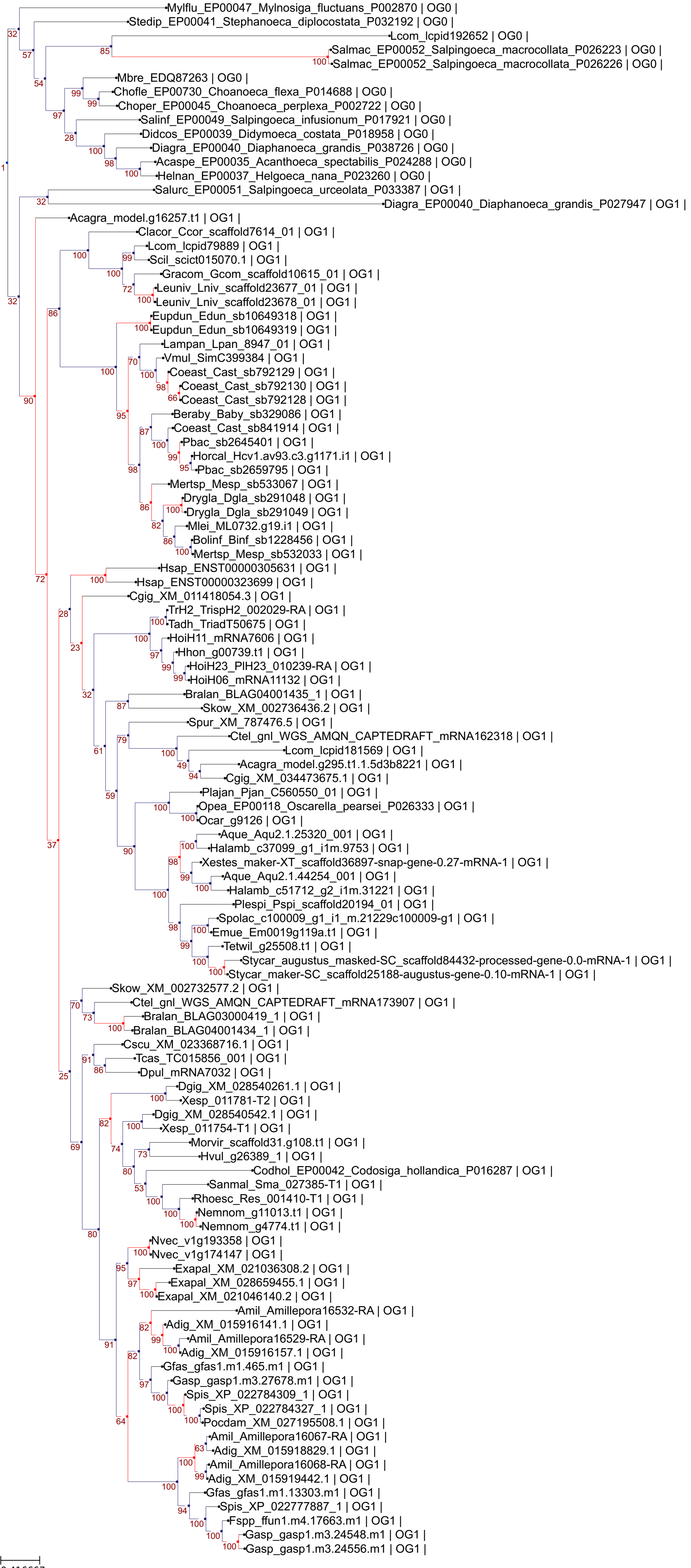
Mylflu_EP00047_Mylnosiga_fluctuans_P002870 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P032192 | OG0 |
Lcom_lcpid192652 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P026223 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P026226 | OG0 |
Mbre_EDQ87263 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P014688 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P002722 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P017921 | OG0 |
Didcos_EP00039_Didymoeca_costata_P018958 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P038726 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P024288 | OG0 |
Helnan_EP00037_Helgoeca_nana_P023260 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P033387 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P027947 | OG1 |
Acagra_model.g16257.t1 | OG1 |
Clacor_Ccor_scaffold7614_01 | OG1 |
Lcom_lcpid79889 | OG1 |
Scil_scict015070.1 | OG1 |
Gracom_Gcom_scaffold10615_01 | OG1 |
Leuniv_Lniv_scaffold23677_01 | OG1 |
Leuniv_Lniv_scaffold23678_01 | OG1 |
Eupdun_Edun_sb10649318 | OG1 |
Eupdun_Edun_sb10649319 | OG1 |
Lampan_Lpan_8947_01 | OG1 |
Vmul_SimC399384 | OG1 |
Coeast_Cast_sb792129 | OG1 |
Coeast_Cast_sb792130 | OG1 |
Coeast_Cast_sb792128 | OG1 |
Beraby_Baby_sb329086 | OG1 |
Coeast_Cast_sb841914 | OG1 |
Pbac_sb2645401 | OG1 |
Horcal_Hcv1.av93.c3.g1171.i1 | OG1 |
Pbac_sb2659795 | OG1 |
Mertsp_Mesp_sb533067 | OG1 |
Drygla_Dgla_sb291048 | OG1 |
Drygla_Dgla_sb291049 | OG1 |
Mlei_ML0732.g19.i1 | OG1 |
Bolinf_Binf_sb1228456 | OG1 |
Mertsp_Mesp_sb532033 | OG1 |
Hsap_ENST00000305631 | OG1 |
Hsap_ENST00000323699 | OG1 |
Cgig_XM_011418054.3 | OG1 |
TrH2_TrispH2_002029-RA | OG1 |
Tadh_TriadT50675 | OG1 |
HoiH11_mRNA7606 | OG1 |
Hhon_g00739.t1 | OG1 |
HoiH23_PlH23_010239-RA | OG1 |
HoiH06_mRNA11132 | OG1 |
Bralan_BLAG04001435_1 | OG1 |
Skow_XM_002736436.2 | OG1 |
Spur_XM_787476.5 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA162318 | OG1 |
Lcom_lcpid181569 | OG1 |
Acagra_model.g295.t1.1.5d3b8221 | OG1 |
Cgig_XM_034473675.1 | OG1 |
Plajan_Pjan_C560550_01 | OG1 |
Opea_EP00118_Oscarella_pearsei_P026333 | OG1 |
Ocar_g9126 | OG1 |
Aque_Aqu2.1.25320_001 | OG1 |
Halamb_c37099_g1_i1m.9753 | OG1 |
Xestes_maker-XT_scaffold36897-snap-gene-0.27-mRNA-1 | OG1 |
Aque_Aqu2.1.44254_001 | OG1 |
Halamb_c51712_g2_i1m.31221 | OG1 |
Plespi_Pspi_scaffold20194_01 | OG1 |
Spolac_c100009_g1_i1_m.21229c100009-g1 | OG1 |
Emue_Em0019g119a.t1 | OG1 |
Tetwil_g25508.t1 | OG1 |
Stycar_augustus_masked-SC_scaffold84432-processed-gene-0.0-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold25188-augustus-gene-0.10-mRNA-1 | OG1 |
Skow_XM_002732577.2 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA173907 | OG1 |
Bralan_BLAG03000419_1 | OG1 |
Bralan_BLAG04001434_1 | OG1 |
Cscu_XM_023368716.1 | OG1 |
Tcas_TC015856_001 | OG1 |
Dpul_mRNA7032 | OG1 |
Dgig_XM_028540261.1 | OG1 |
Xesp_011781-T2 | OG1 |
Dgig_XM_028540542.1 | OG1 |
Xesp_011754-T1 | OG1 |
Morvir_scaffold31.g108.t1 | OG1 |
Hvul_g26389_1 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P016287 | OG1 |
Sanmal_Sma_027385-T1 | OG1 |
Rhoesc_Res_001410-T1 | OG1 |
Nemnom_g11013.t1 | OG1 |
Nemnom_g4774.t1 | OG1 |
Nvec_v1g193358 | OG1 |
Nvec_v1g174147 | OG1 |
Exapal_XM_021036308.2 | OG1 |
Exapal_XM_028659455.1 | OG1 |
Exapal_XM_021046140.2 | OG1 |
Amil_Amillepora16532-RA | OG1 |
Adig_XM_015916141.1 | OG1 |
Amil_Amillepora16529-RA | OG1 |
Adig_XM_015916157.1 | OG1 |
Gfas_gfas1.m1.465.m1 | OG1 |
Gasp_gasp1.m3.27678.m1 | OG1 |
Spis_XP_022784309_1 | OG1 |
Spis_XP_022784327_1 | OG1 |
Pocdam_XM_027195508.1 | OG1 |
Amil_Amillepora16067-RA | OG1 |
Adig_XM_015918829.1 | OG1 |
Amil_Amillepora16068-RA | OG1 |
Adig_XM_015919442.1 | OG1 |
Gfas_gfas1.m1.13303.m1 | OG1 |
Spis_XP_022777887_1 | OG1 |
Fspp_ffun1.m4.17663.m1 | OG1 |
Gasp_gasp1.m3.24548.m1 | OG1 |
Gasp_gasp1.m3.24556.m1 | OG1 |
0.416667