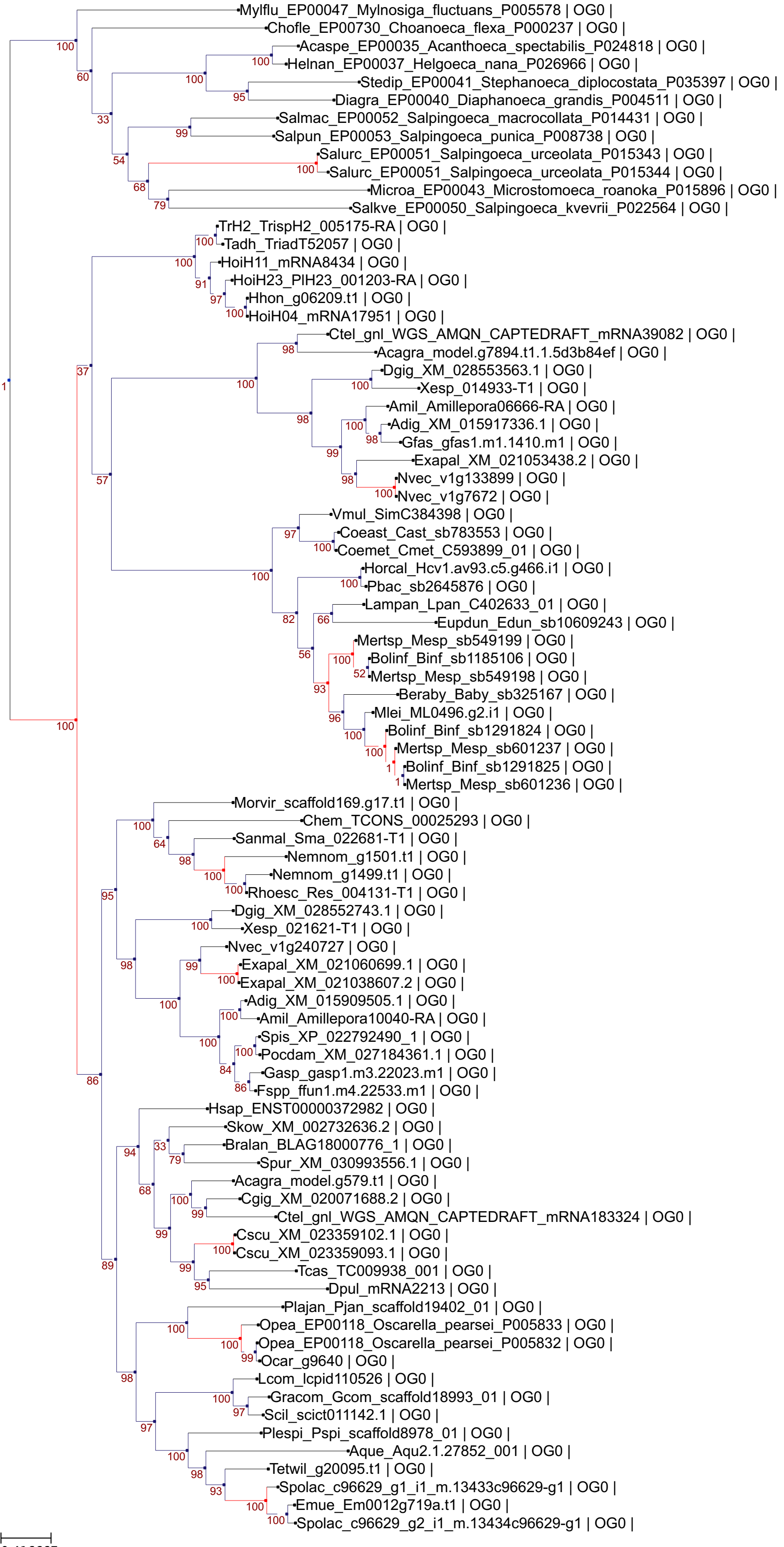
Mylflu_EP00047_Mylnosiga_fluctuans_P005578 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P000237 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P024818 | OG0 |
Helnan_EP00037_Helgoeca_nana_P026966 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P035397 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P004511 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P014431 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P008738 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P015343 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P015344 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P015896 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P022564 | OG0 |
TrH2_TrispH2_005175-RA | OG0 |
Tadh_TriadT52057 | OG0 |
HoiH11_mRNA8434 | OG0 |
HoiH23_PlH23_001203-RA | OG0 |
Hhon_g06209.t1 | OG0 |
HoiH04_mRNA17951 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA39082 | OG0 |
Acagra_model.g7894.t1.1.5d3b84ef | OG0 |
Dgig_XM_028553563.1 | OG0 |
Xesp_014933-T1 | OG0 |
Amil_Amillepora06666-RA | OG0 |
Adig_XM_015917336.1 | OG0 |
Gfas_gfas1.m1.1410.m1 | OG0 |
Exapal_XM_021053438.2 | OG0 |
Nvec_v1g133899 | OG0 |
Nvec_v1g7672 | OG0 |
Vmul_SimC384398 | OG0 |
Coeast_Cast_sb783553 | OG0 |
Coemet_Cmet_C593899_01 | OG0 |
Horcal_Hcv1.av93.c5.g466.i1 | OG0 |
Pbac_sb2645876 | OG0 |
Lampan_Lpan_C402633_01 | OG0 |
Eupdun_Edun_sb10609243 | OG0 |
Mertsp_Mesp_sb549199 | OG0 |
Bolinf_Binf_sb1185106 | OG0 |
Mertsp_Mesp_sb549198 | OG0 |
Beraby_Baby_sb325167 | OG0 |
Mlei_ML0496.g2.i1 | OG0 |
Bolinf_Binf_sb1291824 | OG0 |
Mertsp_Mesp_sb601237 | OG0 |
Bolinf_Binf_sb1291825 | OG0 |
Mertsp_Mesp_sb601236 | OG0 |
Morvir_scaffold169.g17.t1 | OG0 |
Chem_TCONS_00025293 | OG0 |
Sanmal_Sma_022681-T1 | OG0 |
Nemnom_g1501.t1 | OG0 |
Nemnom_g1499.t1 | OG0 |
Rhoesc_Res_004131-T1 | OG0 |
Dgig_XM_028552743.1 | OG0 |
Xesp_021621-T1 | OG0 |
Nvec_v1g240727 | OG0 |
Exapal_XM_021060699.1 | OG0 |
Exapal_XM_021038607.2 | OG0 |
Adig_XM_015909505.1 | OG0 |
Amil_Amillepora10040-RA | OG0 |
Spis_XP_022792490_1 | OG0 |
Pocdam_XM_027184361.1 | OG0 |
Gasp_gasp1.m3.22023.m1 | OG0 |
Fspp_ffun1.m4.22533.m1 | OG0 |
Hsap_ENST00000372982 | OG0 |
Skow_XM_002732636.2 | OG0 |
Bralan_BLAG18000776_1 | OG0 |
Spur_XM_030993556.1 | OG0 |
Acagra_model.g579.t1 | OG0 |
Cgig_XM_020071688.2 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA183324 | OG0 |
Cscu_XM_023359102.1 | OG0 |
Cscu_XM_023359093.1 | OG0 |
Tcas_TC009938_001 | OG0 |
Dpul_mRNA2213 | OG0 |
Plajan_Pjan_scaffold19402_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P005833 | OG0 |
Opea_EP00118_Oscarella_pearsei_P005832 | OG0 |
Ocar_g9640 | OG0 |
Lcom_lcpid110526 | OG0 |
Gracom_Gcom_scaffold18993_01 | OG0 |
Scil_scict011142.1 | OG0 |
Plespi_Pspi_scaffold8978_01 | OG0 |
Aque_Aqu2.1.27852_001 | OG0 |
Tetwil_g20095.t1 | OG0 |
Spolac_c96629_g1_i1_m.13433c96629-g1 | OG0 |
Emue_Em0012g719a.t1 | OG0 |
Spolac_c96629_g2_i1_m.13434c96629-g1 | OG0 |
0.416667