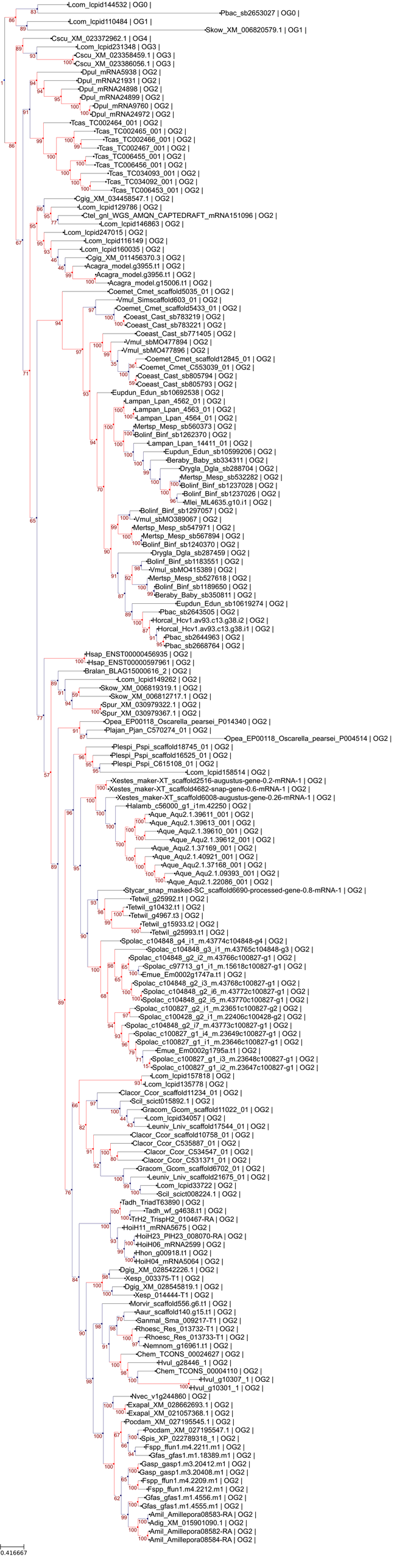
Lcom_lcpid144532 | OG0 |
Pbac_sb2653027 | OG0 |
Lcom_lcpid110484 | OG1 |
Skow_XM_006820579.1 | OG1 |
Cscu_XM_023372962.1 | OG4 |
Lcom_lcpid231348 | OG3 |
Cscu_XM_023358459.1 | OG3 |
Cscu_XM_023386056.1 | OG3 |
Dpul_mRNA5938 | OG2 |
Dpul_mRNA21931 | OG2 |
Dpul_mRNA24898 | OG2 |
Dpul_mRNA24899 | OG2 |
Dpul_mRNA9760 | OG2 |
Dpul_mRNA24972 | OG2 |
Tcas_TC002464_001 | OG2 |
Tcas_TC002465_001 | OG2 |
Tcas_TC002466_001 | OG2 |
Tcas_TC002467_001 | OG2 |
Tcas_TC006455_001 | OG2 |
Tcas_TC006456_001 | OG2 |
Tcas_TC034093_001 | OG2 |
Tcas_TC034092_001 | OG2 |
Tcas_TC006453_001 | OG2 |
Cgig_XM_034458547.1 | OG2 |
Lcom_lcpid129786 | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA151096 | OG2 |
Lcom_lcpid146863 | OG2 |
Lcom_lcpid247015 | OG2 |
Lcom_lcpid116149 | OG2 |
Lcom_lcpid160035 | OG2 |
Cgig_XM_011456370.3 | OG2 |
Acagra_model.g3955.t1 | OG2 |
Acagra_model.g3956.t1 | OG2 |
Acagra_model.g15006.t1 | OG2 |
Coemet_Cmet_scaffold5035_01 | OG2 |
Vmul_Simscaffold603_01 | OG2 |
Coemet_Cmet_scaffold5433_01 | OG2 |
Coeast_Cast_sb783219 | OG2 |
Coeast_Cast_sb783221 | OG2 |
Coeast_Cast_sb771405 | OG2 |
Vmul_sbMO477894 | OG2 |
Vmul_sbMO477896 | OG2 |
Coemet_Cmet_scaffold12845_01 | OG2 |
Coemet_Cmet_C553039_01 | OG2 |
Coeast_Cast_sb805794 | OG2 |
Coeast_Cast_sb805793 | OG2 |
Eupdun_Edun_sb10692538 | OG2 |
Lampan_Lpan_4562_01 | OG2 |
Lampan_Lpan_4563_01 | OG2 |
Lampan_Lpan_4564_01 | OG2 |
Mertsp_Mesp_sb560373 | OG2 |
Bolinf_Binf_sb1262370 | OG2 |
Lampan_Lpan_14411_01 | OG2 |
Eupdun_Edun_sb10599206 | OG2 |
Beraby_Baby_sb334311 | OG2 |
Drygla_Dgla_sb288704 | OG2 |
Mertsp_Mesp_sb532282 | OG2 |
Bolinf_Binf_sb1237028 | OG2 |
Bolinf_Binf_sb1237026 | OG2 |
Mlei_ML4635.g10.i1 | OG2 |
Bolinf_Binf_sb1297057 | OG2 |
Vmul_sbMO389067 | OG2 |
Mertsp_Mesp_sb547971 | OG2 |
Mertsp_Mesp_sb567894 | OG2 |
Bolinf_Binf_sb1240370 | OG2 |
Drygla_Dgla_sb287459 | OG2 |
Bolinf_Binf_sb1183551 | OG2 |
Vmul_sbMO415389 | OG2 |
Mertsp_Mesp_sb527618 | OG2 |
Bolinf_Binf_sb1189650 | OG2 |
Beraby_Baby_sb350811 | OG2 |
Eupdun_Edun_sb10619274 | OG2 |
Pbac_sb2643505 | OG2 |
Horcal_Hcv1.av93.c13.g38.i2 | OG2 |
Horcal_Hcv1.av93.c13.g38.i1 | OG2 |
Pbac_sb2644963 | OG2 |
Pbac_sb2668764 | OG2 |
Hsap_ENST00000456935 | OG2 |
Hsap_ENST00000597961 | OG2 |
Bralan_BLAG15000616_2 | OG2 |
Lcom_lcpid149262 | OG2 |
Skow_XM_006819319.1 | OG2 |
Skow_XM_006812717.1 | OG2 |
Spur_XM_030979322.1 | OG2 |
Spur_XM_030979367.1 | OG2 |
Opea_EP00118_Oscarella_pearsei_P014340 | OG2 |
Plajan_Pjan_C570274_01 | OG2 |
Opea_EP00118_Oscarella_pearsei_P004514 | OG2 |
Plespi_Pspi_scaffold18745_01 | OG2 |
Plespi_Pspi_scaffold16525_01 | OG2 |
Plespi_Pspi_C615108_01 | OG2 |
Lcom_lcpid158514 | OG2 |
Xestes_maker-XT_scaffold2516-augustus-gene-0.2-mRNA-1 | OG2 |
Xestes_maker-XT_scaffold4682-snap-gene-0.6-mRNA-1 | OG2 |
Xestes_maker-XT_scaffold6008-augustus-gene-0.26-mRNA-1 | OG2 |
Halamb_c56000_g1_i1m.42250 | OG2 |
Aque_Aqu2.1.39611_001 | OG2 |
Aque_Aqu2.1.39613_001 | OG2 |
Aque_Aqu2.1.39610_001 | OG2 |
Aque_Aqu2.1.39612_001 | OG2 |
Aque_Aqu2.1.37169_001 | OG2 |
Aque_Aqu2.1.40921_001 | OG2 |
Aque_Aqu2.1.37168_001 | OG2 |
Aque_Aqu2.1.09393_001 | OG2 |
Aque_Aqu2.1.22086_001 | OG2 |
Stycar_snap_masked-SC_scaffold6690-processed-gene-0.8-mRNA-1 | OG2 |
Tetwil_g25992.t1 | OG2 |
Tetwil_g10432.t1 | OG2 |
Tetwil_g4967.t3 | OG2 |
Tetwil_g15933.t2 | OG2 |
Tetwil_g25993.t1 | OG2 |
Spolac_c104848_g4_i1_m.43774c104848-g4 | OG2 |
Spolac_c104848_g3_i1_m.43765c104848-g3 | OG2 |
Spolac_c104848_g2_i2_m.43766c100827-g1 | OG2 |
Spolac_c97713_g1_i1_m.15618c100827-g1 | OG2 |
Emue_Em0002g1747a.t1 | OG2 |
Spolac_c104848_g2_i3_m.43768c100827-g1 | OG2 |
Spolac_c104848_g2_i6_m.43772c100827-g1 | OG2 |
Spolac_c104848_g2_i5_m.43770c100827-g1 | OG2 |
Spolac_c100827_g2_i1_m.23651c100827-g2 | OG2 |
Spolac_c100428_g2_i1_m.22406c100428-g2 | OG2 |
Spolac_c104848_g2_i7_m.43773c100827-g1 | OG2 |
Spolac_c100827_g1_i4_m.23649c100827-g1 | OG2 |
Spolac_c100827_g1_i1_m.23646c100827-g1 | OG2 |
Emue_Em0002g1795a.t1 | OG2 |
Spolac_c100827_g1_i3_m.23648c100827-g1 | OG2 |
Spolac_c100827_g1_i2_m.23647c100827-g1 | OG2 |
Lcom_lcpid157818 | OG2 |
Lcom_lcpid135778 | OG2 |
Clacor_Ccor_scaffold11234_01 | OG2 |
Scil_scict015892.1 | OG2 |
Gracom_Gcom_scaffold11022_01 | OG2 |
Lcom_lcpid34057 | OG2 |
Leuniv_Lniv_scaffold17544_01 | OG2 |
Clacor_Ccor_scaffold10758_01 | OG2 |
Clacor_Ccor_C535887_01 | OG2 |
Clacor_Ccor_C534547_01 | OG2 |
Clacor_Ccor_C531371_01 | OG2 |
Gracom_Gcom_scaffold6702_01 | OG2 |
Leuniv_Lniv_scaffold21675_01 | OG2 |
Lcom_lcpid33722 | OG2 |
Scil_scict008224.1 | OG2 |
Tadh_TriadT63890 | OG2 |
Tadh_wf_g4638.t1 | OG2 |
TrH2_TrispH2_010467-RA | OG2 |
HoiH11_mRNA5675 | OG2 |
HoiH23_PlH23_008070-RA | OG2 |
HoiH06_mRNA2599 | OG2 |
Hhon_g00918.t1 | OG2 |
HoiH04_mRNA5064 | OG2 |
Dgig_XM_028542226.1 | OG2 |
Xesp_003375-T1 | OG2 |
Dgig_XM_028545819.1 | OG2 |
Xesp_014444-T1 | OG2 |
Morvir_scaffold556.g6.t1 | OG2 |
Aaur_scaffold140.g15.t1 | OG2 |
Sanmal_Sma_009217-T1 | OG2 |
Rhoesc_Res_013732-T1 | OG2 |
Rhoesc_Res_013733-T1 | OG2 |
Nemnom_g16961.t1 | OG2 |
Chem_TCONS_00024627 | OG2 |
Hvul_g28446_1 | OG2 |
Chem_TCONS_00004110 | OG2 |
Hvul_g10307_1 | OG2 |
Hvul_g10301_1 | OG2 |
Nvec_v1g244860 | OG2 |
Exapal_XM_028662693.1 | OG2 |
Exapal_XM_021057368.1 | OG2 |
Pocdam_XM_027195545.1 | OG2 |
Pocdam_XM_027195547.1 | OG2 |
Spis_XP_022789318_1 | OG2 |
Fspp_ffun1.m4.2211.m1 | OG2 |
Gfas_gfas1.m1.18389.m1 | OG2 |
Gasp_gasp1.m3.20412.m1 | OG2 |
Gasp_gasp1.m3.20408.m1 | OG2 |
Fspp_ffun1.m4.2209.m1 | OG2 |
Fspp_ffun1.m4.2212.m1 | OG2 |
Gfas_gfas1.m1.4556.m1 | OG2 |
Gfas_gfas1.m1.4555.m1 | OG2 |
Amil_Amillepora08583-RA | OG2 |
Adig_XM_015901090.1 | OG2 |
Amil_Amillepora08582-RA | OG2 |
Amil_Amillepora08584-RA | OG2 |
0.416667