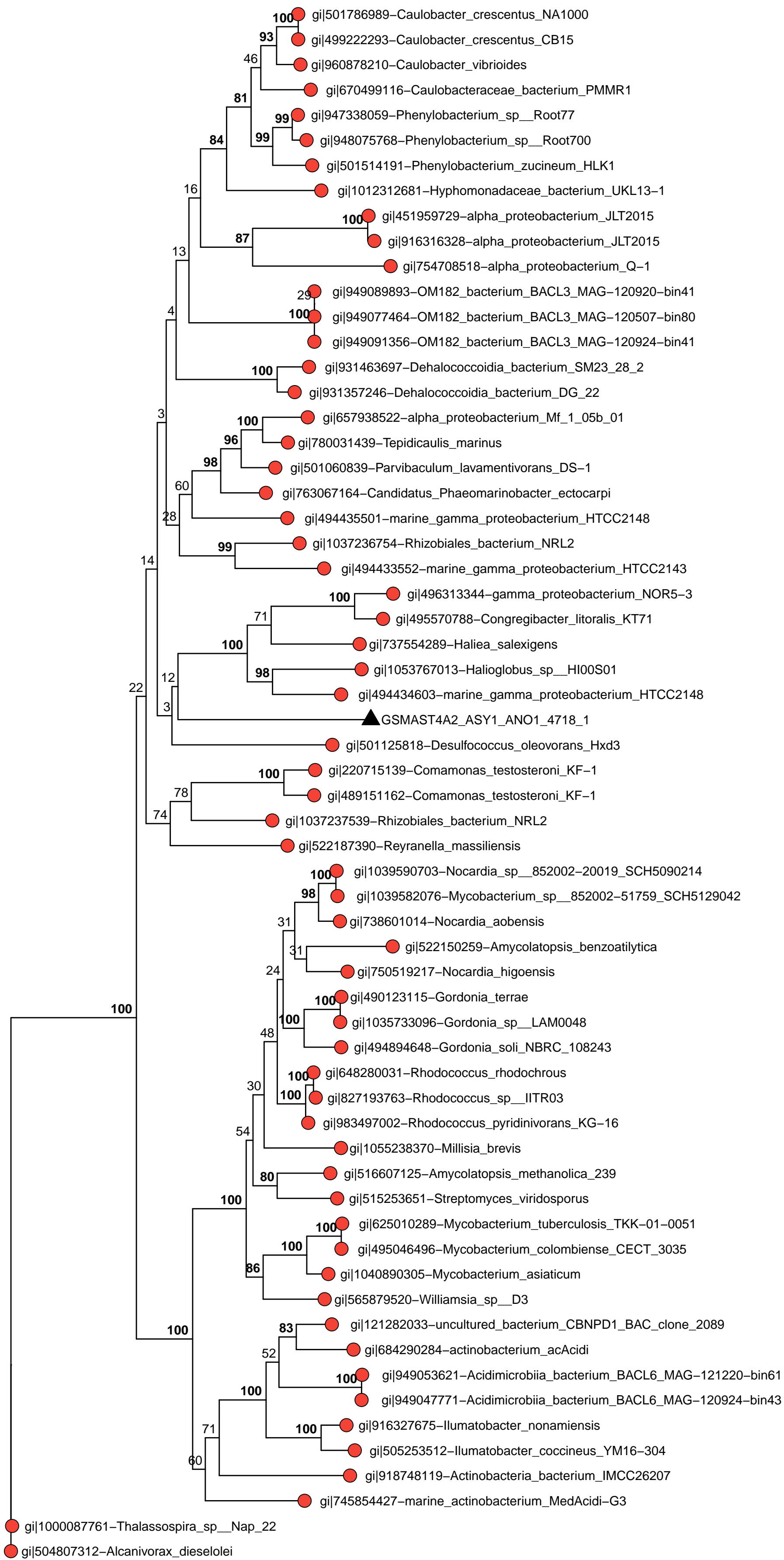


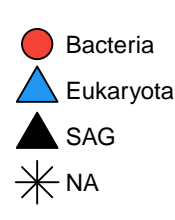
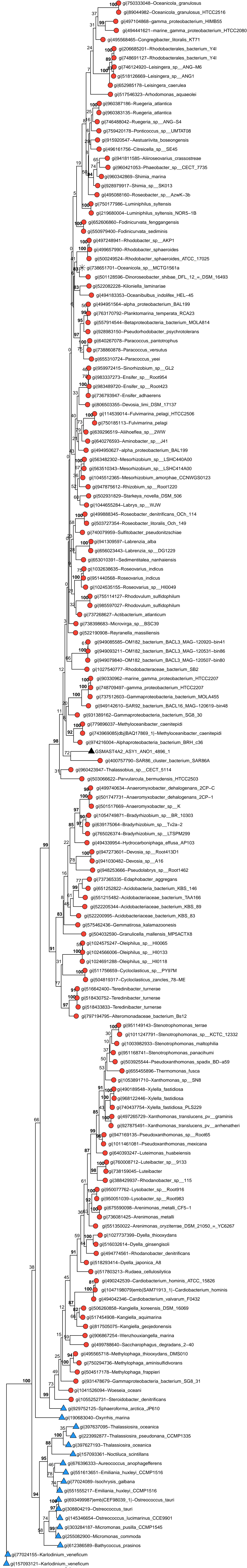
- Bacteria
- ▲ SAG
- ✱ NA

0.1

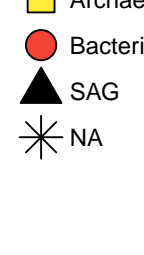
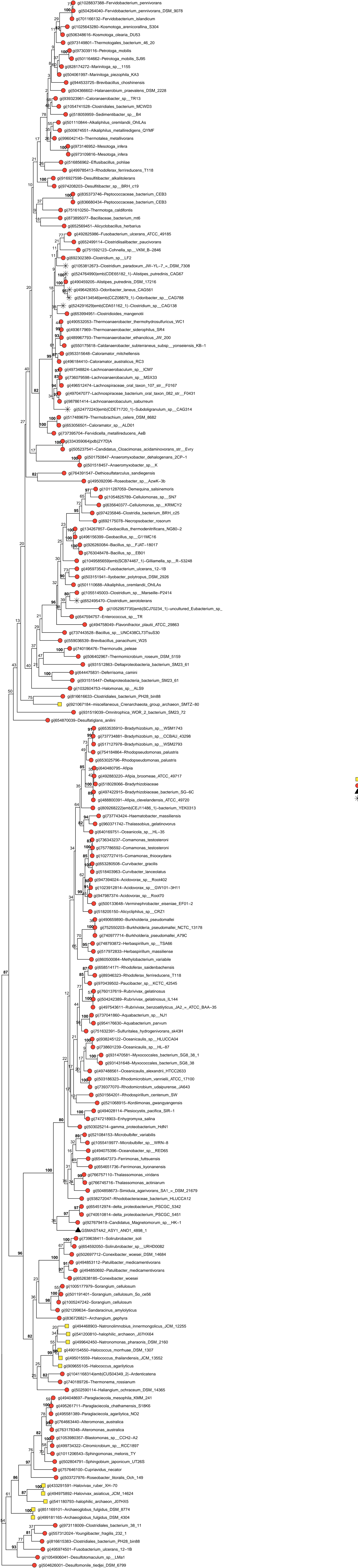




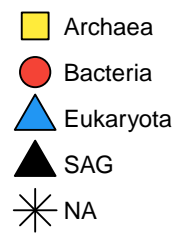
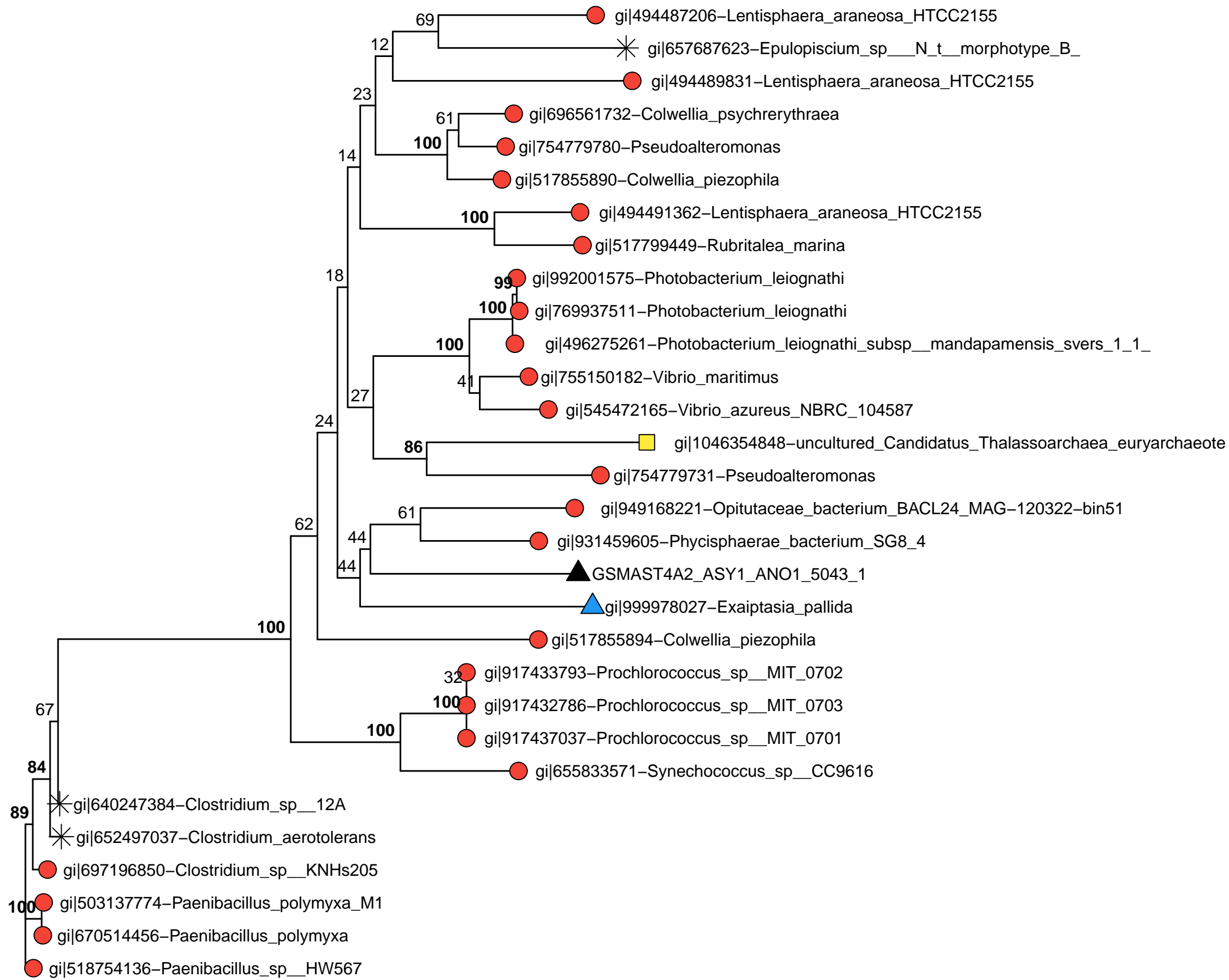


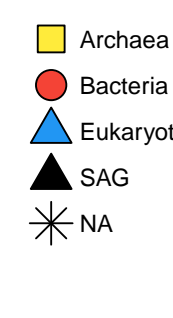
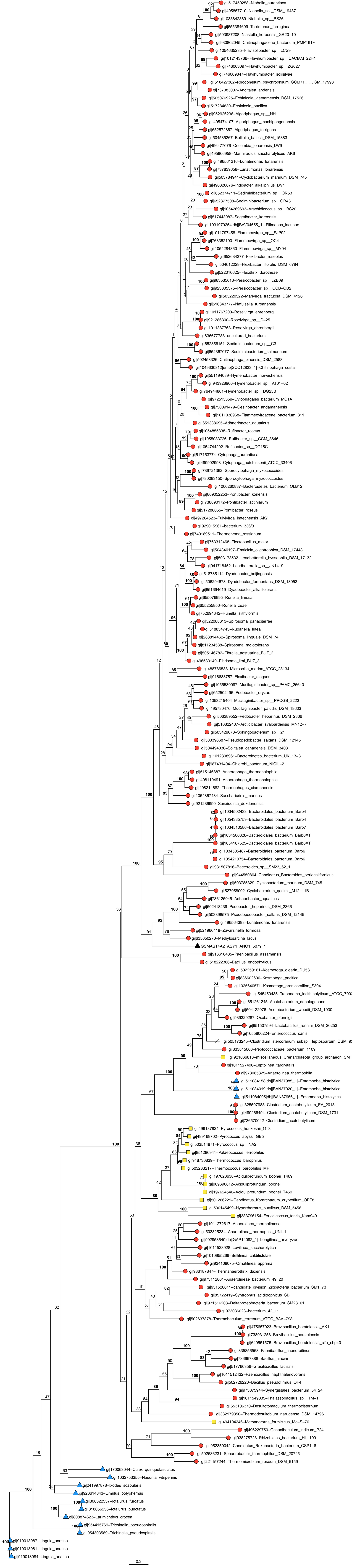


● Bacteria
 ▲ Eukaryota
 ▲ SAG
 ✱ NA

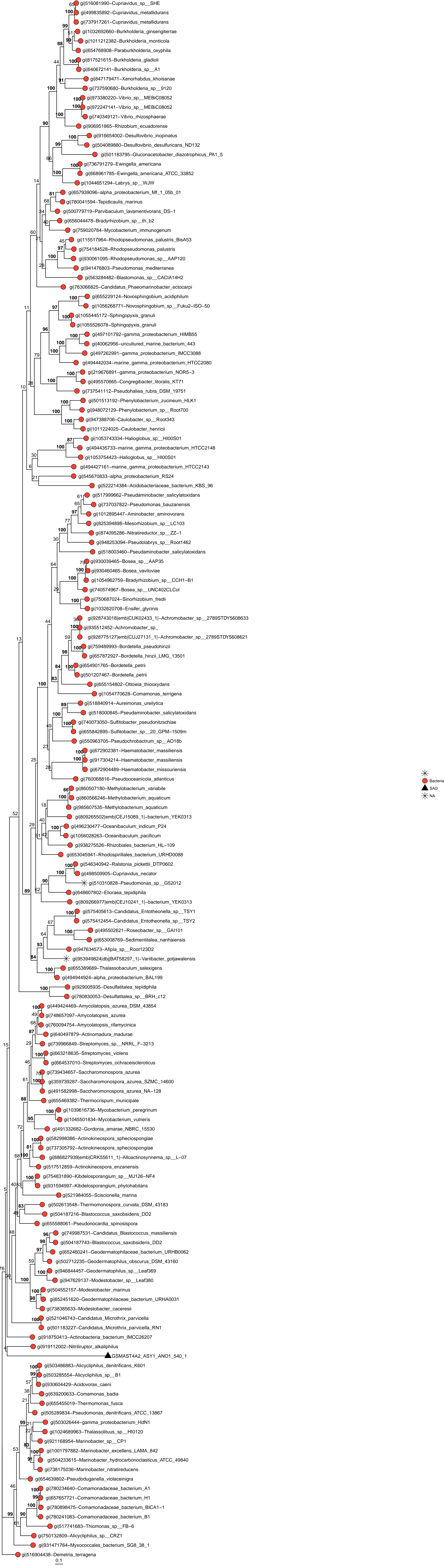


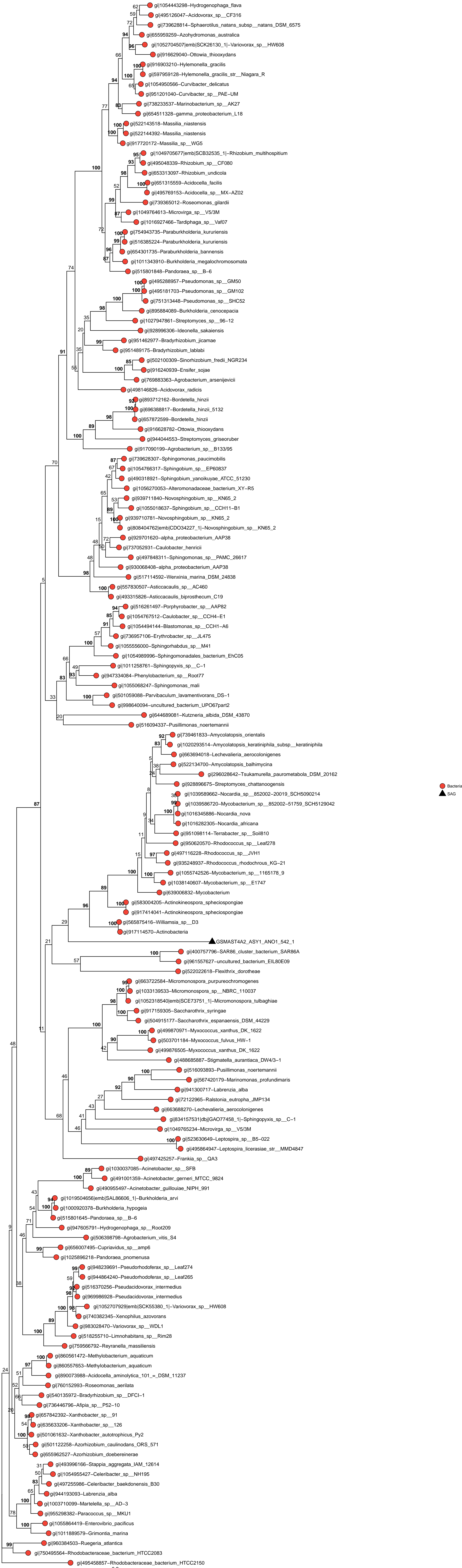
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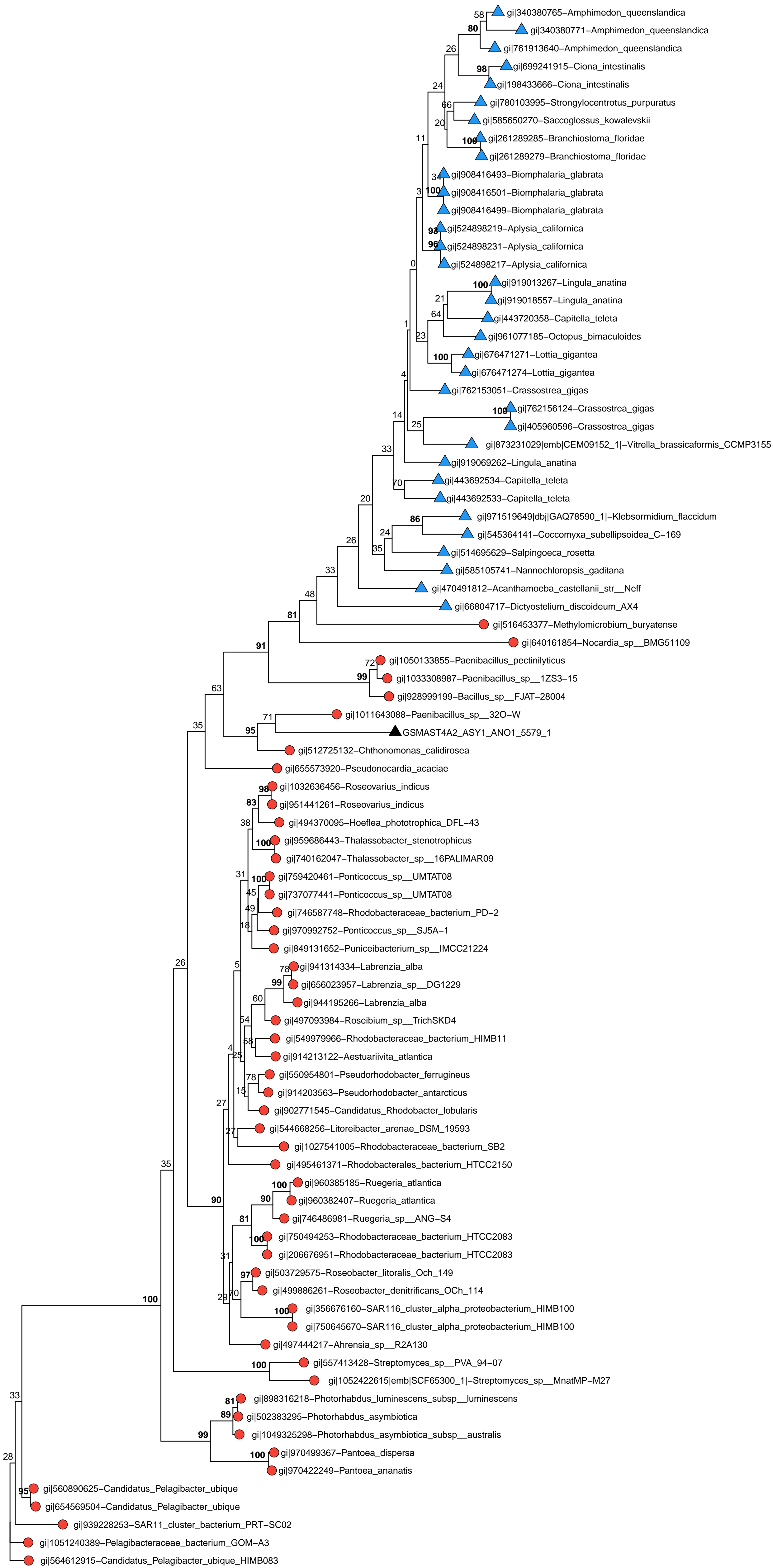




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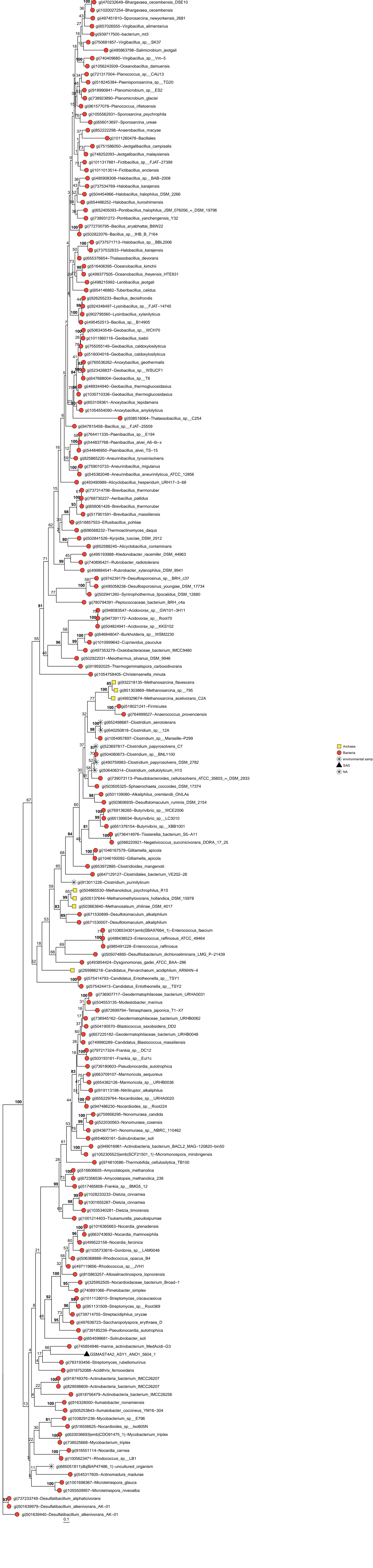


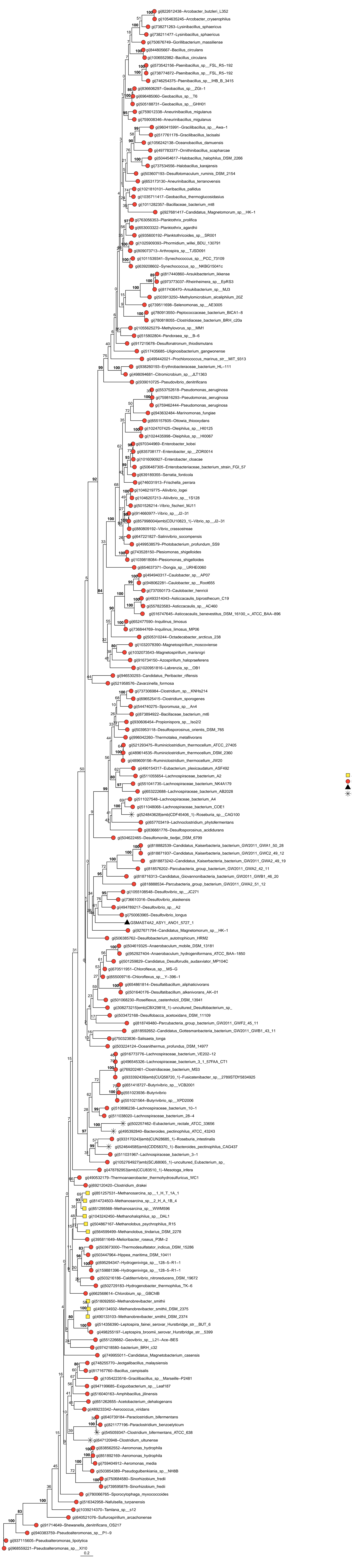


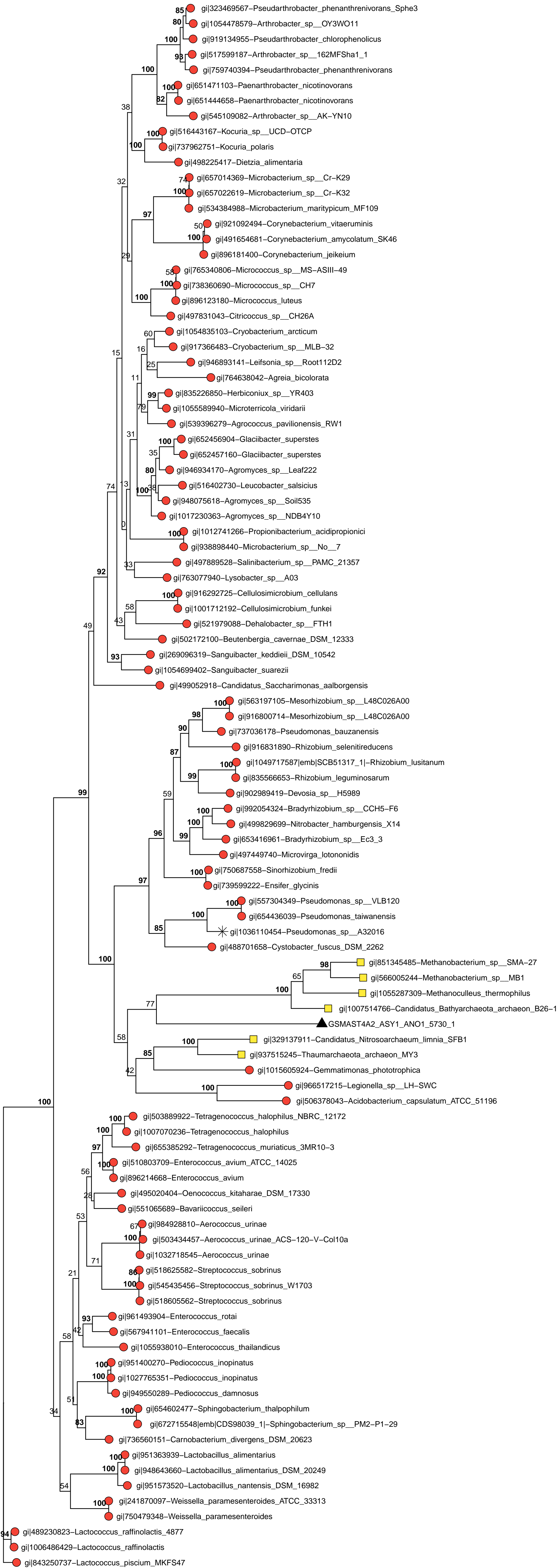


● Bacteria
 ▲ Eukaryota
 ▲ SAG

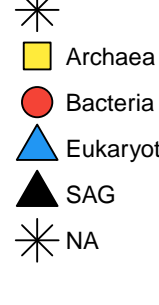
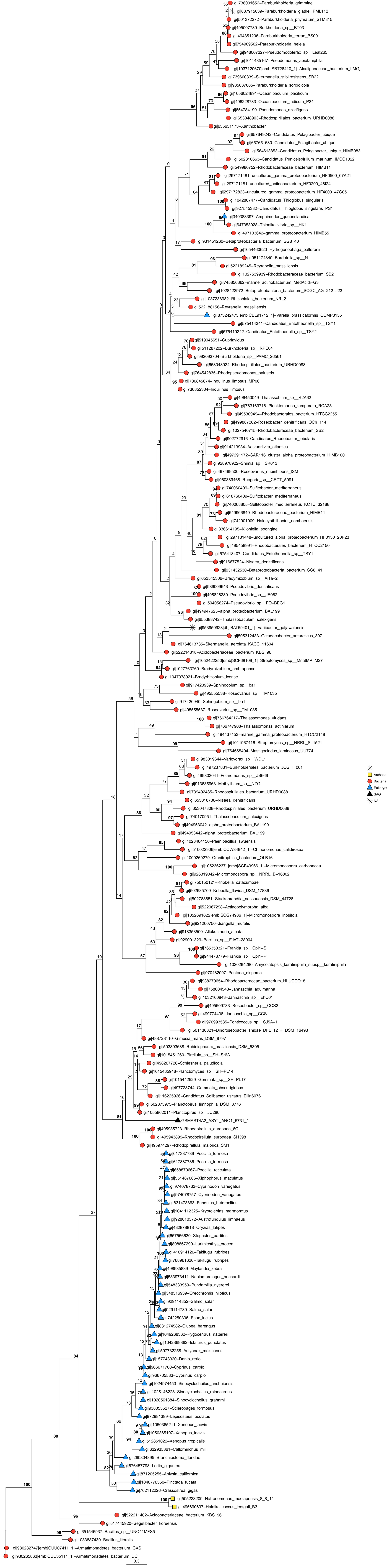
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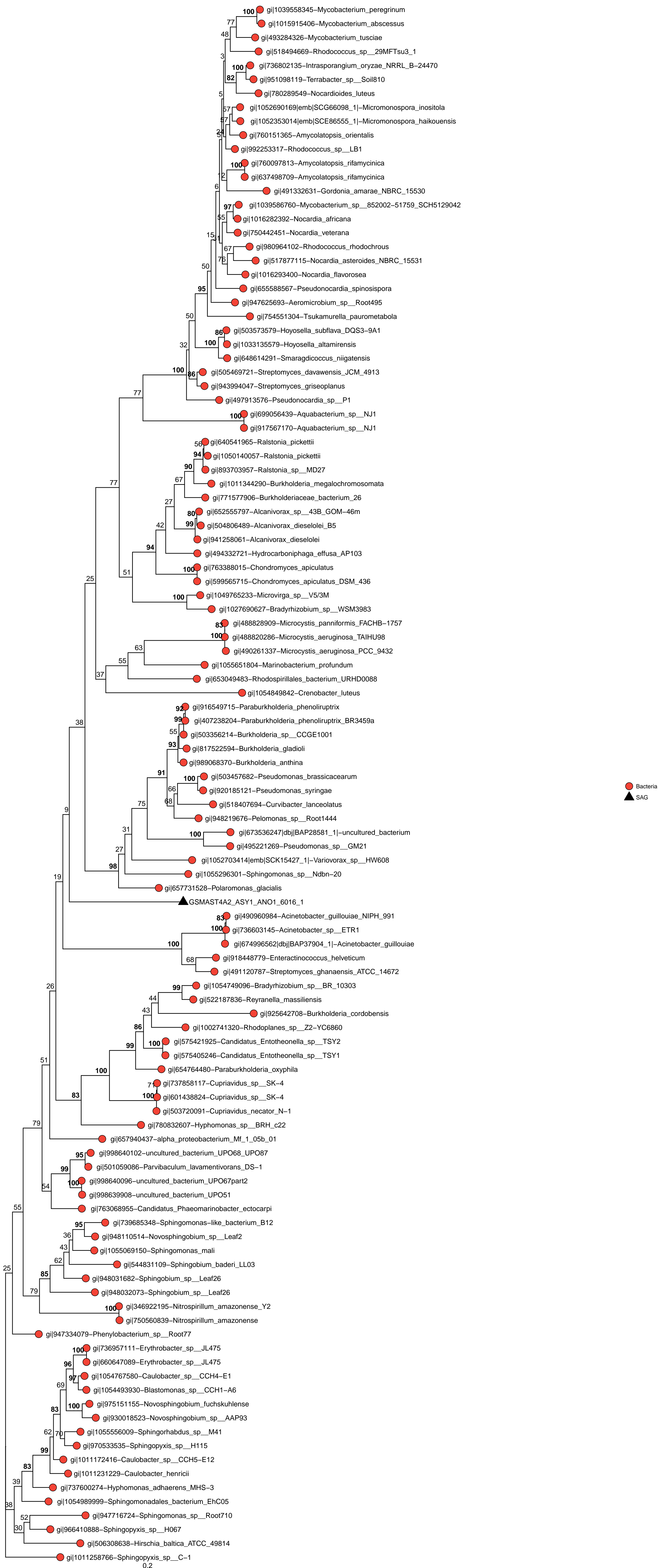


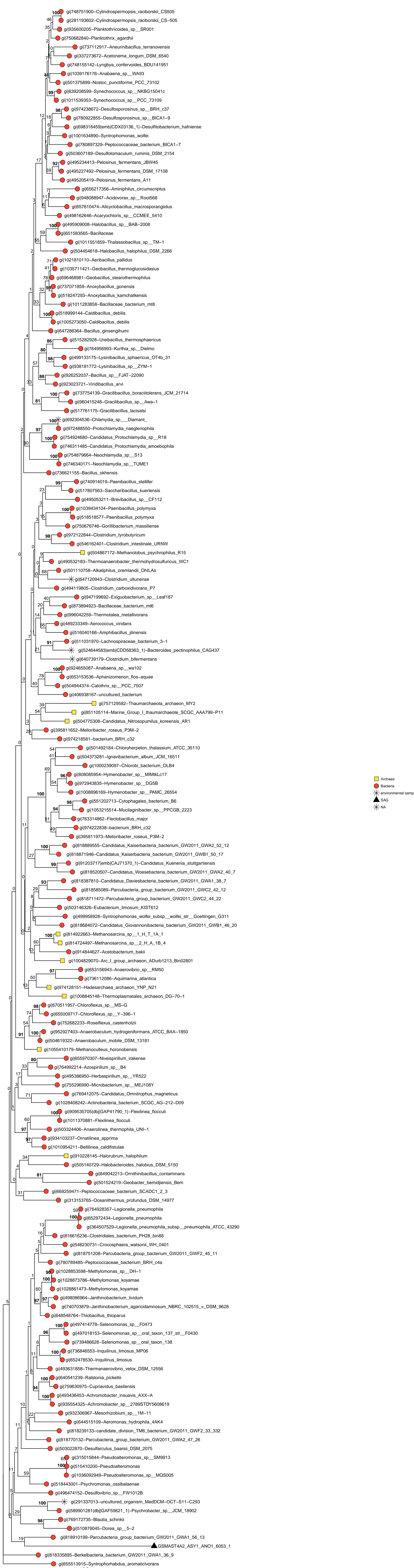


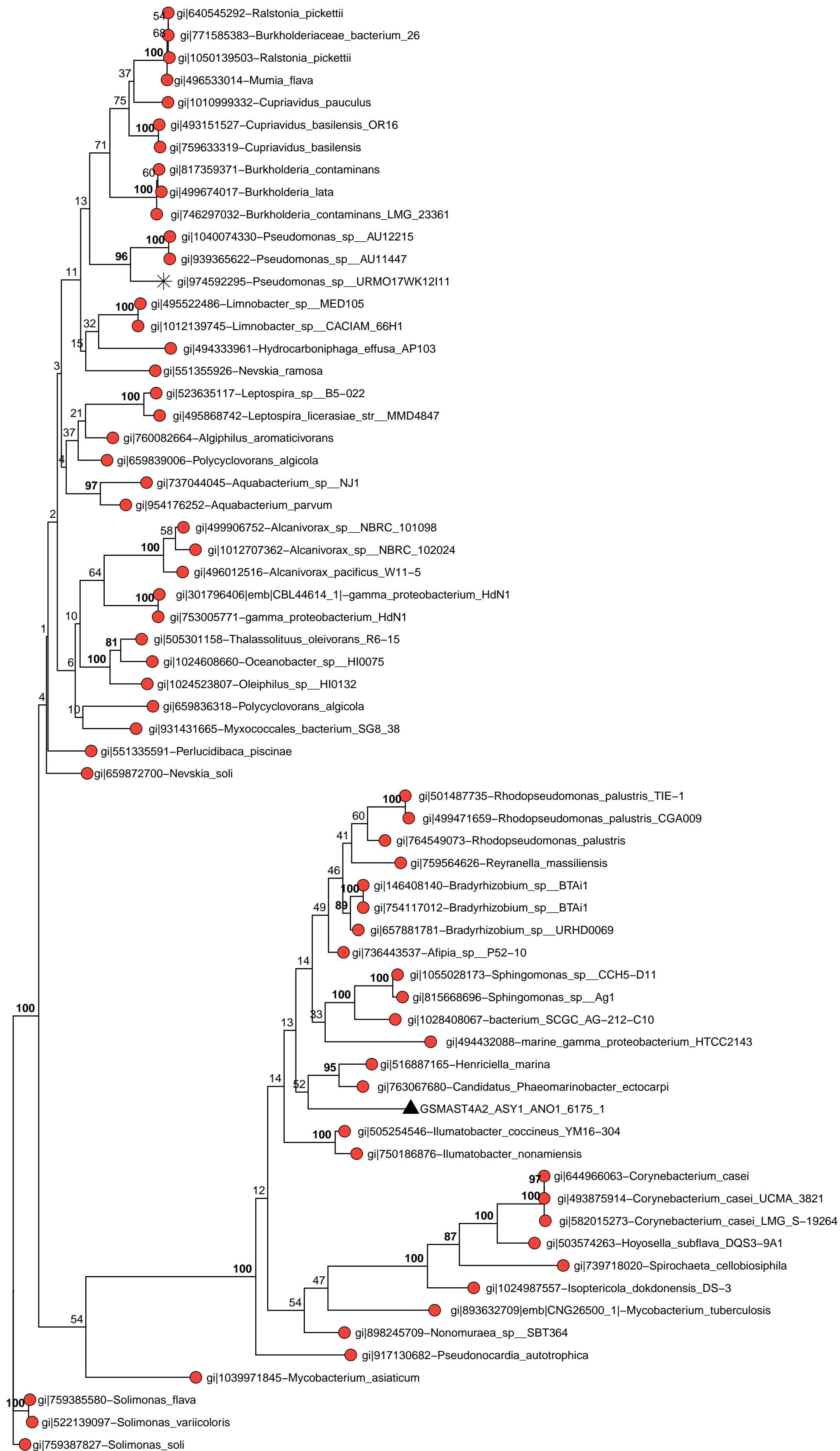
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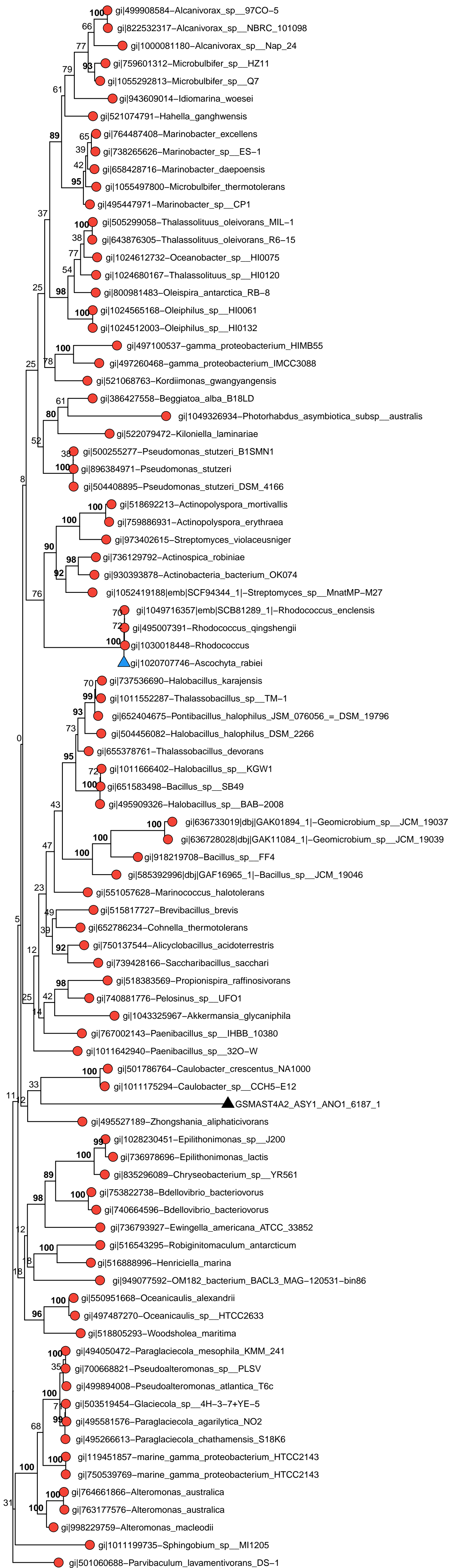




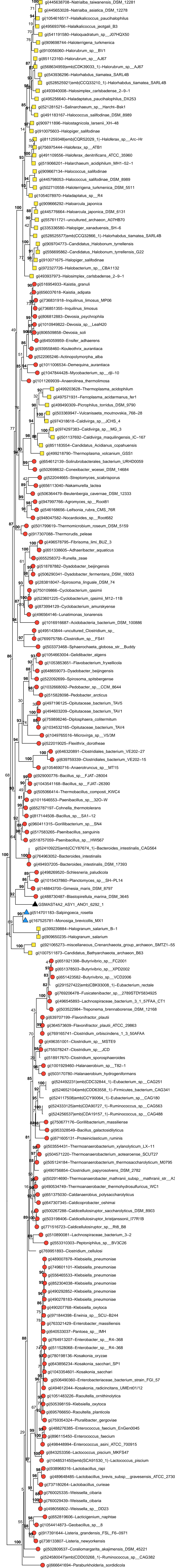


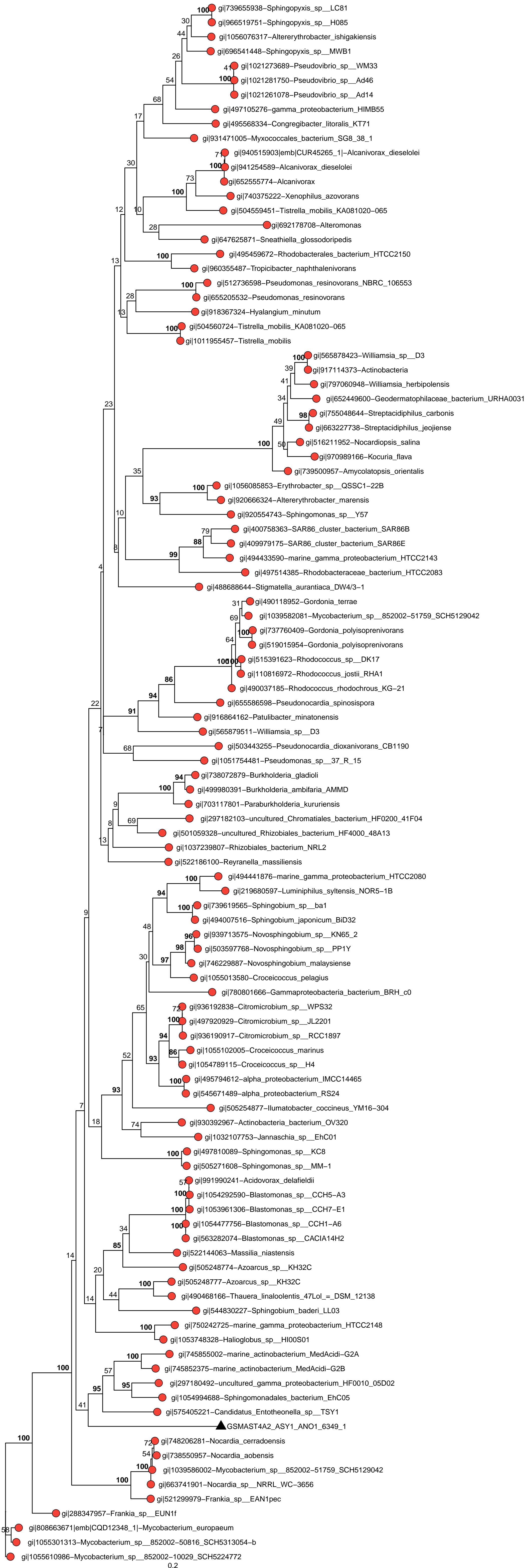
● Bacteria
▲ SAG
✱ NA

0.2

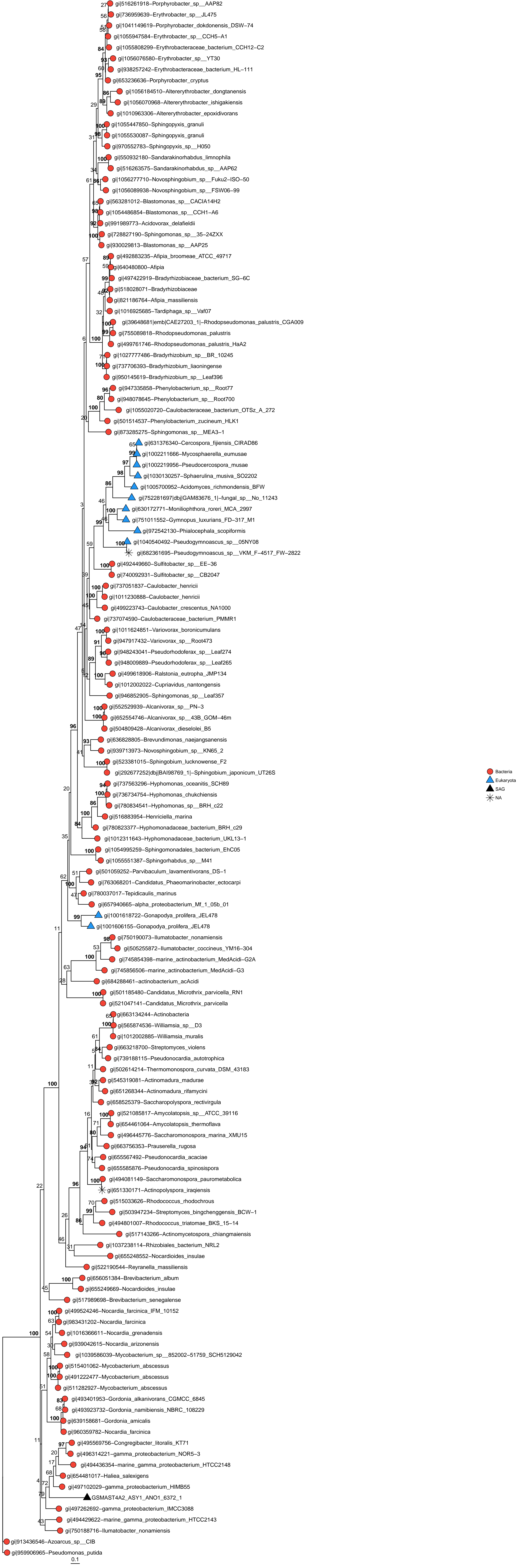


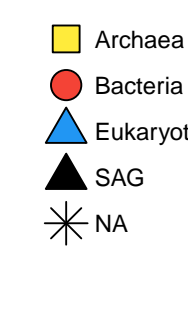
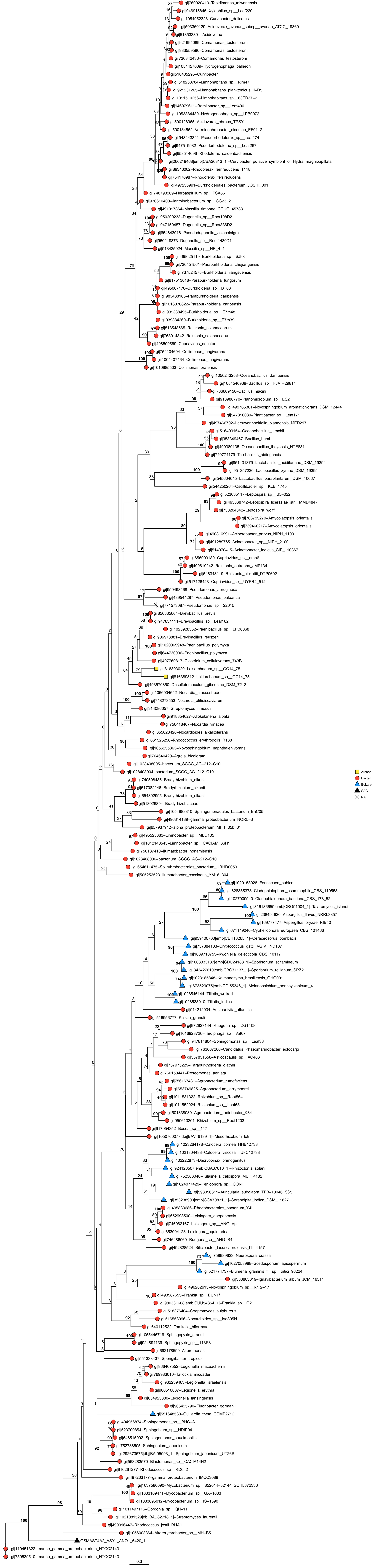
● Bacteria
▲ Eukaryota
▲ SAG



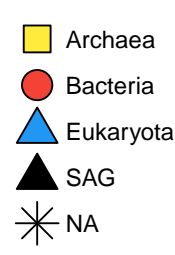
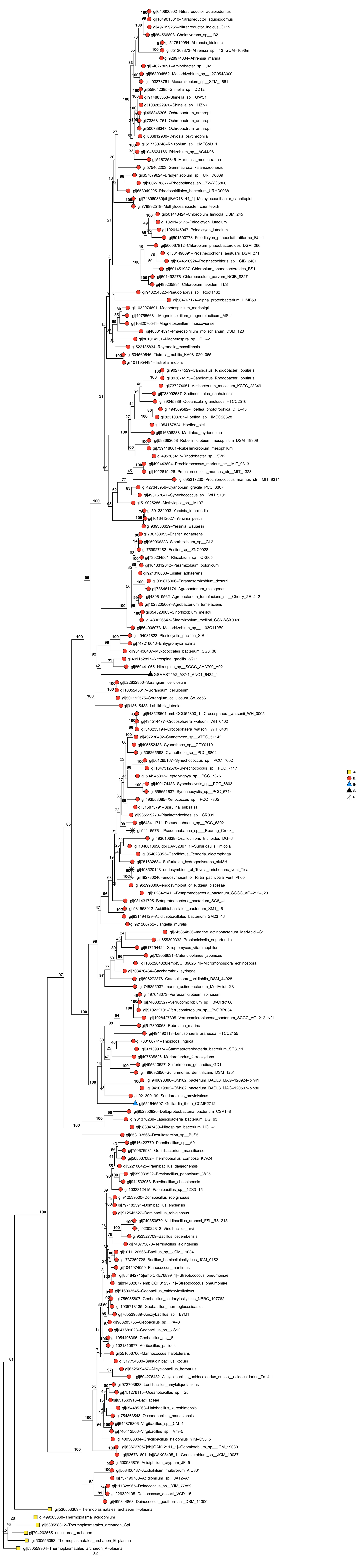


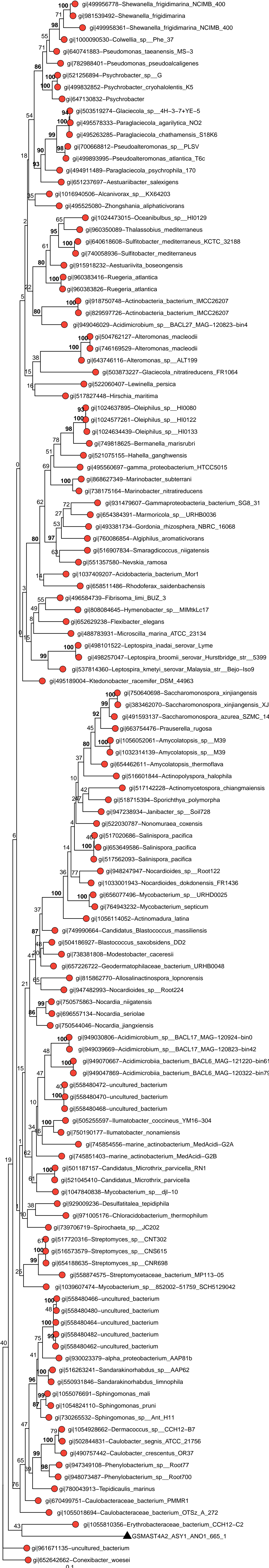
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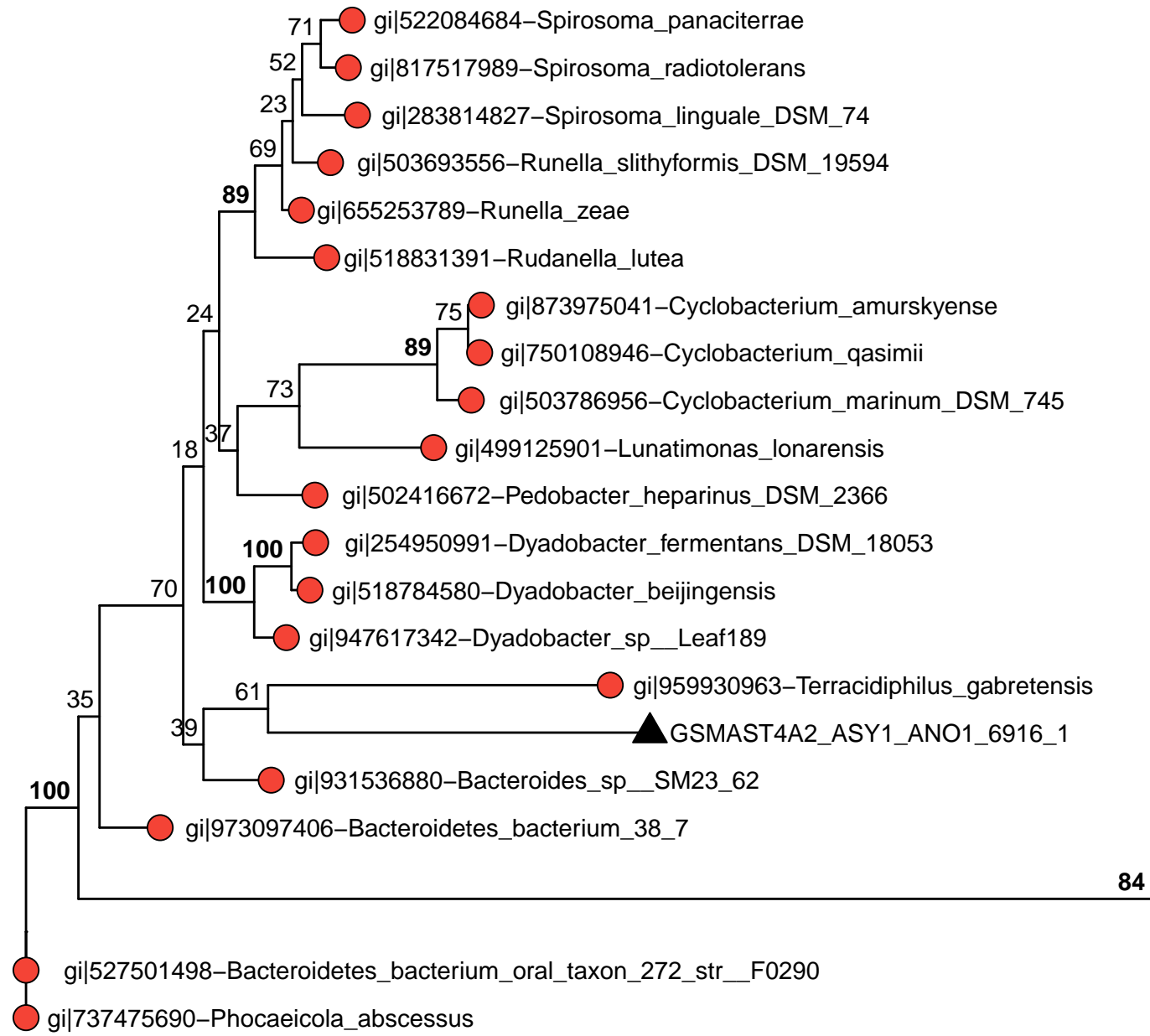
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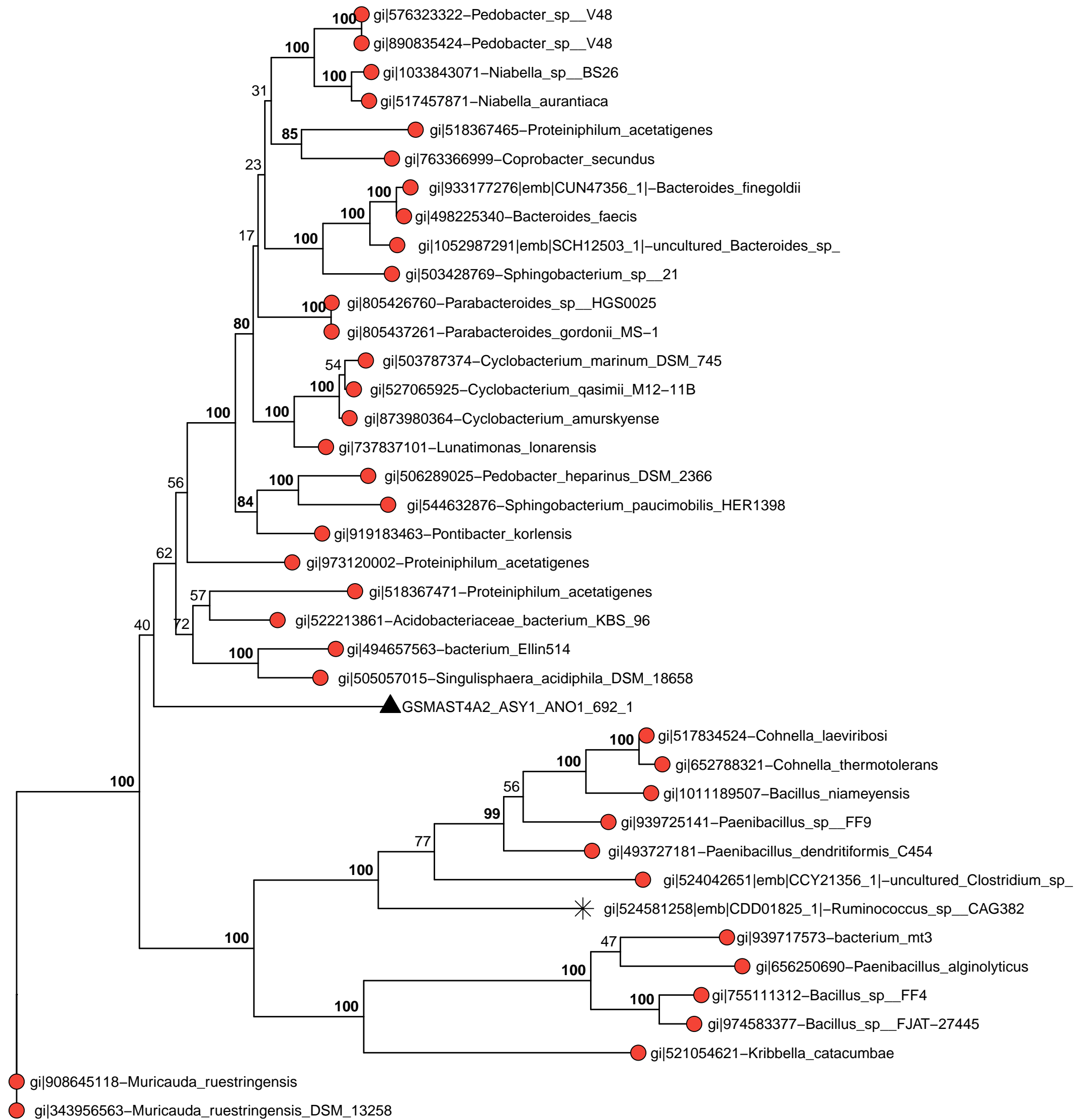
● Bacteria
▲ SAG

0.1

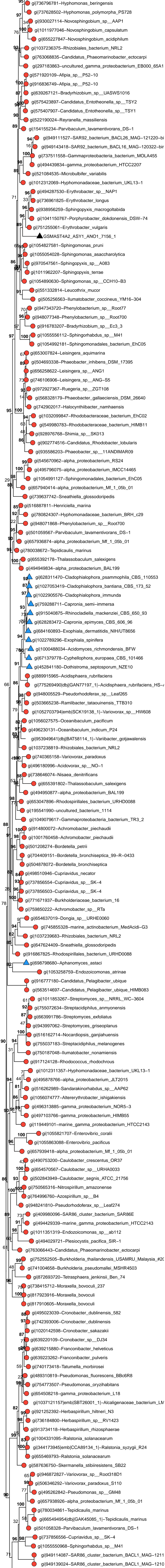


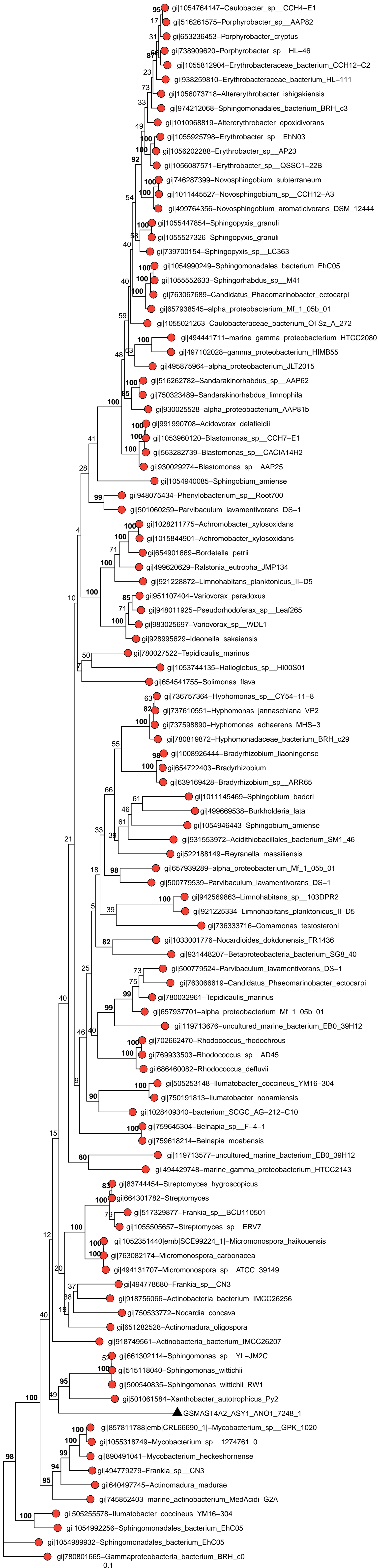
● Bacteria
▲ SAG

0.2

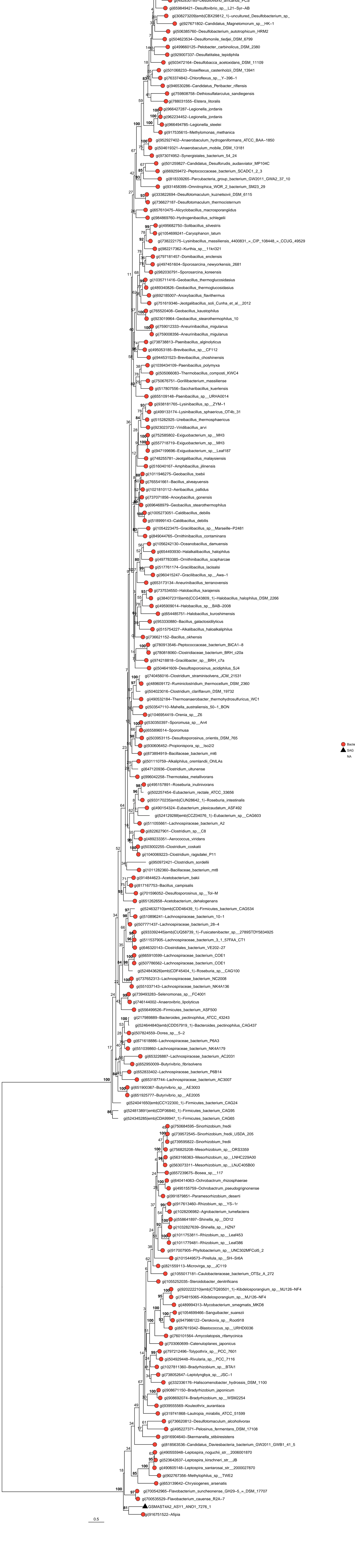


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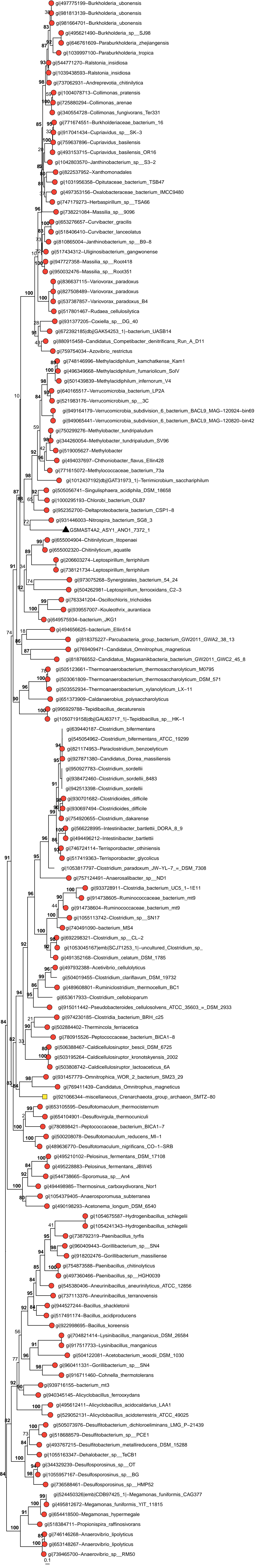


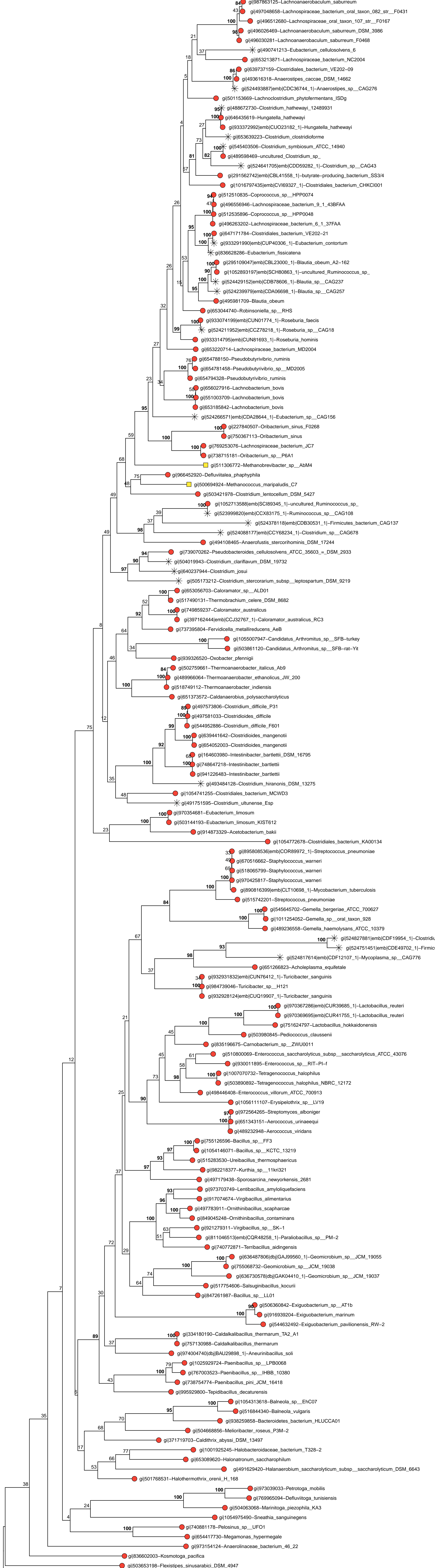


● Bacteria
▲ SAG



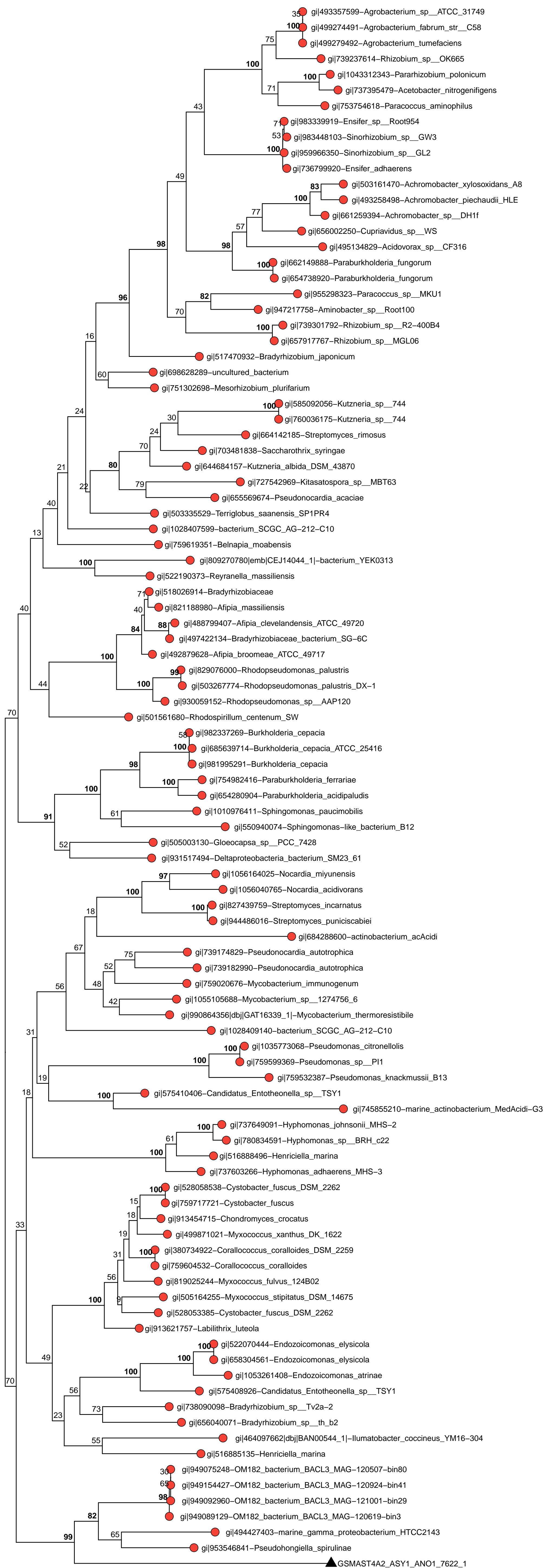
● Bacteria
▲ SA
● NA

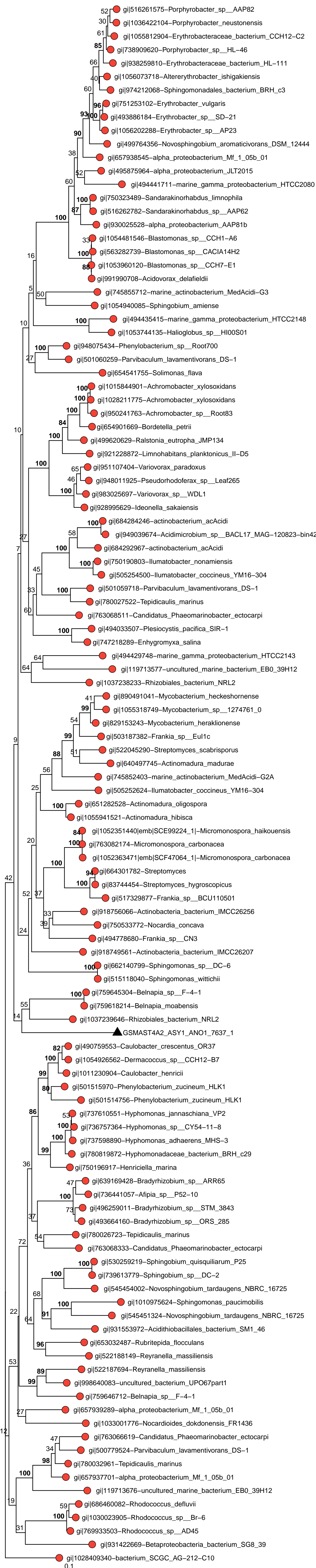


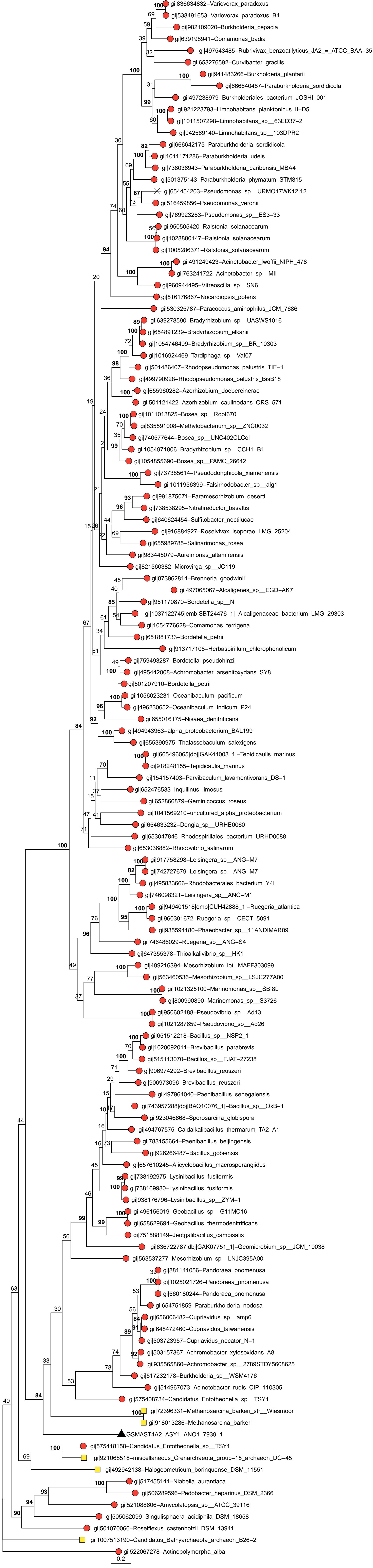


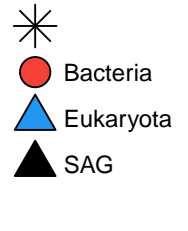
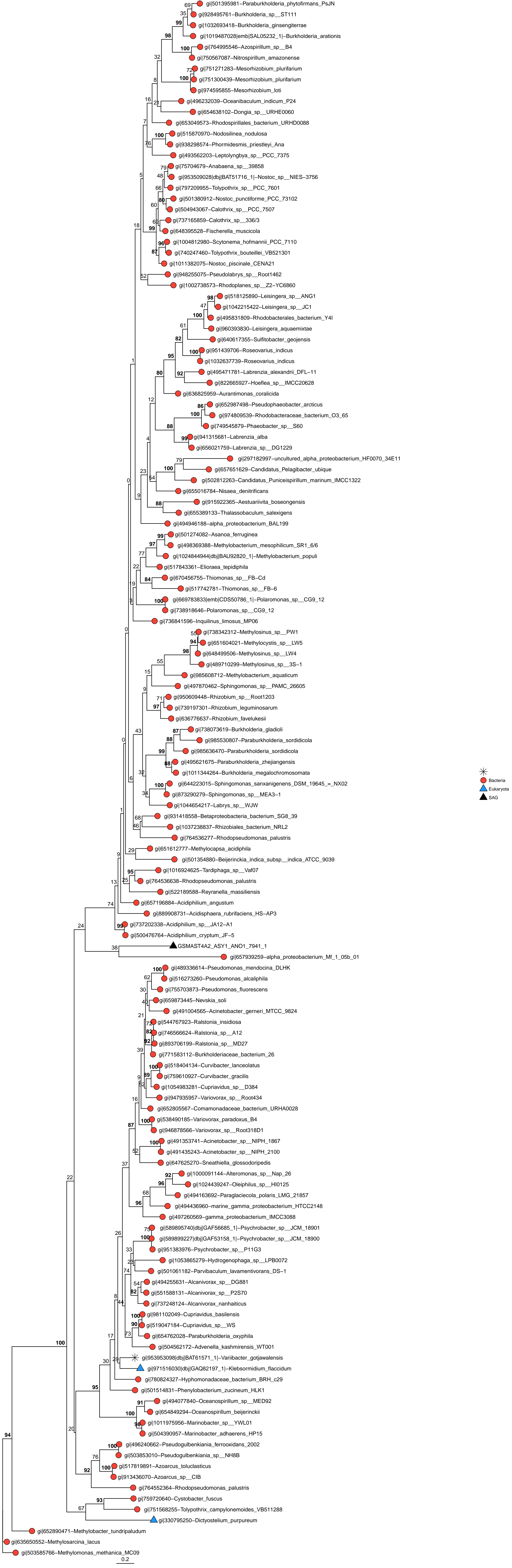
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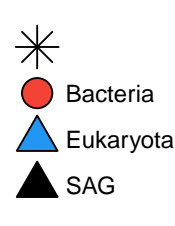
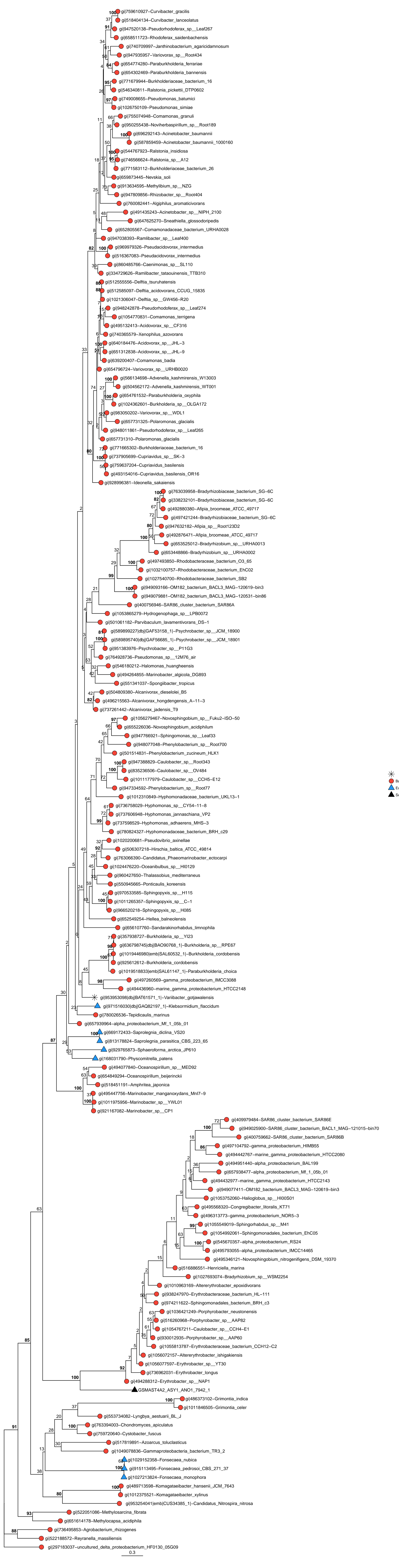






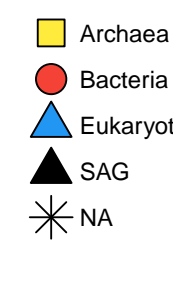
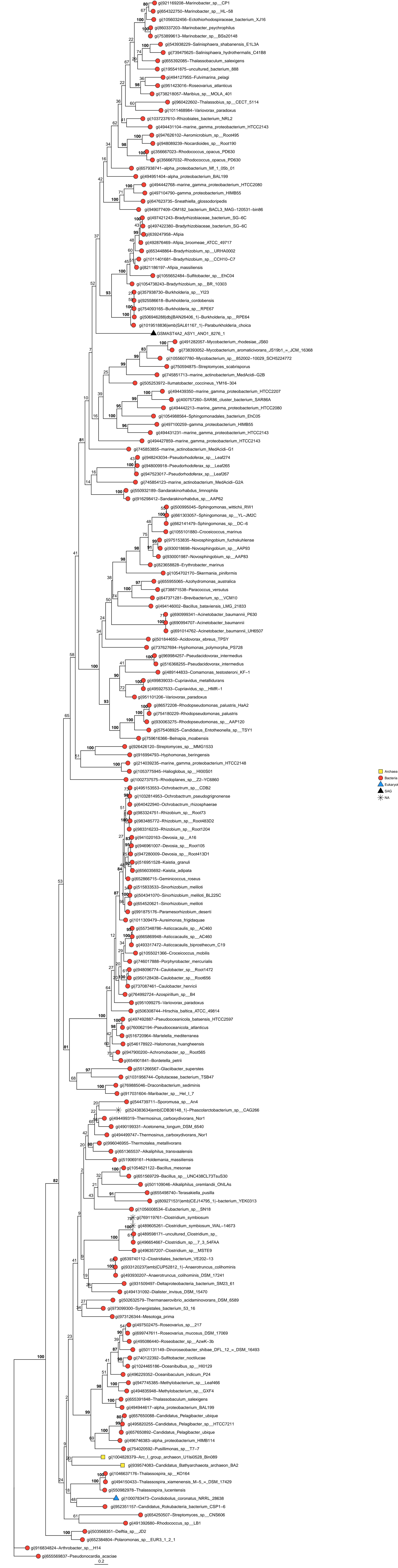


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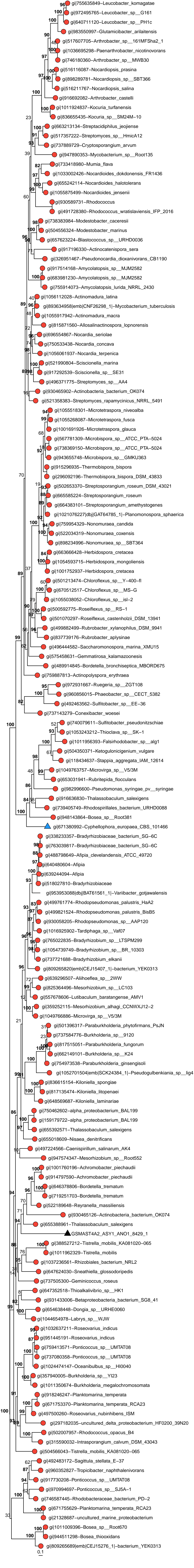


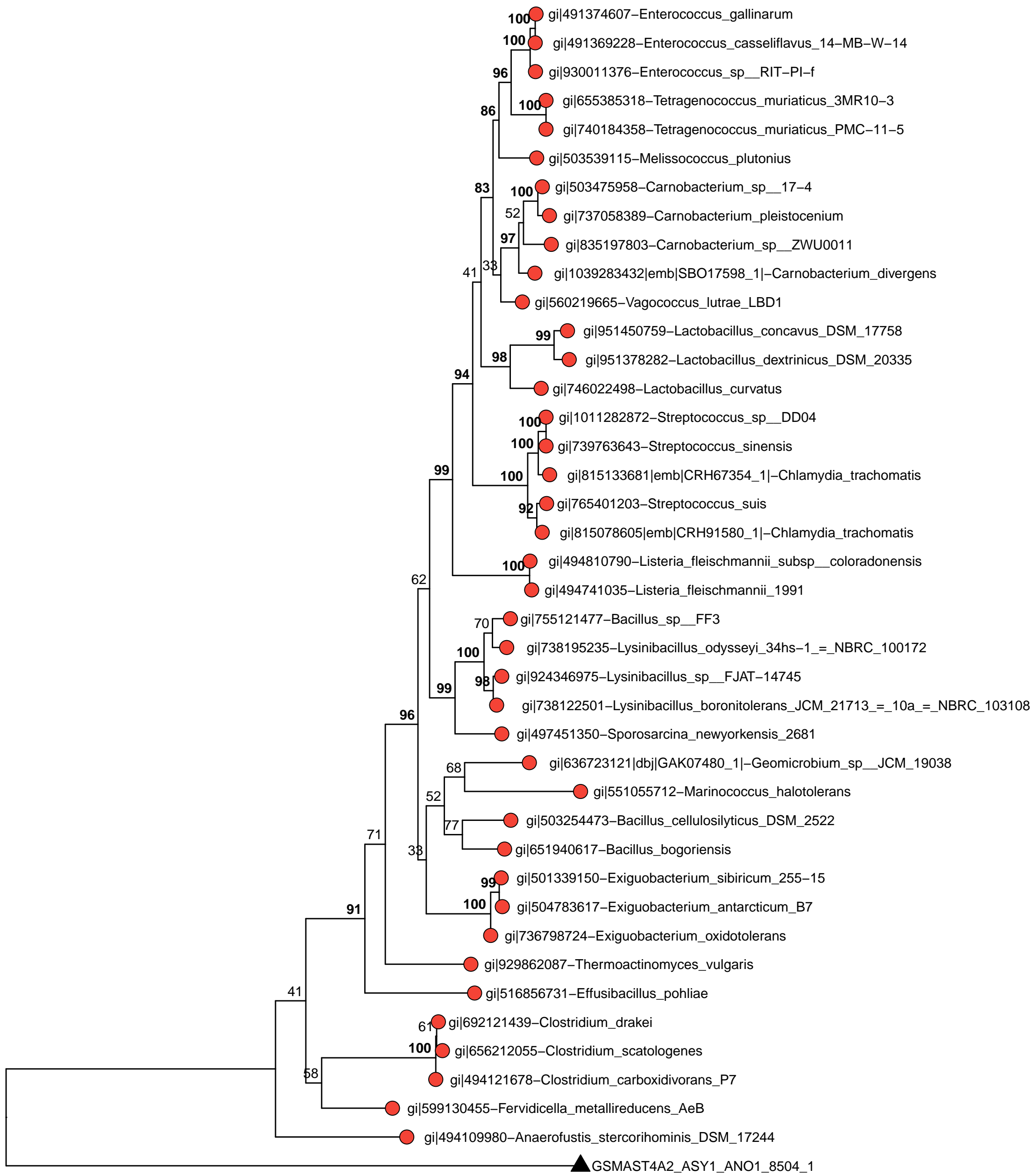
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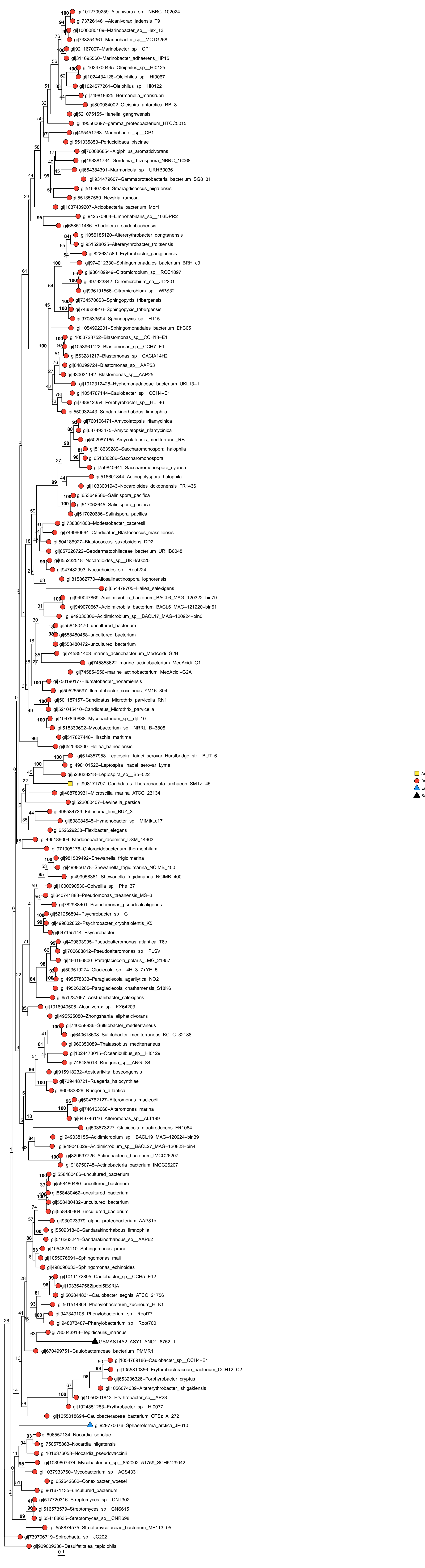
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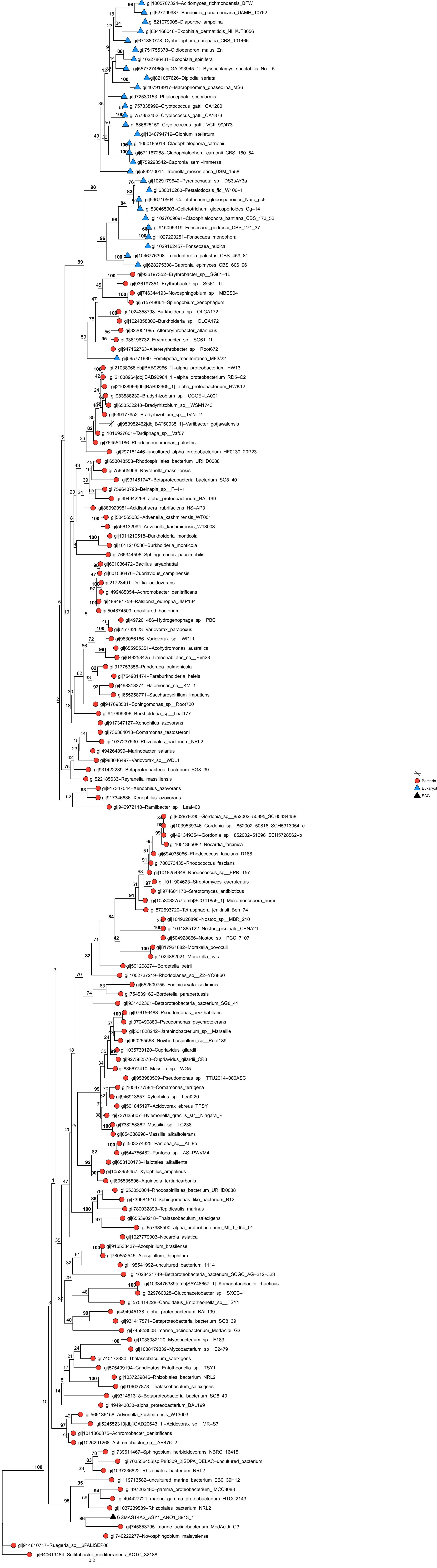


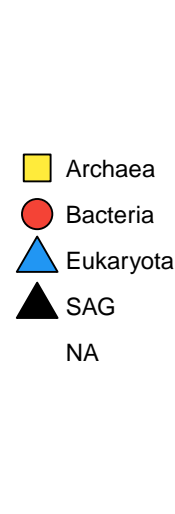
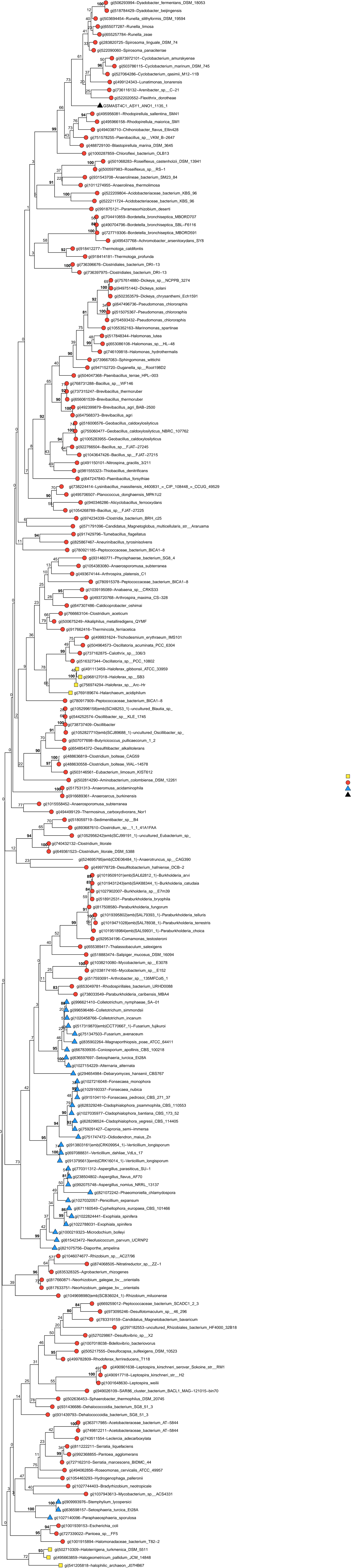
● Bacteria
▲ SAG

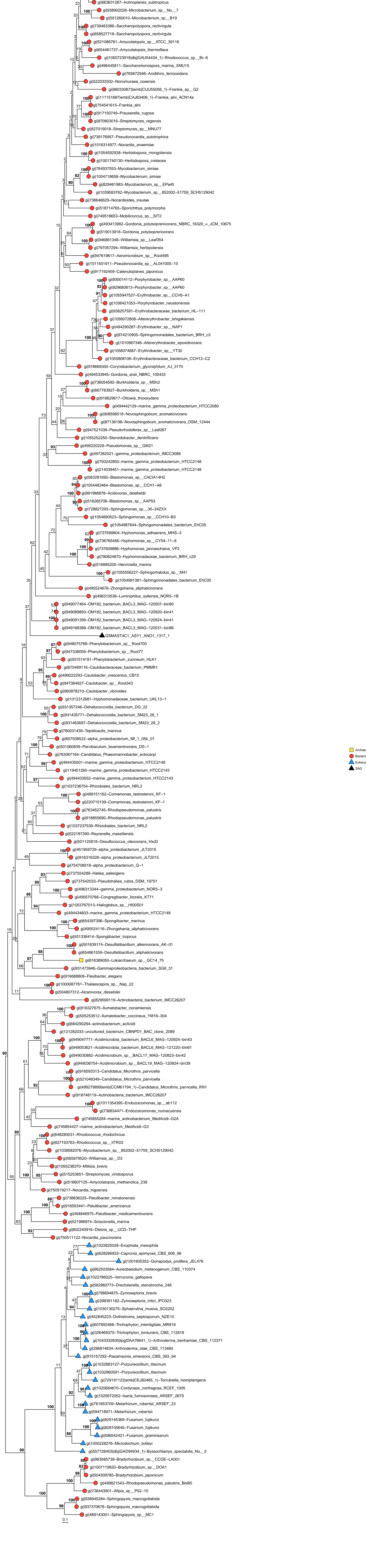
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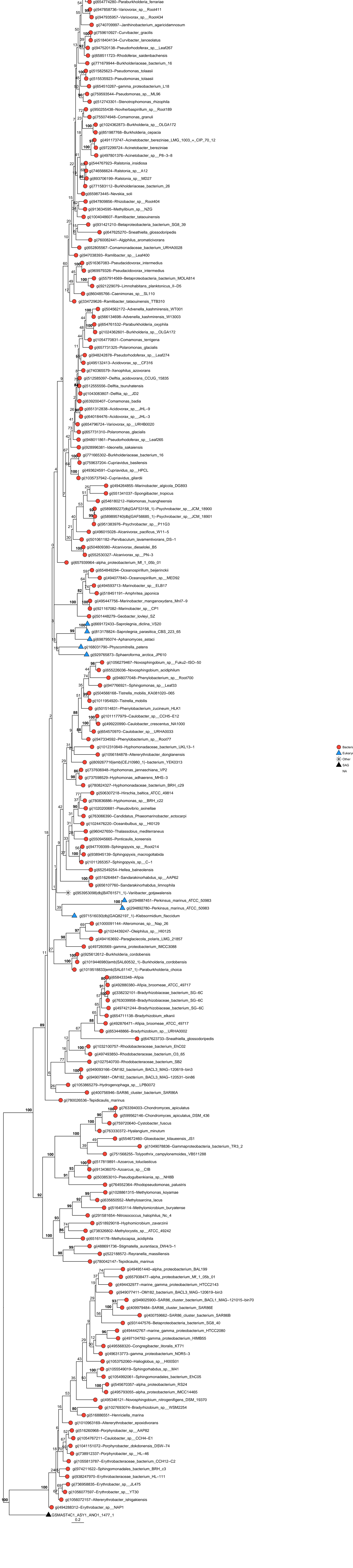


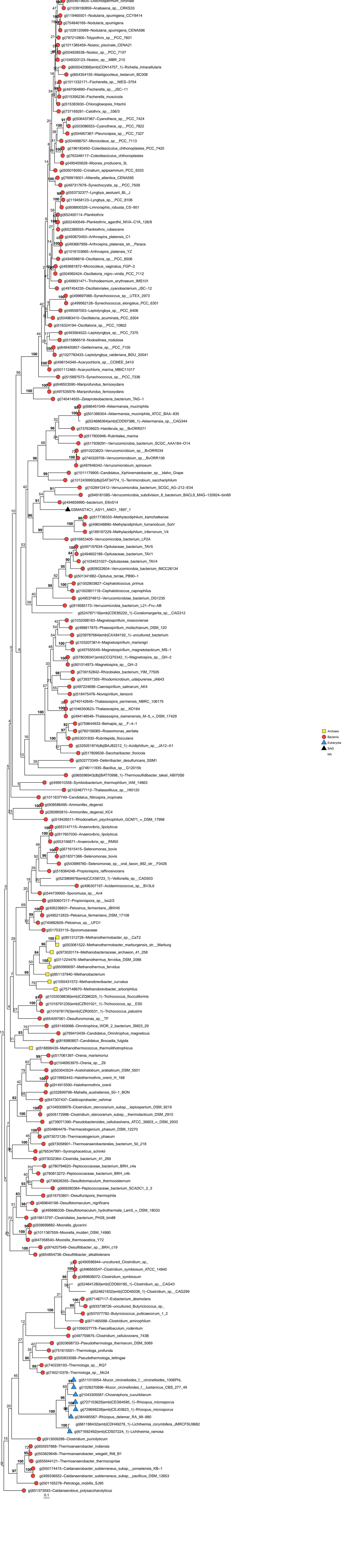




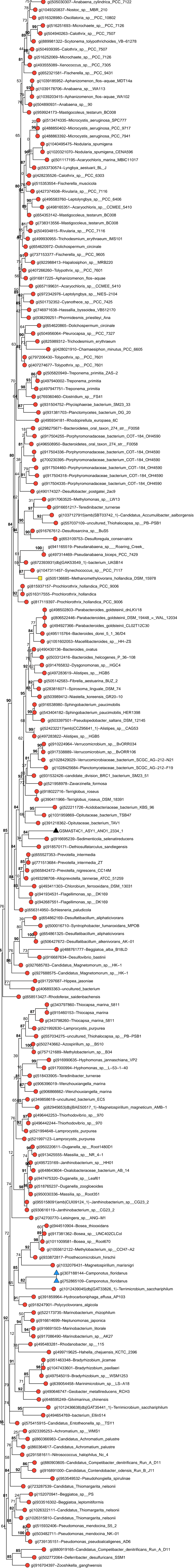


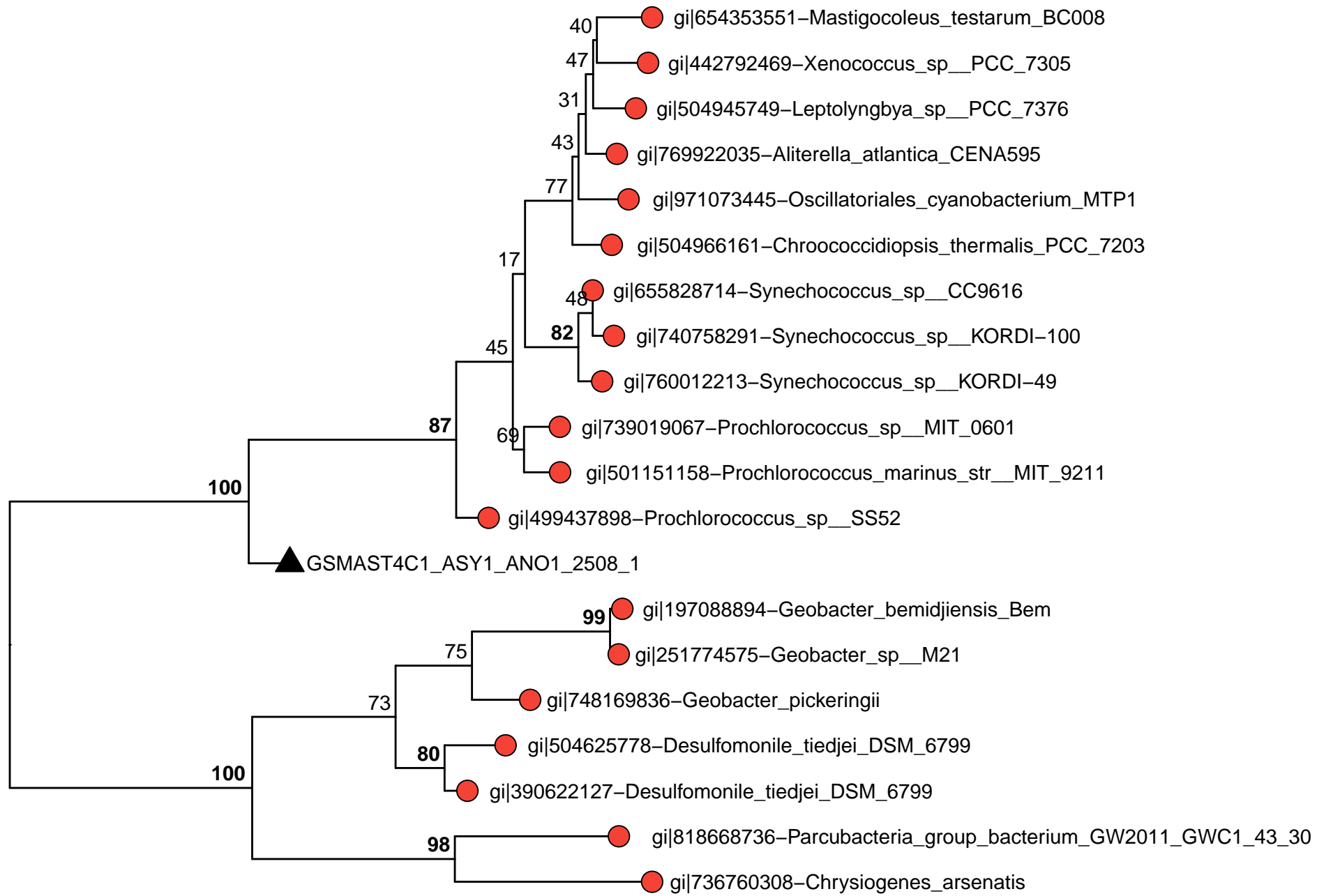


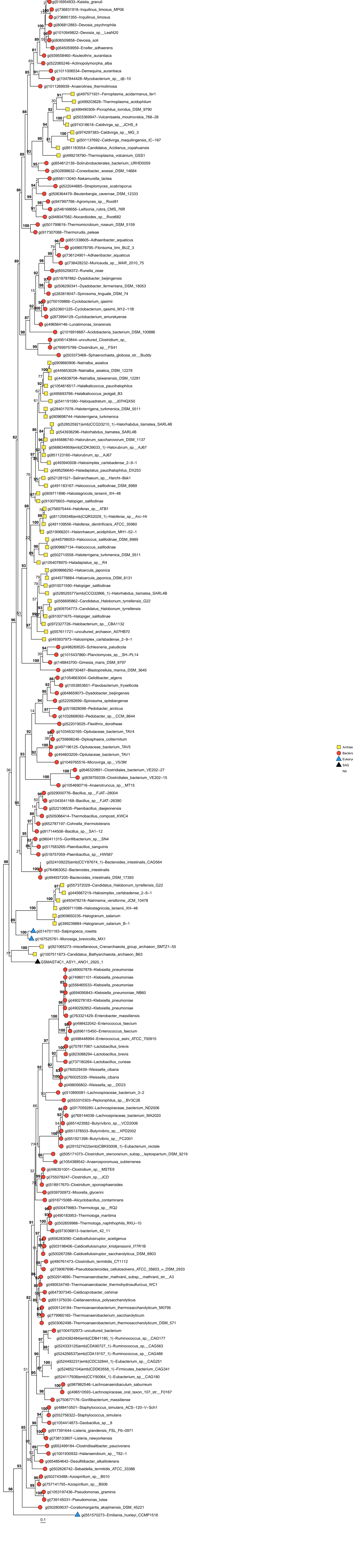


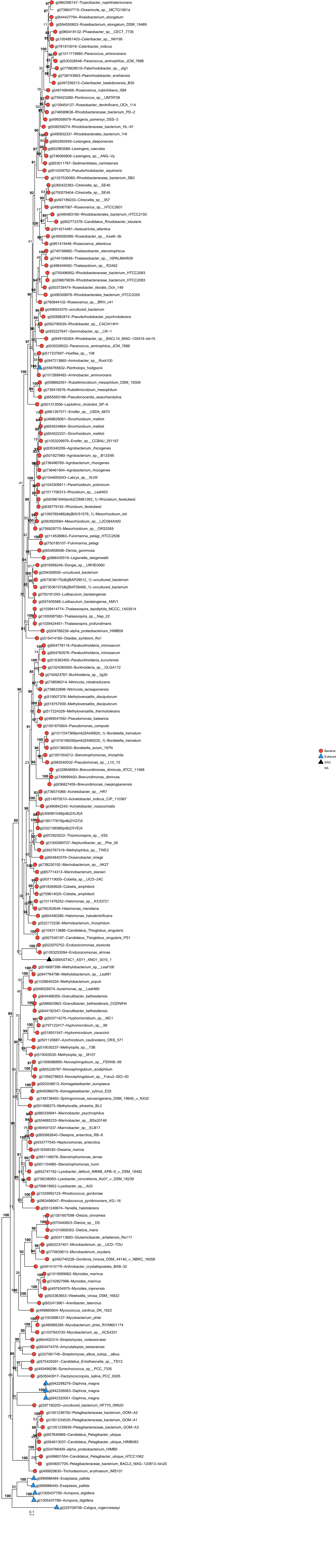


■ Archaea
● Bacteria
▲ Eukaryota
▲ SAG
 NA

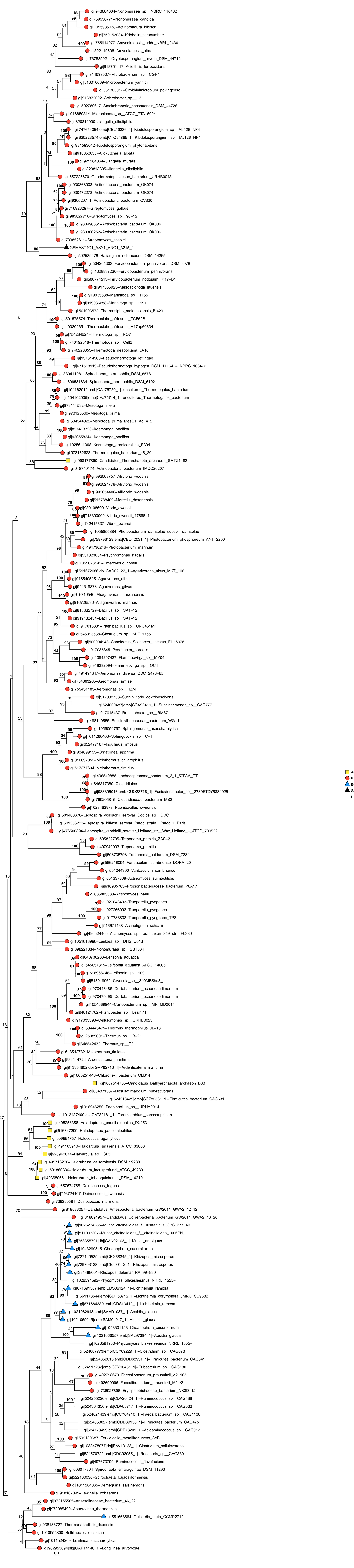


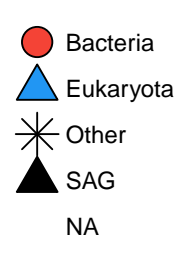
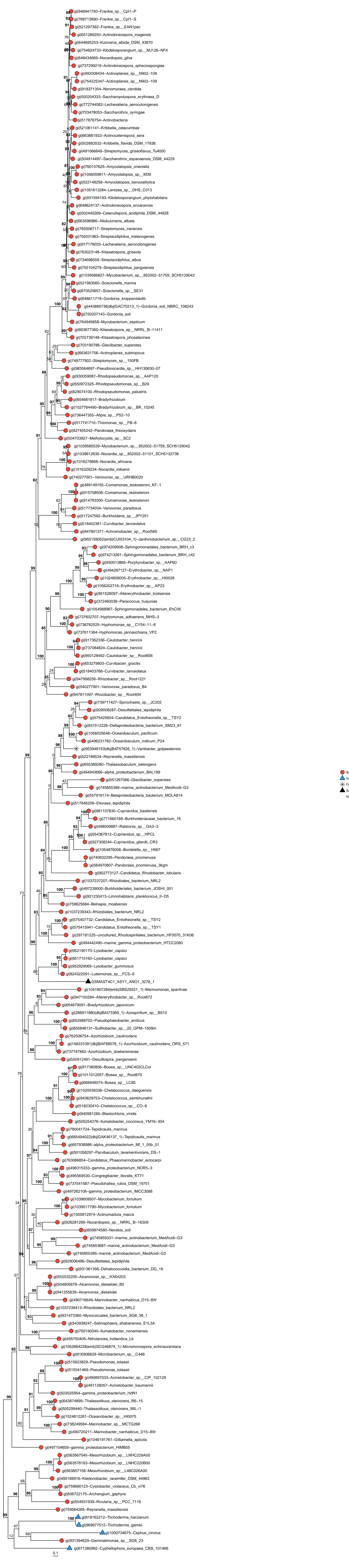






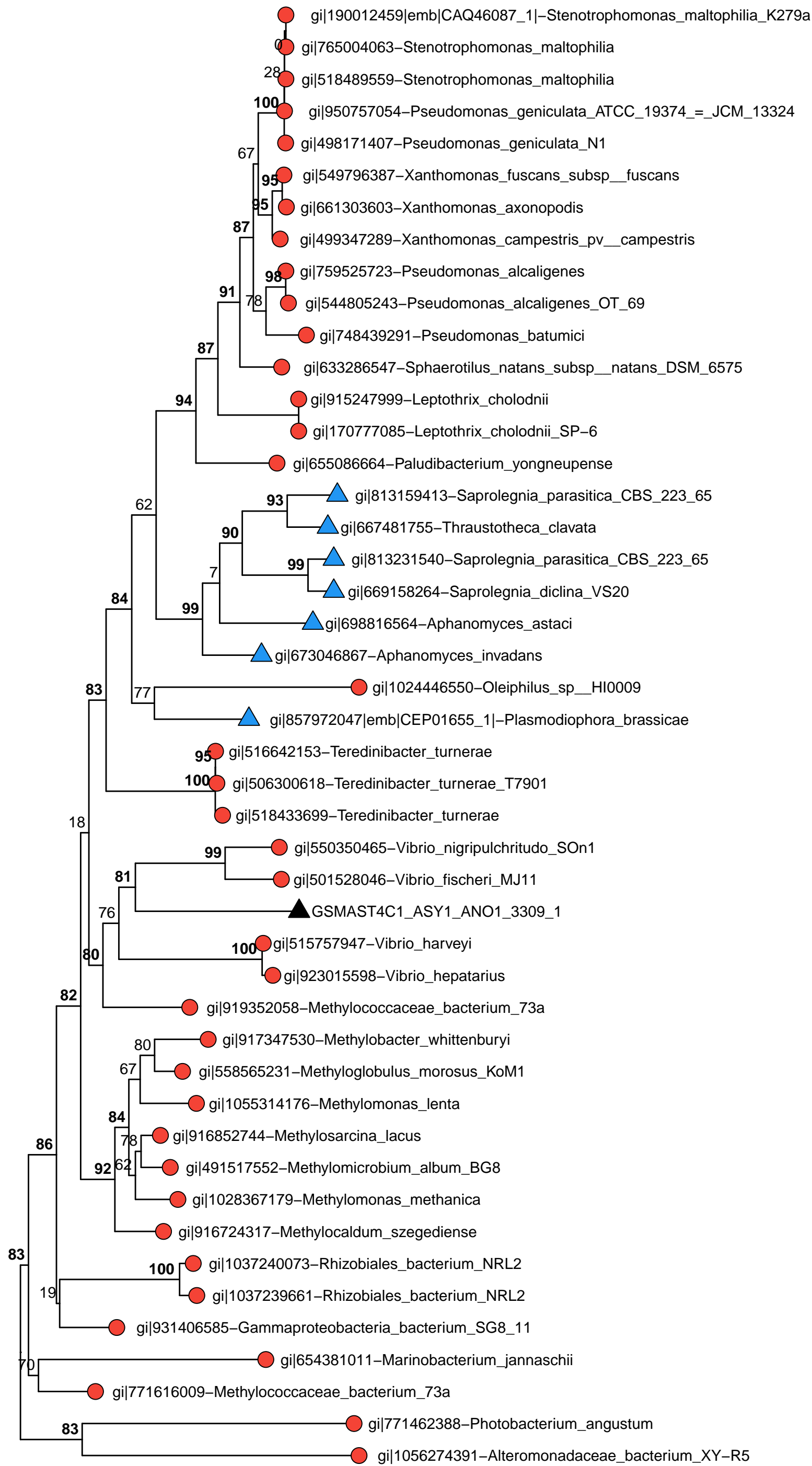
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 gi|960418132-Phaeobacter_sp__CECT_7735
 gi|1054951403-Celeribacter_sp__NH195
 gi|761810016-Celeribacter_indicus
 gi|1011719980-Paracoccus_aminovorans
 gi|530328548-Paracoccus_aminophilus_JCM_7686
 gi|77928519-Falsirhodobacter_sp__alg1
 gi|738743893-Paenirhodobacter_enshiensis
 gi|497256213-Celeribacter_baekdonensis_B30
 gi|497499456-Roseovarius_nubinihibens_ISM
 gi|759423289-Ponticoccus_sp__UMTAT08
 gi|109454127-Roseobacter_denitrificans_OCh_114
 gi|746588638-Rhodobacteraceae_bacterium_PD-2
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 gi|495832331-Rhodobacterales_bacterium_Y41
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 gi|652983588-Leisingera_caerulea
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 gi|1027539360-Rhodobacteraceae_bacterium_SB2
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 gi|951419446-Roseovarius_atlanticus
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 gi|947213660-Aminobacter_sp__Root100
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 gi|501313556-Leptothrix_cholodnii_SP-6
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 gi|501827980-Agrobacterium_sp__B133/95
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 gi|937119005-Cobetia_sp__UCD-24C
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 gi|1001657598-Dietzia_cinnamea
 gi|570440823-Dietzia_sp__D5
 gi|1010958352-Dietzia_maris
 gi|503113683-Glutamicibacter_arilaitensis_Re117
 gi|602237401-Microbacterium_sp__UCD-TDU
 gi|770639513-Microbacterium_oxydans
 gi|491410179-Arthrobacter_hirsallotopietes_DSM_44140_=_NBRC_16056
 gi|1019589062-Myroides_marinus
 gi|742827996-Myroides_marinus
 gi|497934975-Myroides_injenensis
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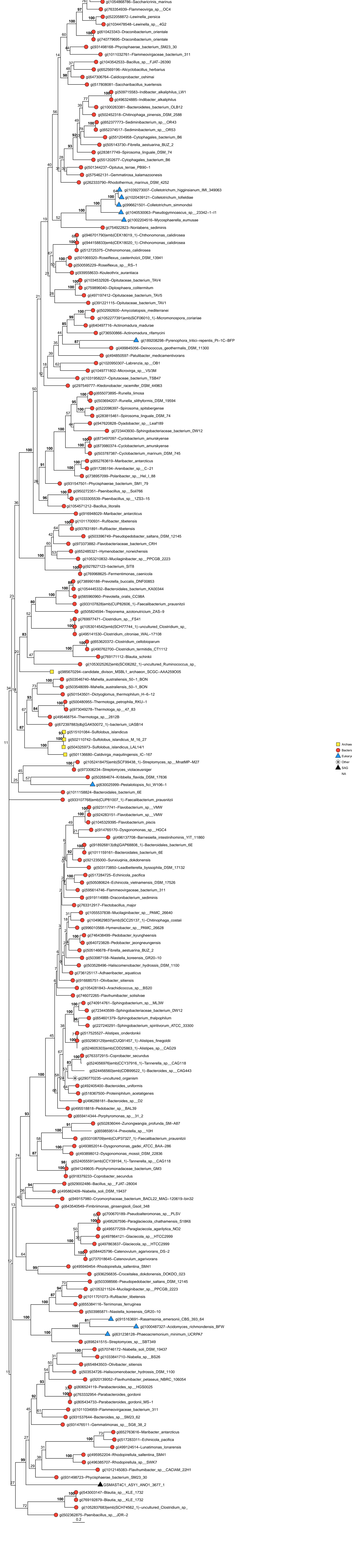
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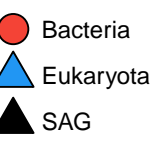
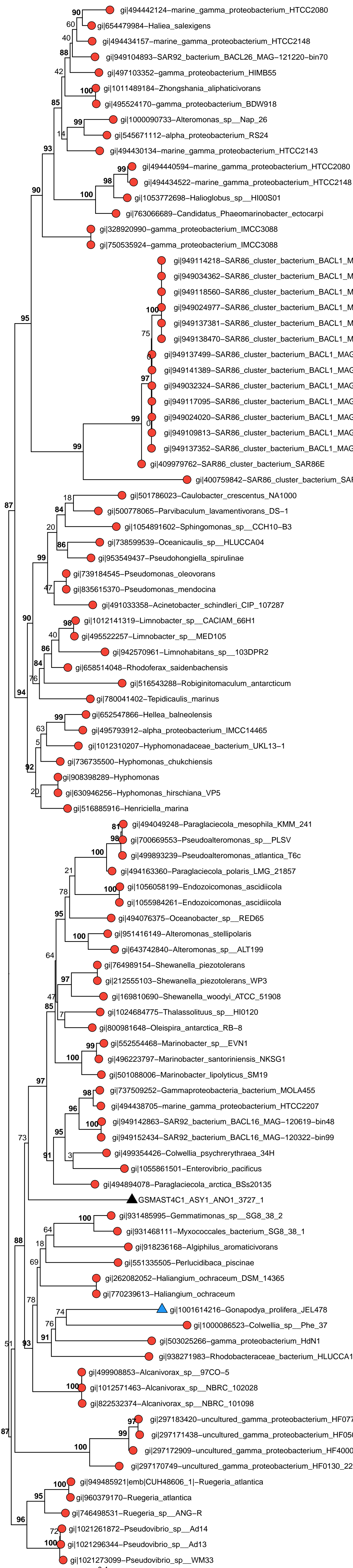
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 gj|651289250--Actinokineospora_inagensis
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 gj|737299219--Actinokineospora_sphaciospongiae
 gj|492008434--Actinoplanes_sp__N902-109
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 gj|772744582--Lechevalieria_aerocolonigenes
 gj|703478053--Saccharothrix_syringae
 gj|517876754--Actinobacteria
 gj|521061141--Kribbella_catacumbae
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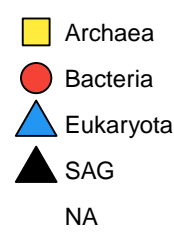
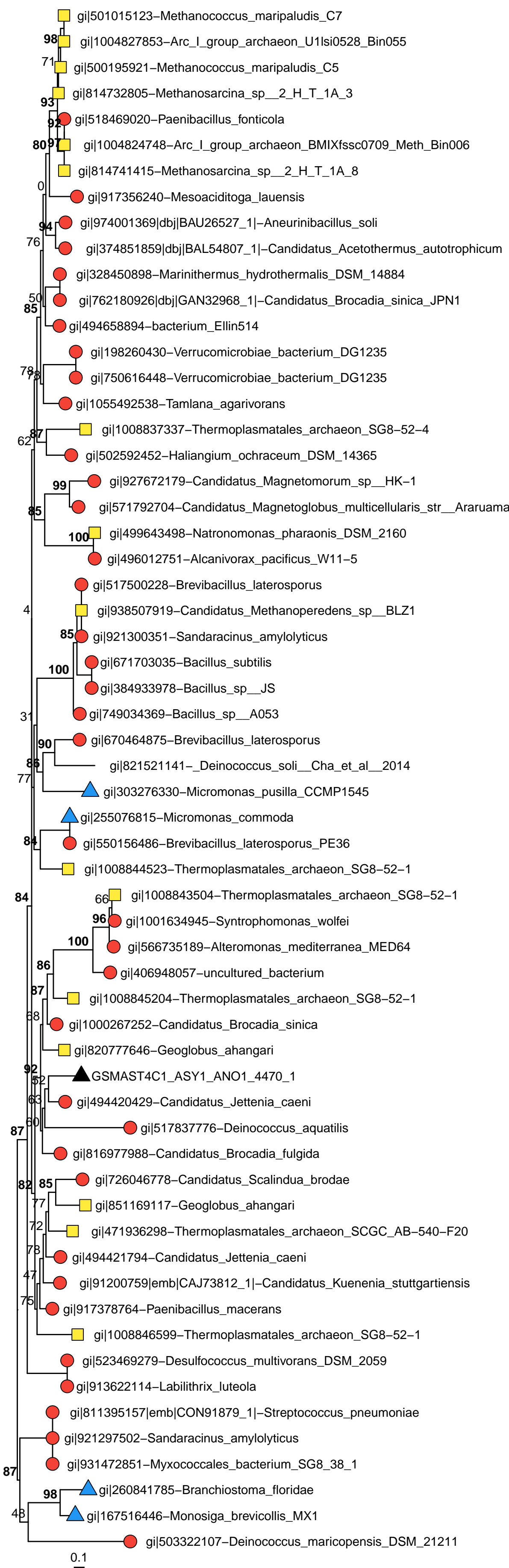


● Bacteria
▲ Eukaryota
▲ SAG

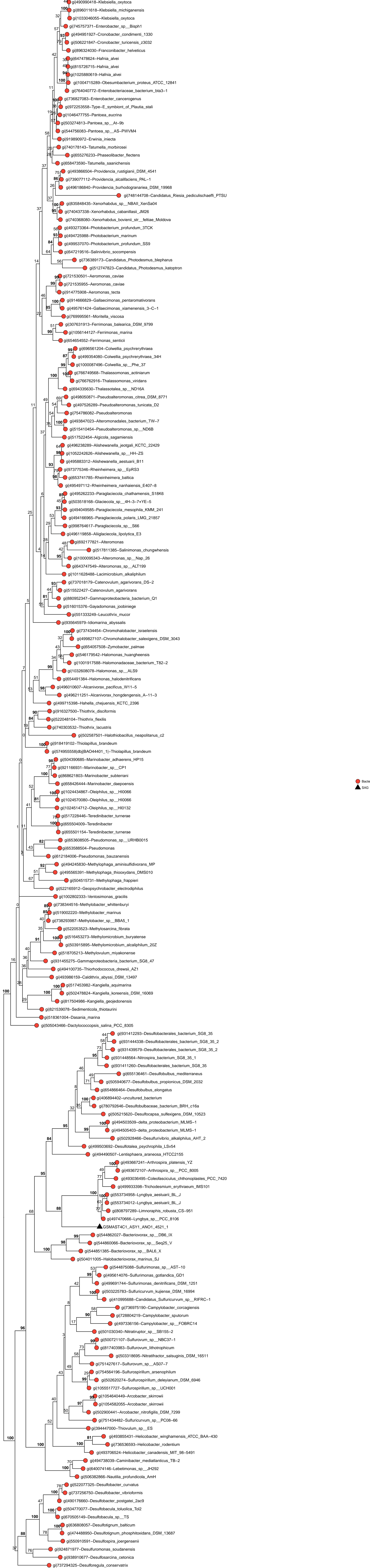
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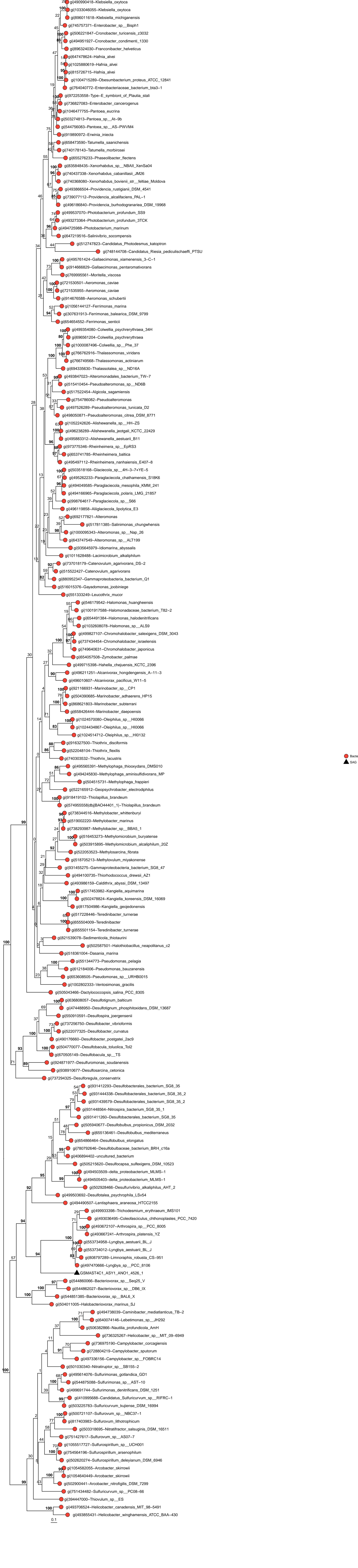




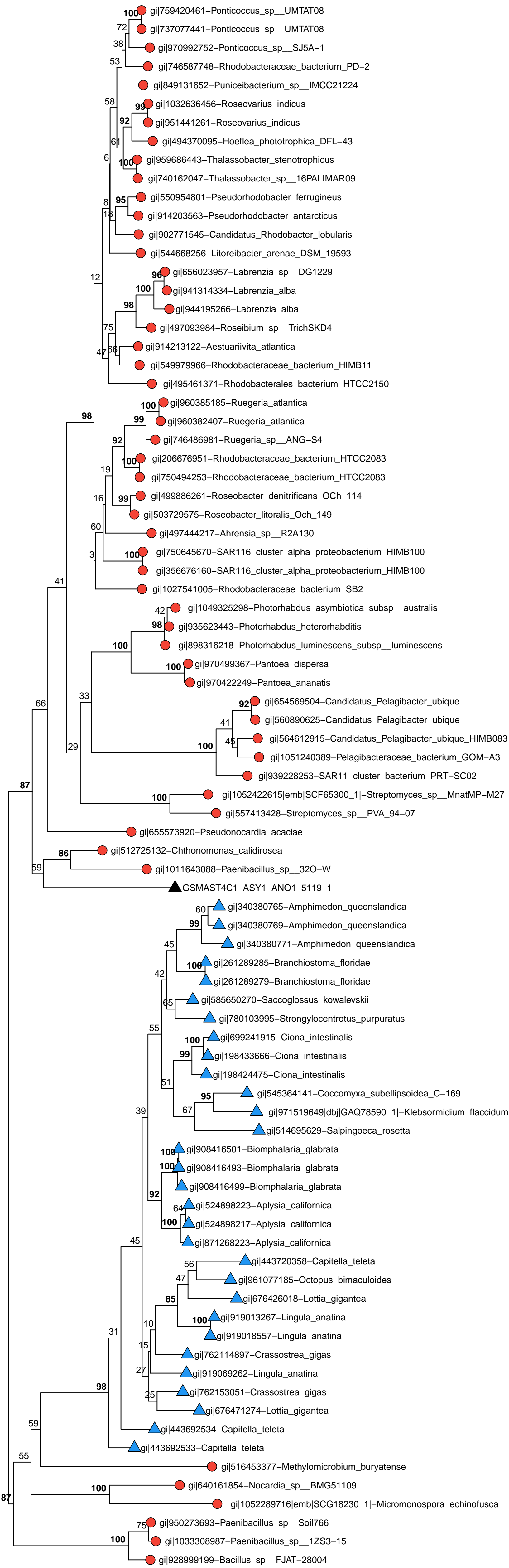


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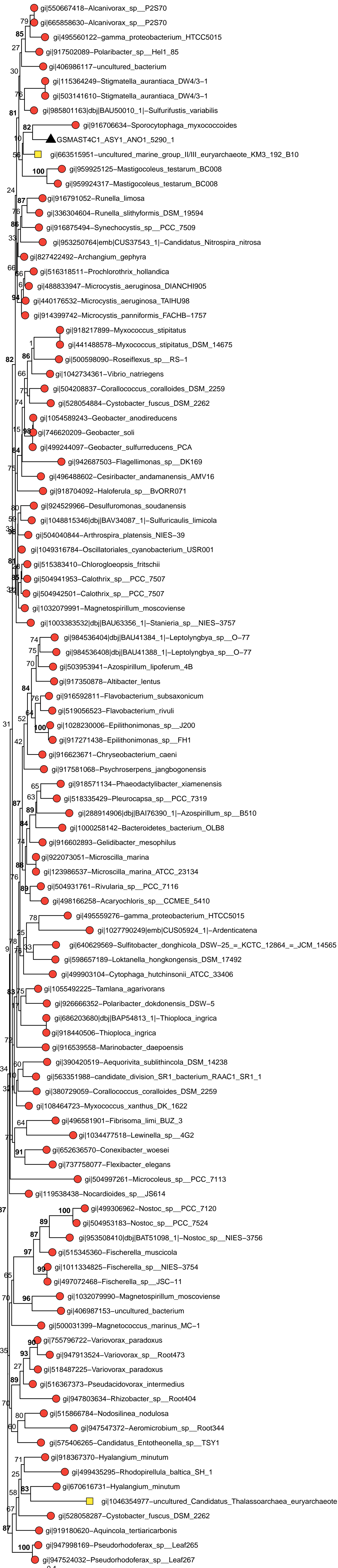


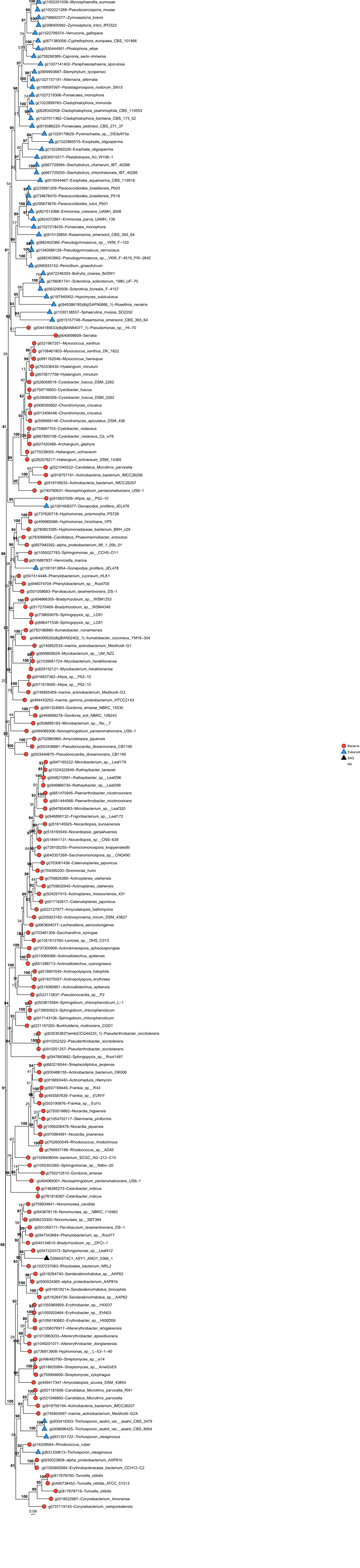


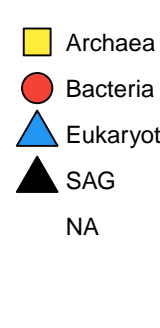
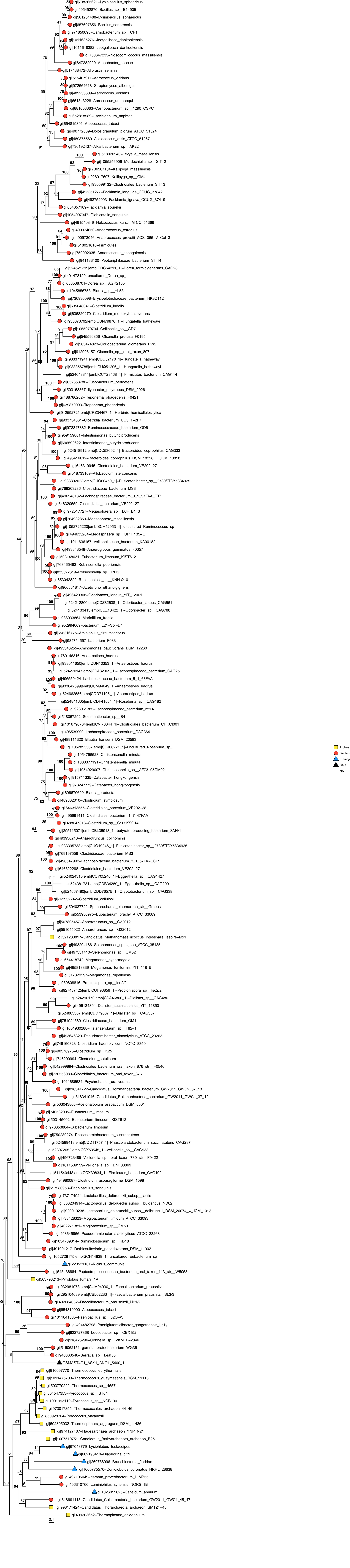


● Bacteria
 ▲ Eukaryota
 ▲ SAG

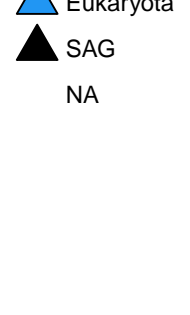
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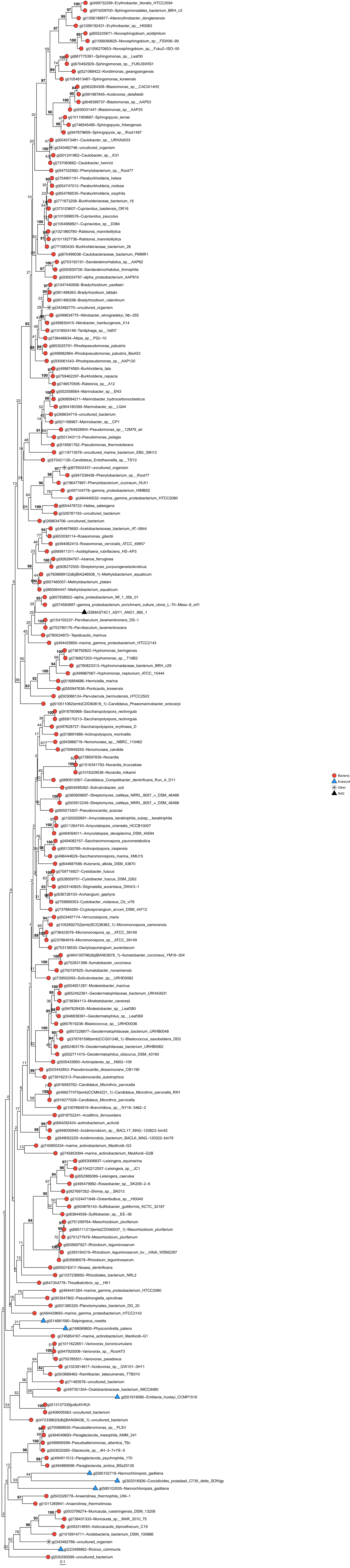


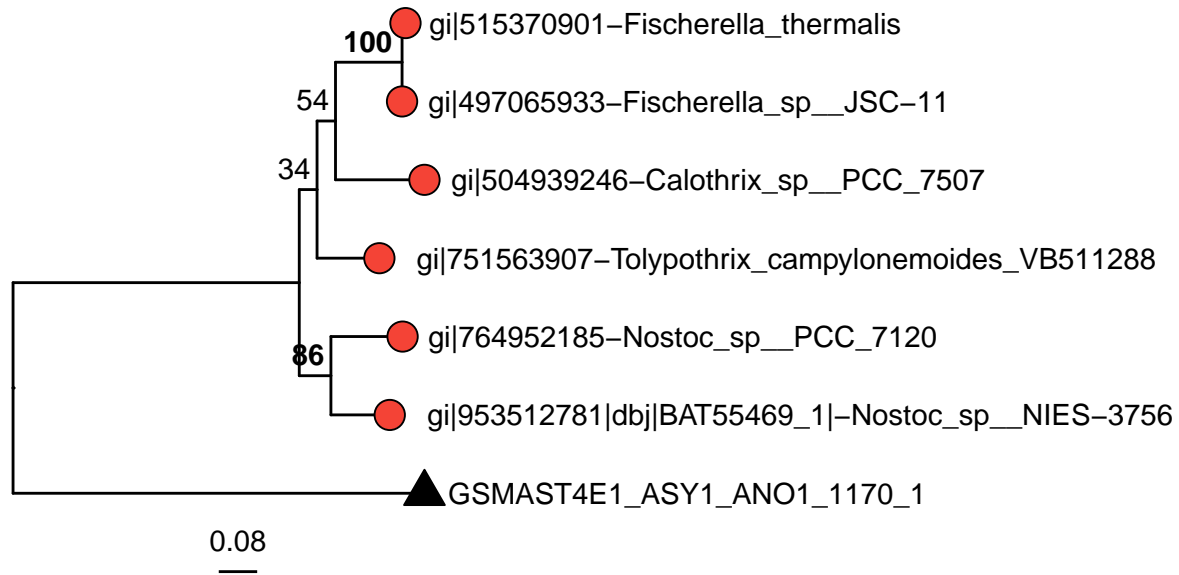


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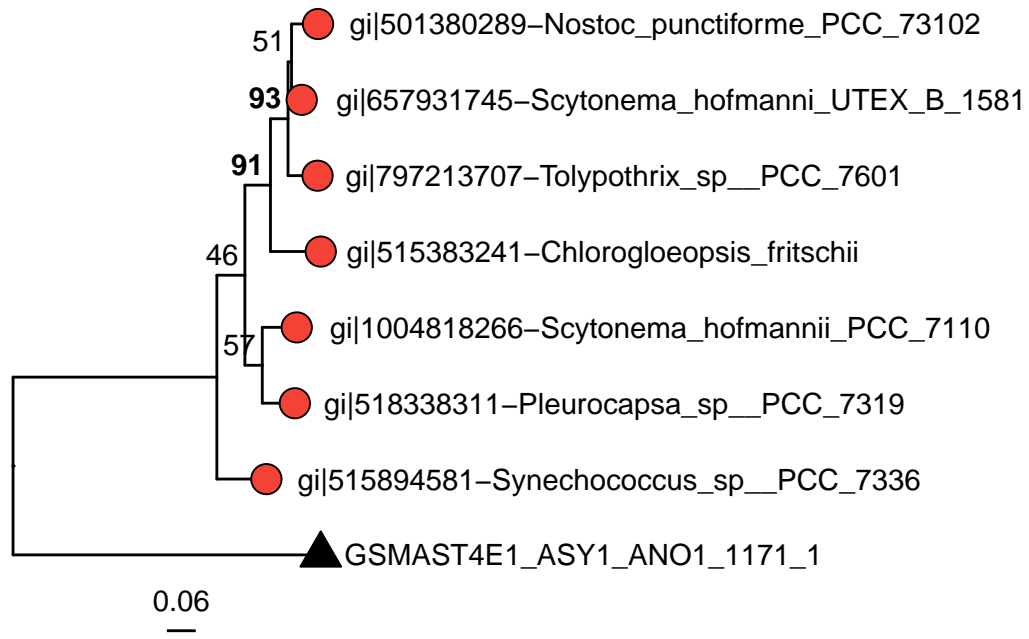


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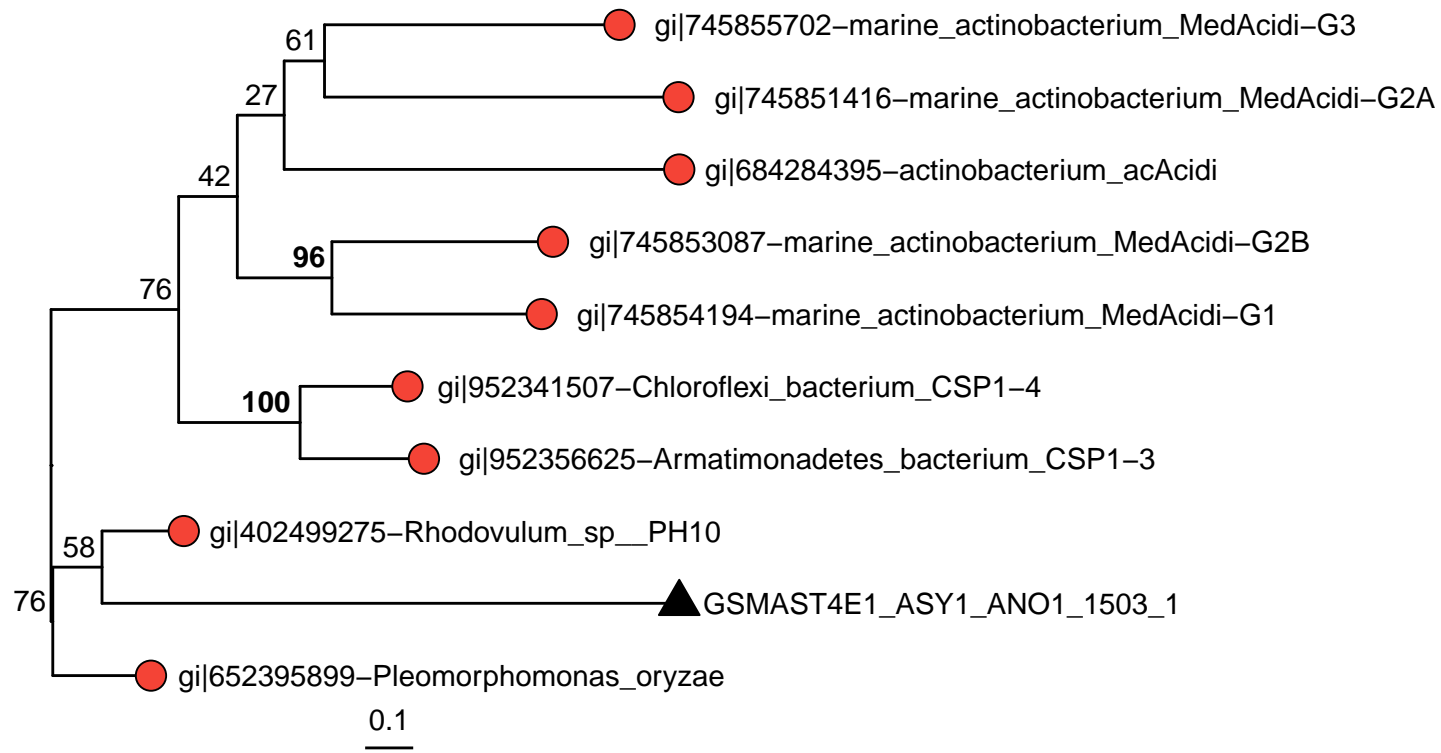


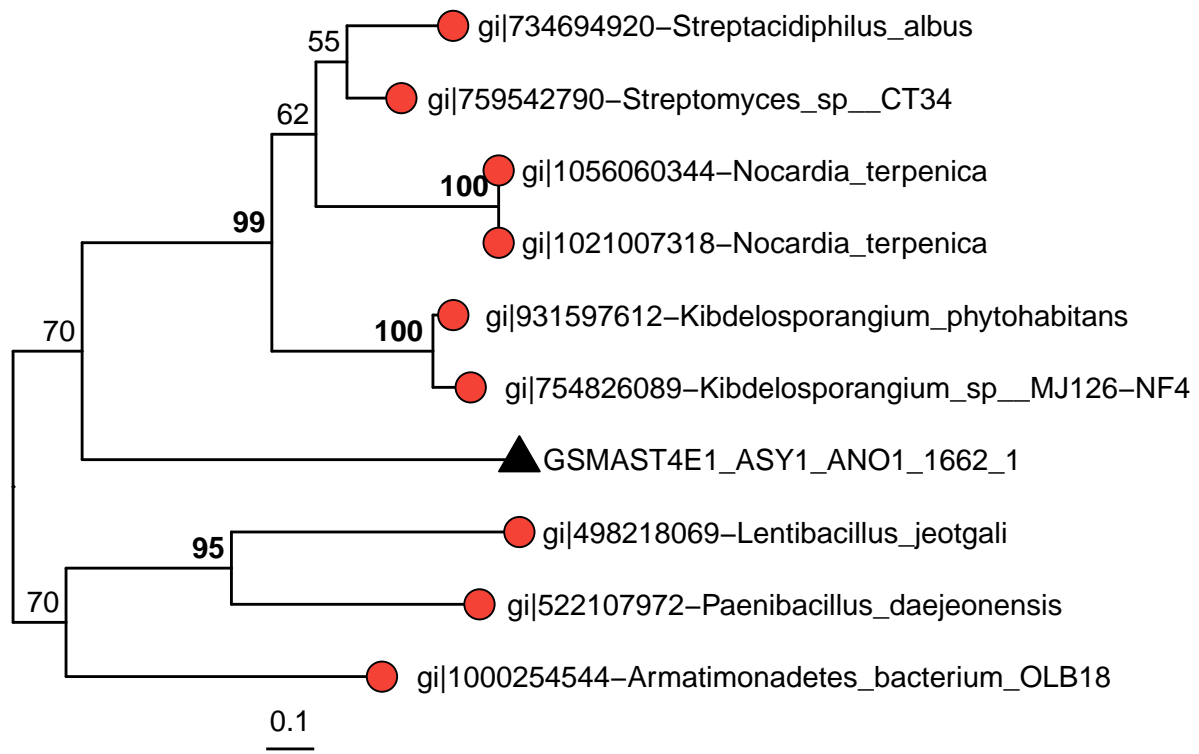


● Bacteria
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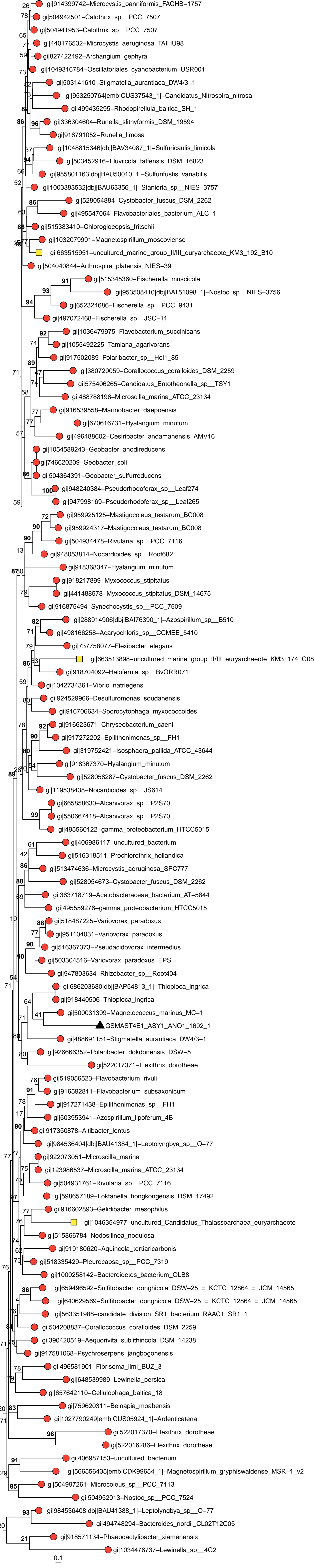


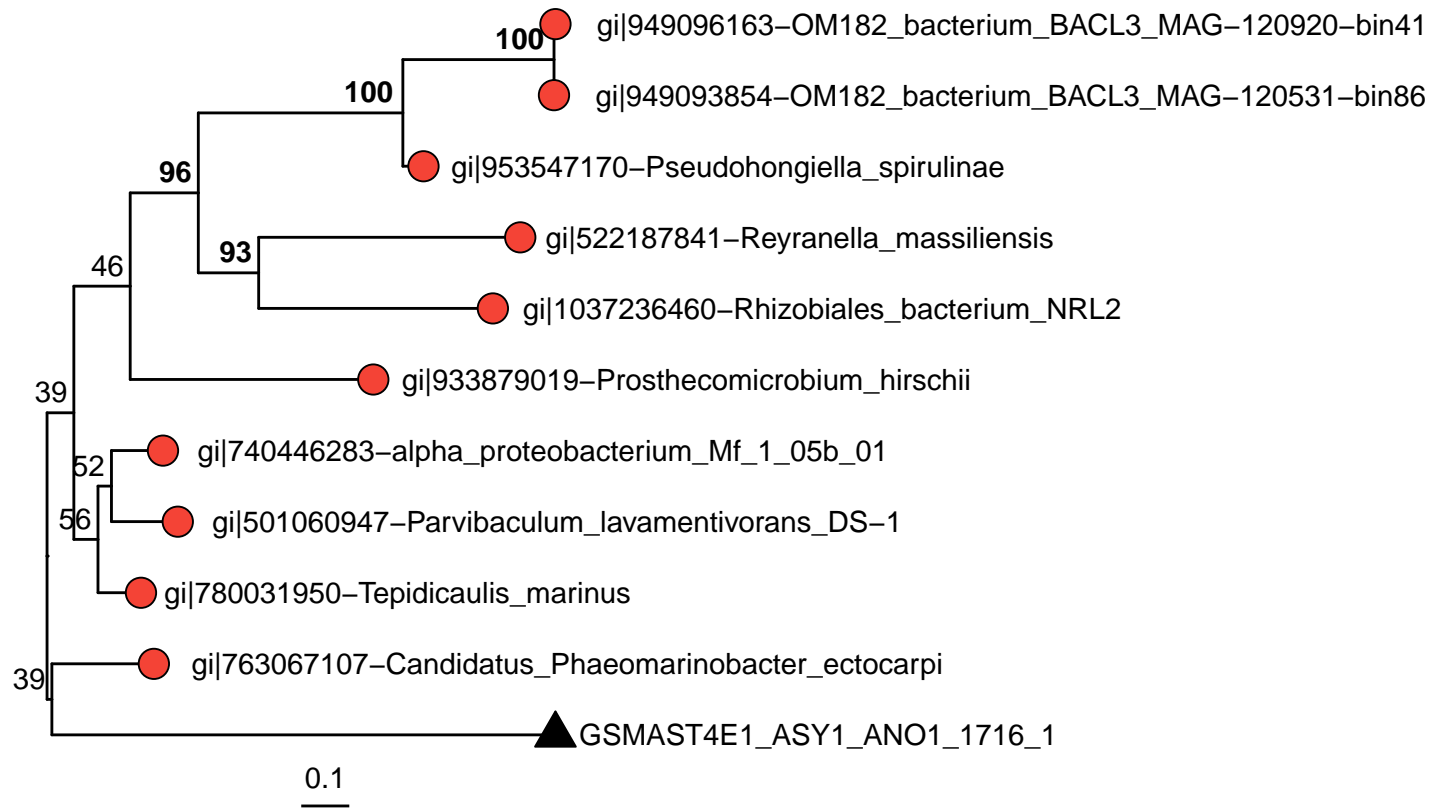
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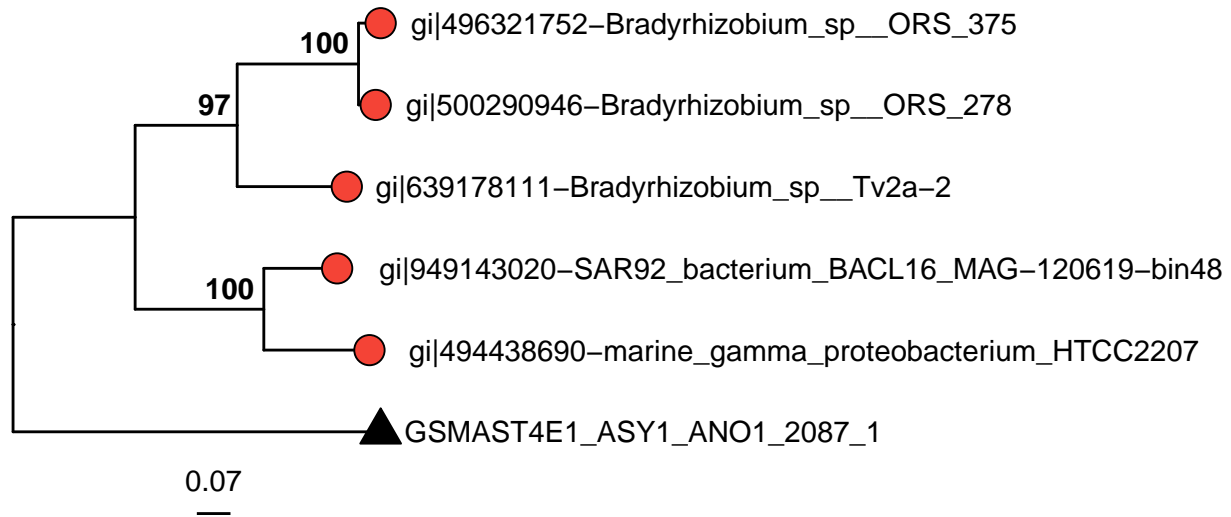


● Bacteria
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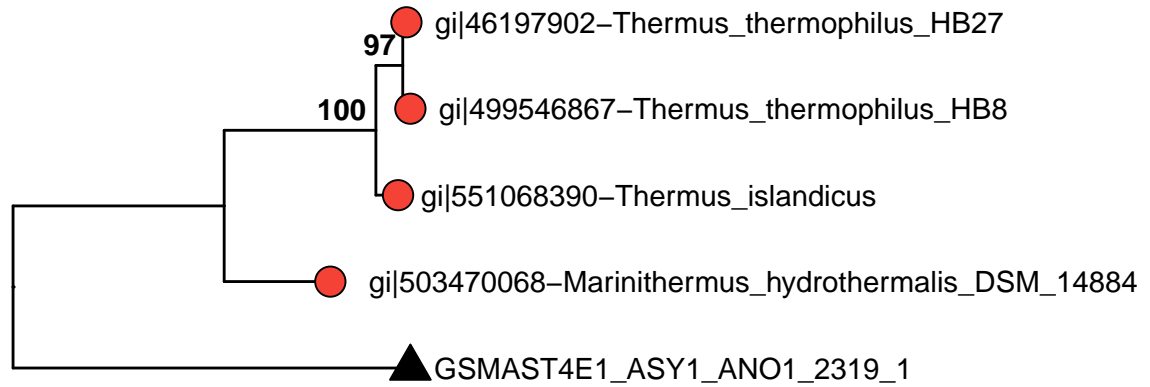




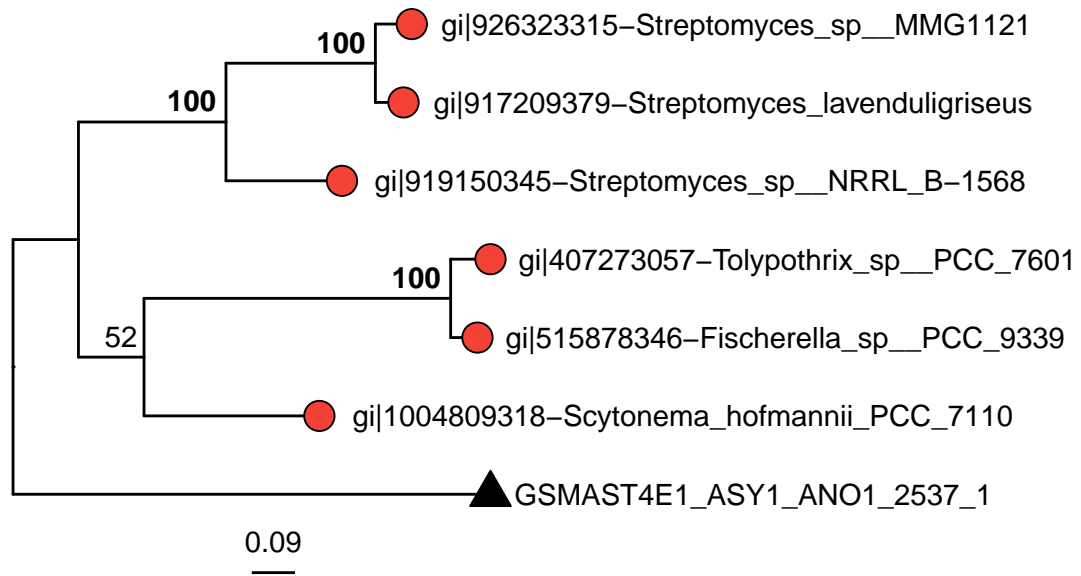
● Bacteria
▲ SAG



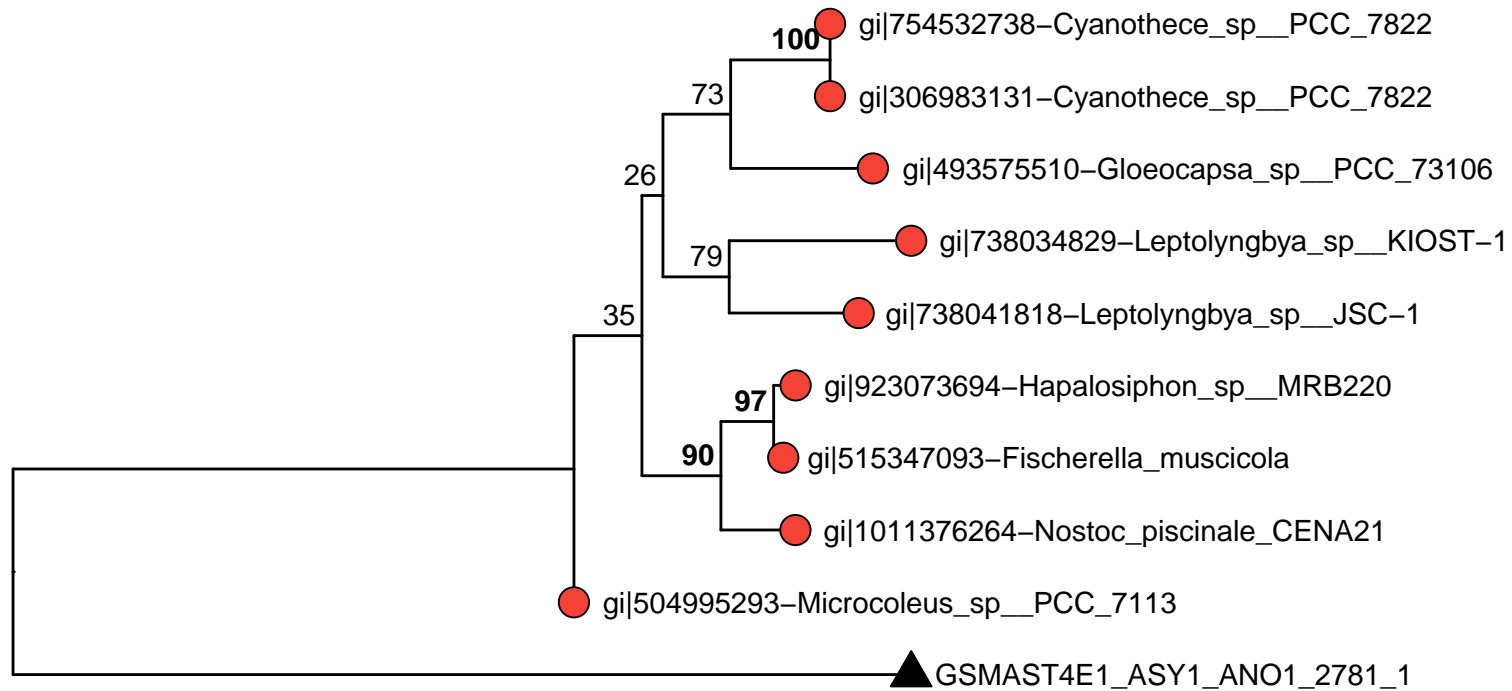
● Bacteria
▲ SAG



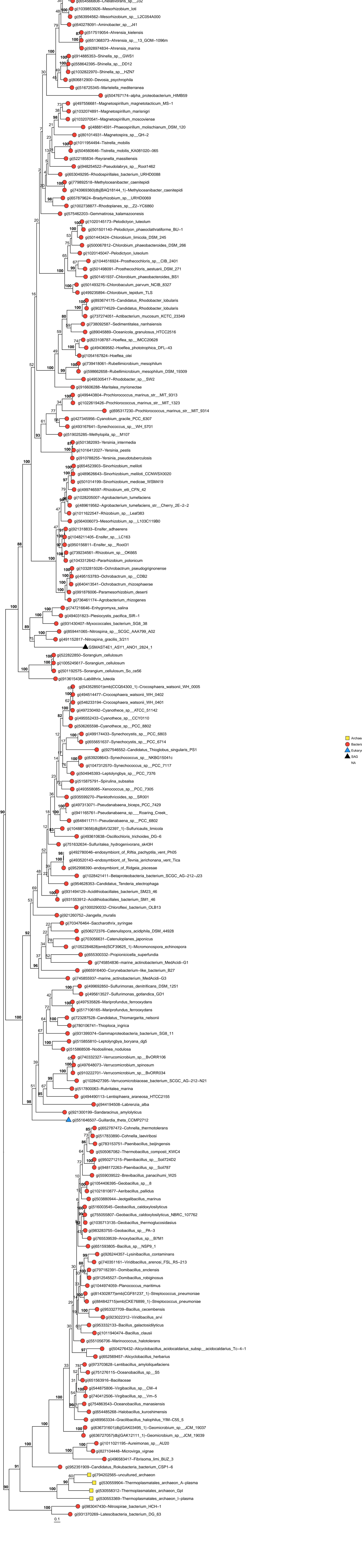
● Bacteria
▲ SAG

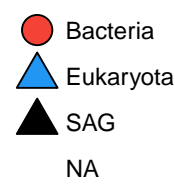
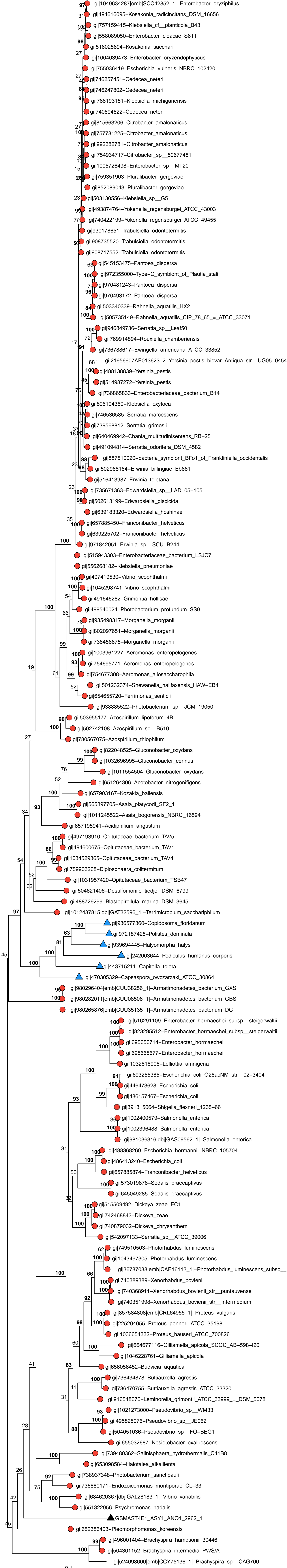


● Bacteria
▲ SAG

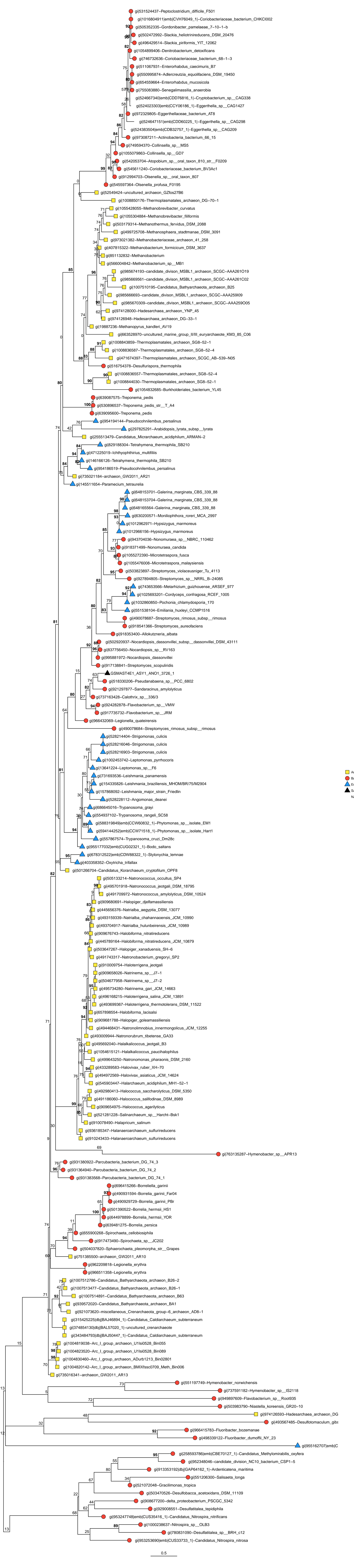


● Bacteria
▲ SAG



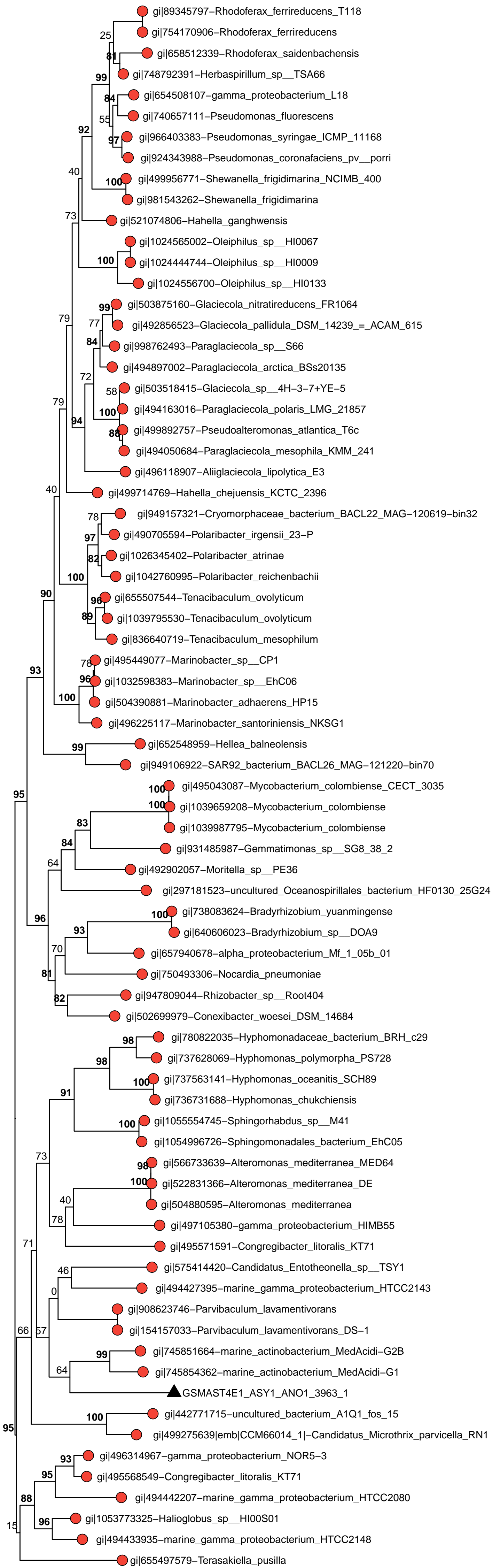


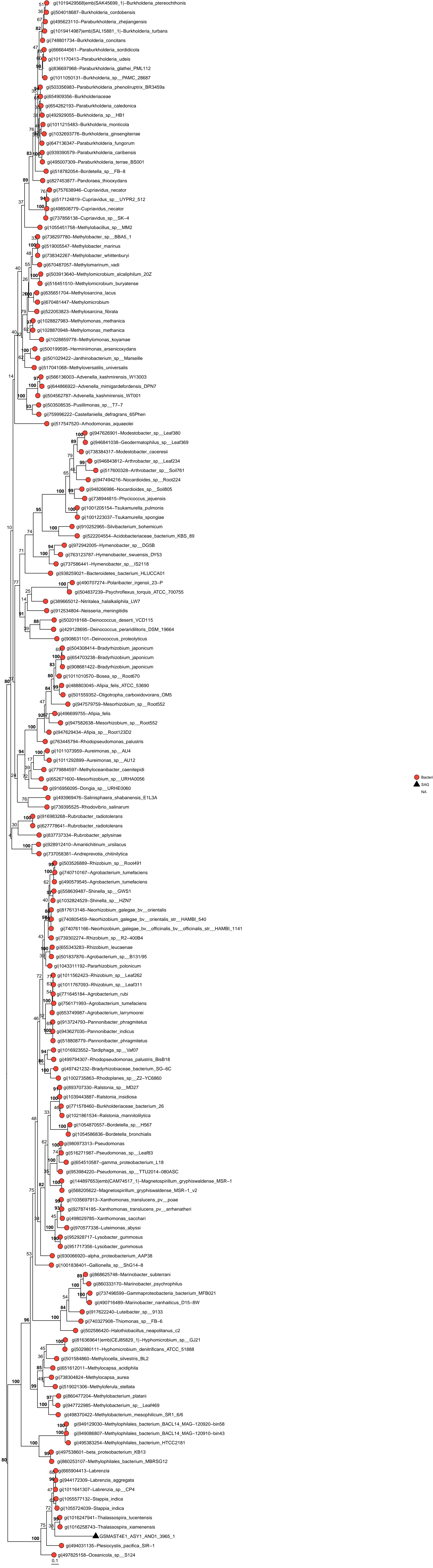
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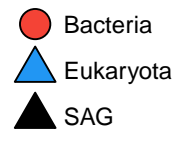
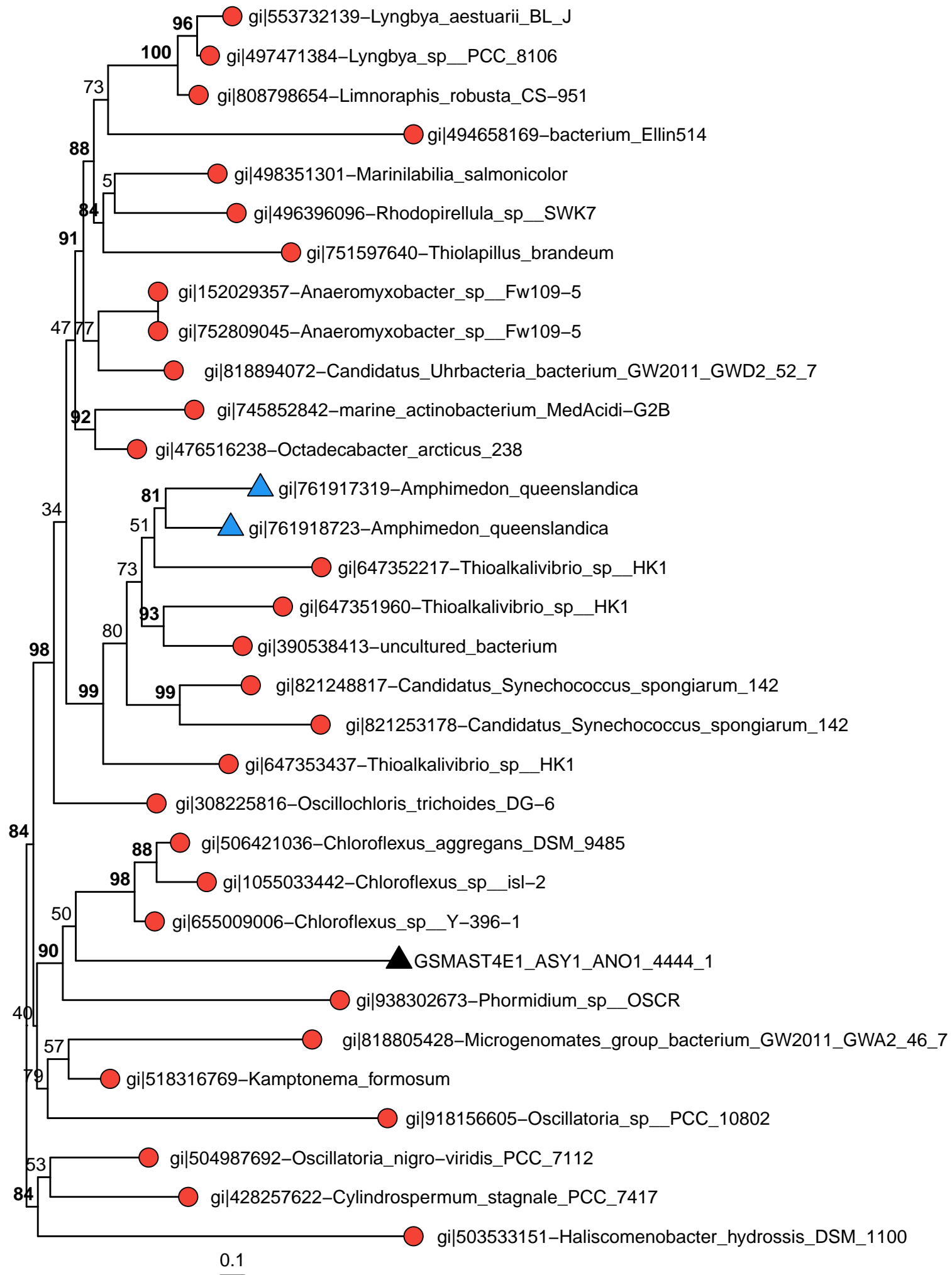


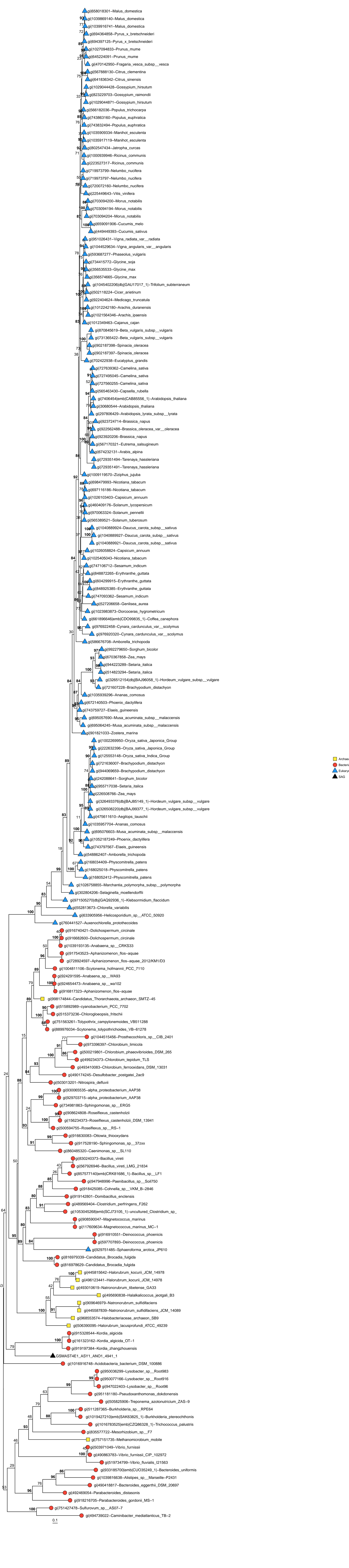
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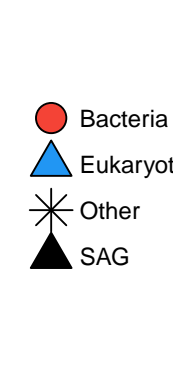
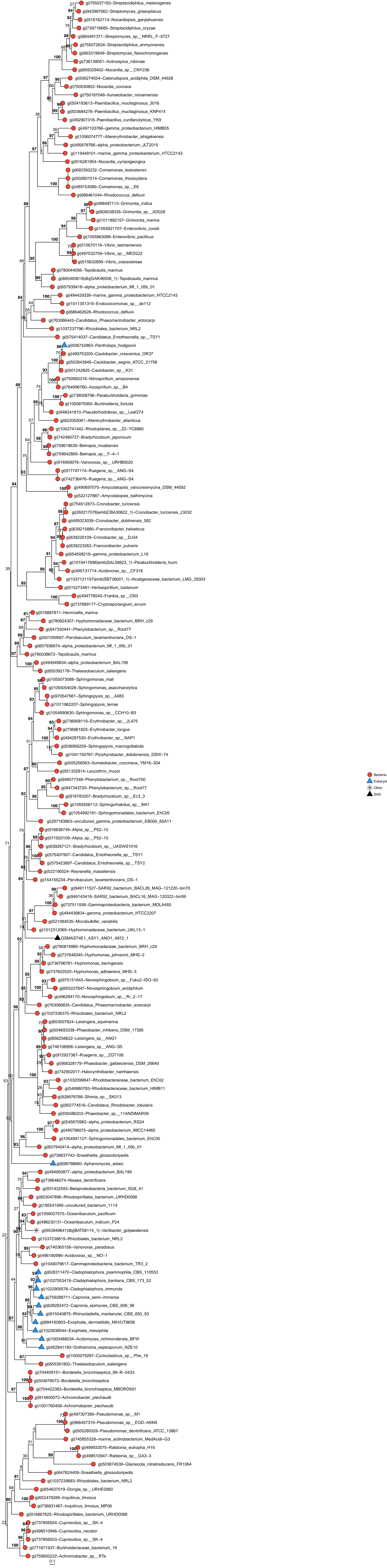


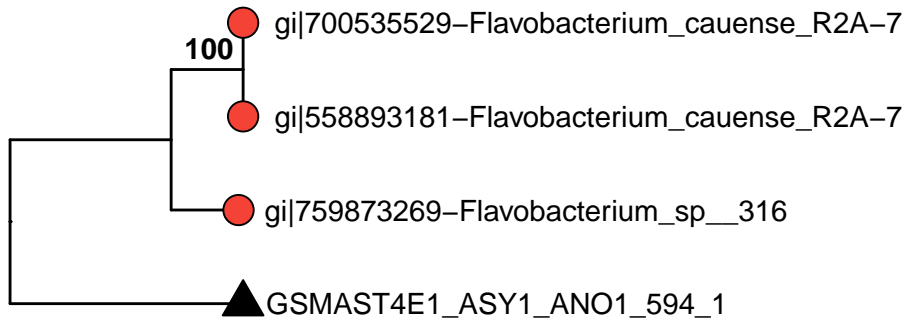




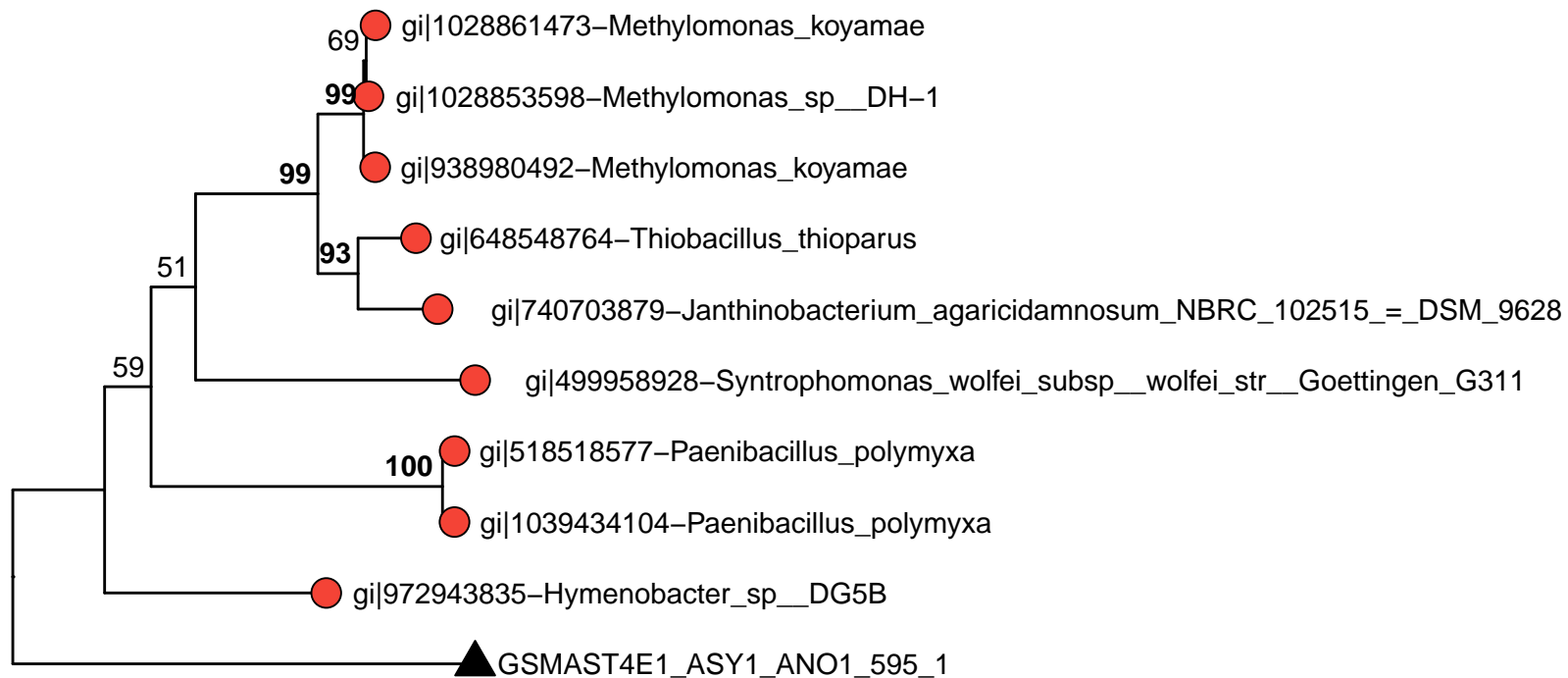






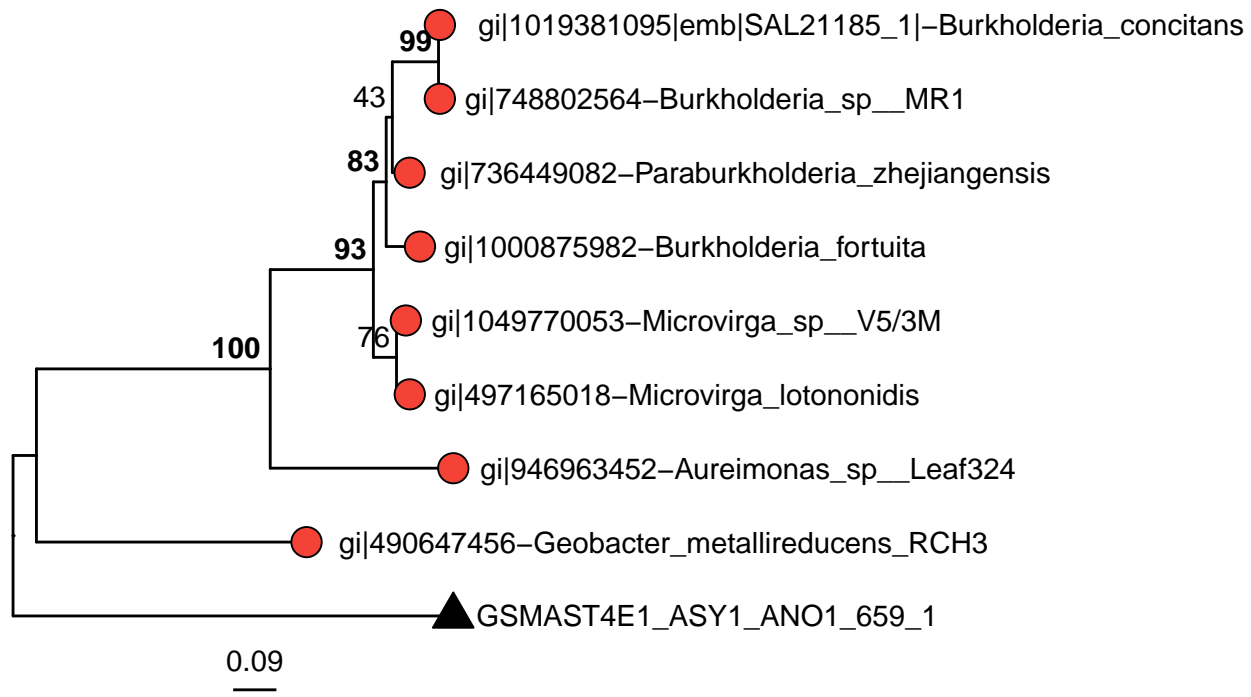


● Bacteria
▲ SAG

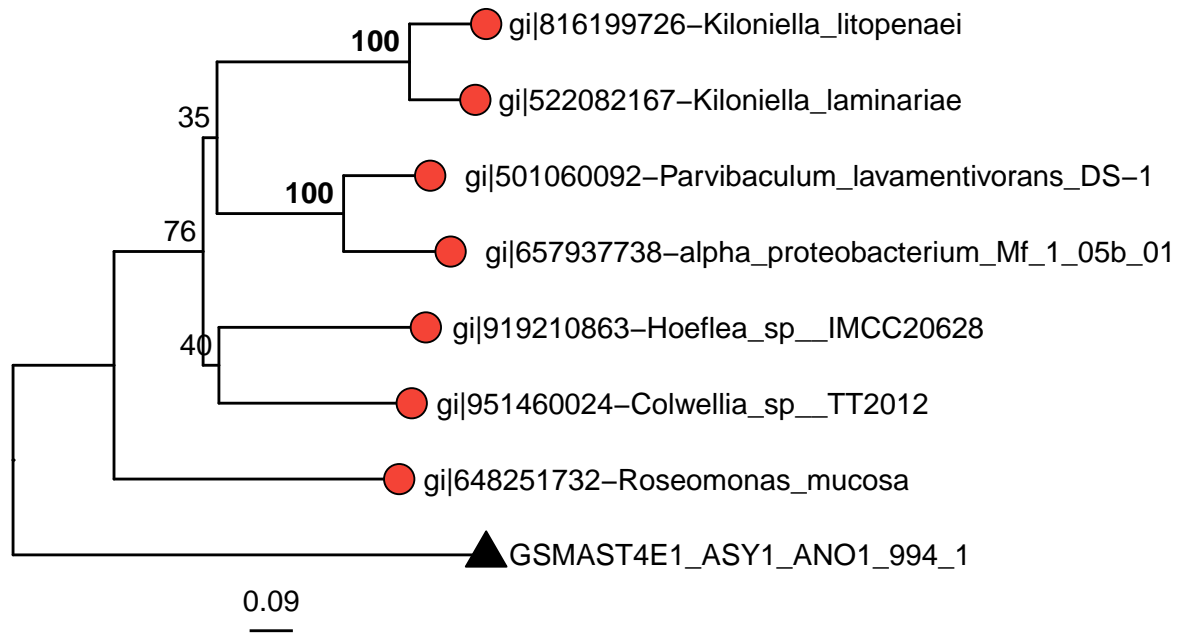


● Bacteria
▲ SAG

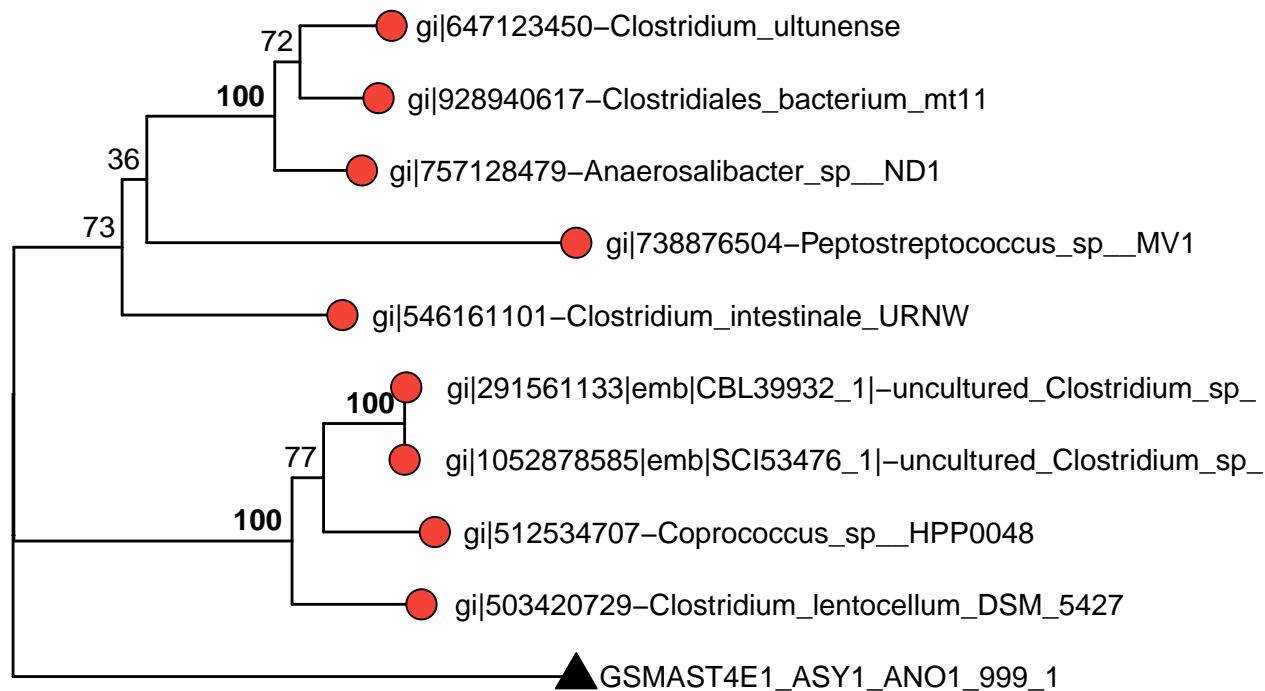
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● Bacteria
▲ SAG



● Bacteria
▲ SAG



● Bacteria
▲ SAG