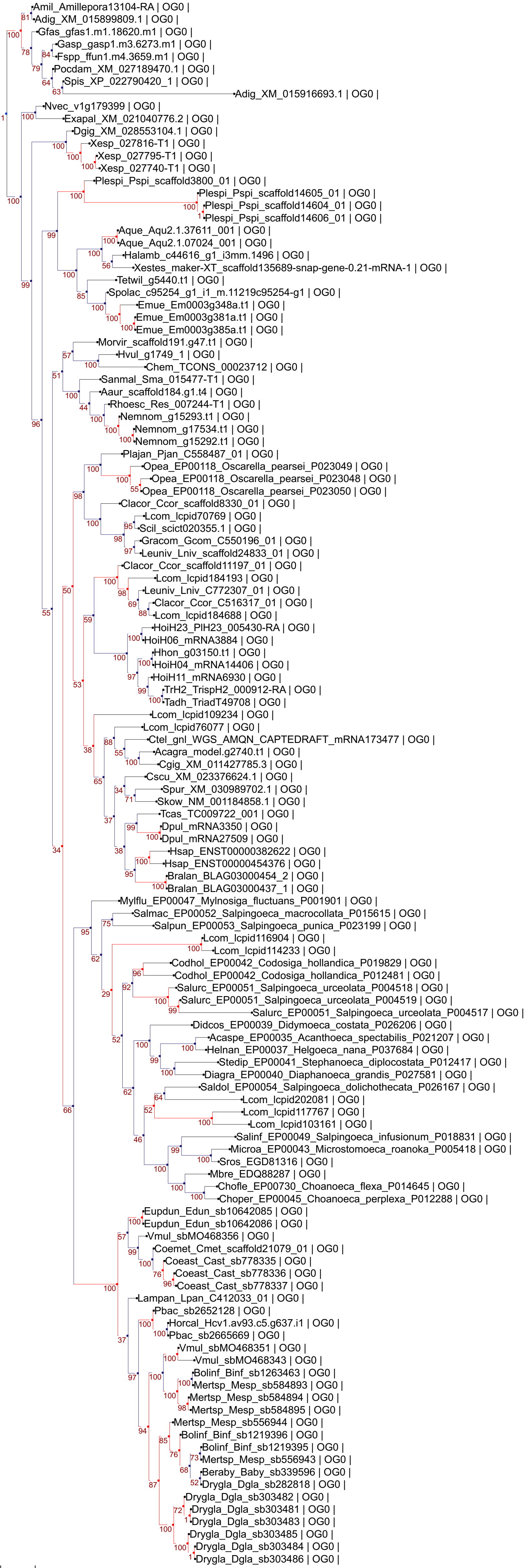
Amil_Amillepora13104-RA | OG0 |
Adig_XM_015899809.1 | OG0 |
Gfas_gfas1.m1.18620.m1 | OG0 |
Gasp_gasp1.m3.6273.m1 | OG0 |
Fspp_ffun1.m4.3659.m1 | OG0 |
Pocdam_XM_027189470.1 | OG0 |
Spis_XP_022790420_1 | OG0 |
Adig_XM_015916693.1 | OG0 |
Nvec_v1g179399 | OG0 |
Exapal_XM_021040776.2 | OG0 |
Dgig_XM_028553104.1 | OG0 |
Xesp_027816-T1 | OG0 |
Xesp_027795-T1 | OG0 |
Xesp_027740-T1 | OG0 |
Plespi_Pspi_scaffold3800_01 | OG0 |
Plespi_Pspi_scaffold14605_01 | OG0 |
Plespi_Pspi_scaffold14604_01 | OG0 |
Plespi_Pspi_scaffold14606_01 | OG0 |
Aque_Aqu2.1.37611_001 | OG0 |
Aque_Aqu2.1.07024_001 | OG0 |
Halamb_c44616_g1_i3mm.1496 | OG0 |
Xestes_maker-XT_scaffold135689-snap-gene-0.21-mRNA-1 | OG0 |
Tetwil_g5440.t1 | OG0 |
Spolac_c95254_g1_i1_m.11219c95254-g1 | OG0 |
Emue_Em0003g348a.t1 | OG0 |
Emue_Em0003g381a.t1 | OG0 |
Emue_Em0003g385a.t1 | OG0 |
Morvir_scaffold191.g47.t1 | OG0 |
Hvul_g1749_1 | OG0 |
Chem_TCONS_00023712 | OG0 |
Sanmal_Sma_015477-T1 | OG0 |
Aaur_scaffold184.g1.t4 | OG0 |
Rhoesc_Res_007244-T1 | OG0 |
Nemnom_g15293.t1 | OG0 |
Nemnom_g17534.t1 | OG0 |
Nemnom_g15292.t1 | OG0 |
Plajan_Pjan_C558487_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P023049 | OG0 |
Opea_EP00118_Oscarella_pearsei_P023048 | OG0 |
Opea_EP00118_Oscarella_pearsei_P023050 | OG0 |
Clacor_Ccor_scaffold8330_01 | OG0 |
Lcom_lcpid70769 | OG0 |
Scil_scict020355.1 | OG0 |
Gracom_Gcom_C550196_01 | OG0 |
Leuniv_Lniv_scaffold24833_01 | OG0 |
Clacor_Ccor_scaffold11197_01 | OG0 |
Lcom_lcpid184193 | OG0 |
Leuniv_Lniv_C772307_01 | OG0 |
Clacor_Ccor_C516317_01 | OG0 |
Lcom_lcpid184688 | OG0 |
HoiH23_PlH23_005430-RA | OG0 |
HoiH06_mRNA3884 | OG0 |
Hhon_g03150.t1 | OG0 |
HoiH04_mRNA14406 | OG0 |
HoiH11_mRNA6930 | OG0 |
TrH2_TrispH2_000912-RA | OG0 |
Tadh_TriadT49708 | OG0 |
Lcom_lcpid109234 | OG0 |
Lcom_lcpid76077 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA173477 | OG0 |
Acagra_model.g2740.t1 | OG0 |
Cgig_XM_011427785.3 | OG0 |
Cscu_XM_023376624.1 | OG0 |
Spur_XM_030989702.1 | OG0 |
Skow_NM_001184858.1 | OG0 |
Tcas_TC009722_001 | OG0 |
Dpul_mRNA3350 | OG0 |
Dpul_mRNA27509 | OG0 |
Hsap_ENST00000382622 | OG0 |
Hsap_ENST00000454376 | OG0 |
Bralan_BLAG03000454_2 | OG0 |
Bralan_BLAG03000437_1 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P001901 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P015615 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P023199 | OG0 |
Lcom_lcpid116904 | OG0 |
Lcom_lcpid114233 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P019829 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P012481 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P004518 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P004519 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P004517 | OG0 |
Didcos_EP00039_Didymoeca_costata_P026206 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P021207 | OG0 |
Helnan_EP00037_Helgoeca_nana_P037684 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P012417 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P027581 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P026167 | OG0 |
Lcom_lcpid202081 | OG0 |
Lcom_lcpid117767 | OG0 |
Lcom_lcpid103161 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P018831 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P005418 | OG0 |
Sros_EGD81316 | OG0 |
Mbre_EDQ88287 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P014645 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P012288 | OG0 |
Eupdun_Edun_sb10642085 | OG0 |
Eupdun_Edun_sb10642086 | OG0 |
Vmul_sbMO468356 | OG0 |
Coemet_Cmet_scaffold21079_01 | OG0 |
Coeast_Cast_sb778335 | OG0 |
Coeast_Cast_sb778336 | OG0 |
Coeast_Cast_sb778337 | OG0 |
Lampan_Lpan_C412033_01 | OG0 |
Pbac_sb2652128 | OG0 |
Horcal_Hcv1.av93.c5.g637.i1 | OG0 |
Pbac_sb2665669 | OG0 |
Vmul_sbMO468351 | OG0 |
Vmul_sbMO468343 | OG0 |
Bolinf_Binf_sb1263463 | OG0 |
Mertsp_Mesp_sb584893 | OG0 |
Mertsp_Mesp_sb584894 | OG0 |
Mertsp_Mesp_sb584895 | OG0 |
Mertsp_Mesp_sb556944 | OG0 |
Bolinf_Binf_sb1219396 | OG0 |
Bolinf_Binf_sb1219395 | OG0 |
Mertsp_Mesp_sb556943 | OG0 |
Beraby_Baby_sb339596 | OG0 |
Drygla_Dgla_sb282818 | OG0 |
Drygla_Dgla_sb303482 | OG0 |
Drygla_Dgla_sb303481 | OG0 |
Drygla_Dgla_sb303483 | OG0 |
Drygla_Dgla_sb303485 | OG0 |
Drygla_Dgla_sb303484 | OG0 |
Drygla_Dgla_sb303486 | OG0 |
0.416667