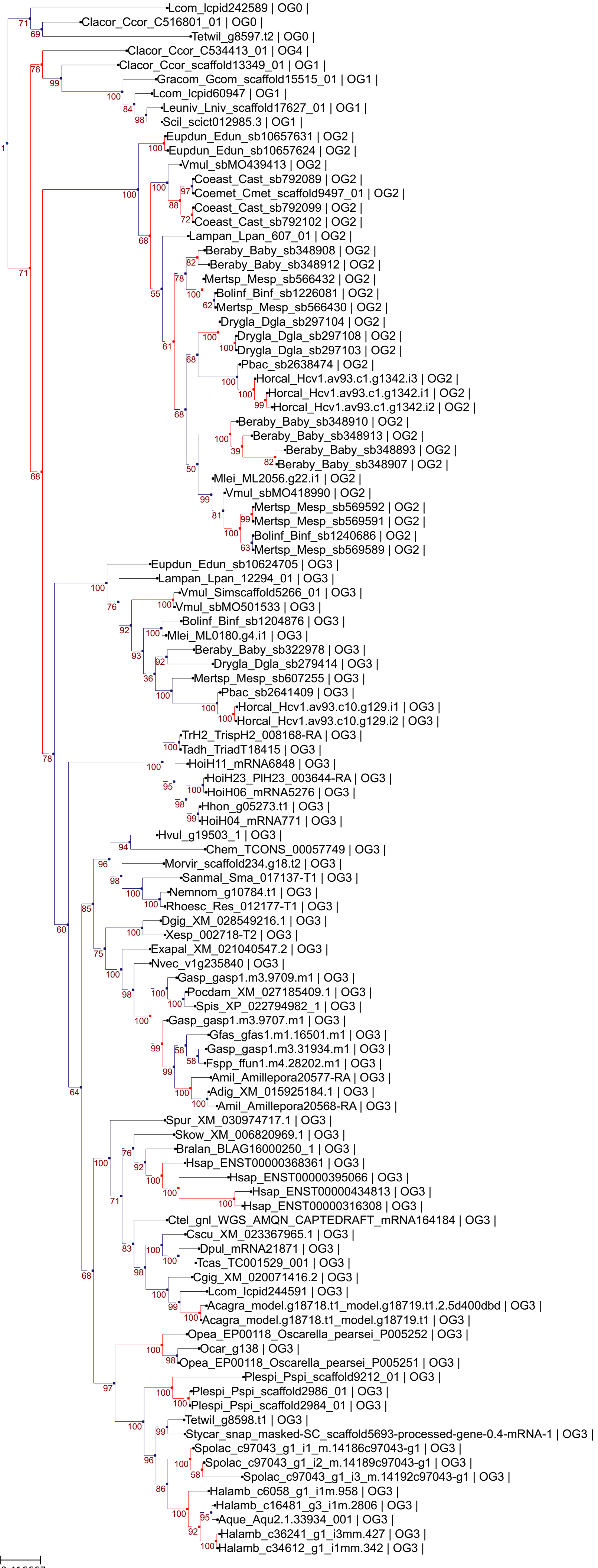
Lcom_lcpid242589 | OG0 |
Clacor_Ccor_C516801_01 | OG0 |
Tetwil_g8597.t2 | OG0 |
Clacor_Ccor_C534413_01 | OG4 |
Clacor_Ccor_scaffold13349_01 | OG1 |
Gracom_Gcom_scaffold15515_01 | OG1 |
Lcom_lcpid60947 | OG1 |
Leuniv_Lniv_scaffold17627_01 | OG1 |
Scil_scict012985.3 | OG1 |
Eupdun_Edun_sb10657631 | OG2 |
Eupdun_Edun_sb10657624 | OG2 |
Vmul_sbMO439413 | OG2 |
Coeast_Cast_sb792089 | OG2 |
Coemet_Cmet_scaffold9497_01 | OG2 |
Coeast_Cast_sb792099 | OG2 |
Coeast_Cast_sb792102 | OG2 |
Lampan_Lpan_607_01 | OG2 |
Beraby_Baby_sb348908 | OG2 |
Beraby_Baby_sb348912 | OG2 |
Mertsp_Mesp_sb566432 | OG2 |
Bolinf_Binf_sb1226081 | OG2 |
Mertsp_Mesp_sb566430 | OG2 |
Drygla_Dgla_sb297104 | OG2 |
Drygla_Dgla_sb297108 | OG2 |
Drygla_Dgla_sb297103 | OG2 |
Pbac_sb2638474 | OG2 |
Horcal_Hcv1.av93.c1.g1342.i3 | OG2 |
Horcal_Hcv1.av93.c1.g1342.i1 | OG2 |
Horcal_Hcv1.av93.c1.g1342.i2 | OG2 |
Beraby_Baby_sb348910 | OG2 |
Beraby_Baby_sb348913 | OG2 |
Beraby_Baby_sb348893 | OG2 |
Beraby_Baby_sb348907 | OG2 |
Mlei_ML2056.g22.i1 | OG2 |
Vmul_sbMO418990 | OG2 |
Mertsp_Mesp_sb569592 | OG2 |
Mertsp_Mesp_sb569591 | OG2 |
Bolinf_Binf_sb1240686 | OG2 |
Mertsp_Mesp_sb569589 | OG2 |
Eupdun_Edun_sb10624705 | OG3 |
Lampan_Lpan_12294_01 | OG3 |
Vmul_Simscaffold5266_01 | OG3 |
Vmul_sbMO501533 | OG3 |
Bolinf_Binf_sb1204876 | OG3 |
Mlei_ML0180.g4.i1 | OG3 |
Beraby_Baby_sb322978 | OG3 |
Drygla_Dgla_sb279414 | OG3 |
Mertsp_Mesp_sb607255 | OG3 |
Pbac_sb2641409 | OG3 |
Horcal_Hcv1.av93.c10.g129.i1 | OG3 |
Horcal_Hcv1.av93.c10.g129.i2 | OG3 |
TrH2_TrispH2_008168-RA | OG3 |
Tadh_TriadT18415 | OG3 |
HoiH11_mRNA6848 | OG3 |
HoiH23_PlH23_003644-RA | OG3 |
HoiH06_mRNA5276 | OG3 |
Hhon_g05273.t1 | OG3 |
HoiH04_mRNA771 | OG3 |
Hvul_g19503_1 | OG3 |
Chem_TCONS_00057749 | OG3 |
Morvir_scaffold234.g18.t2 | OG3 |
Sanmal_Sma_017137-T1 | OG3 |
Nemnom_g10784.t1 | OG3 |
Rhoesc_Res_012177-T1 | OG3 |
Dgig_XM_028549216.1 | OG3 |
Xesp_002718-T2 | OG3 |
Exapal_XM_021040547.2 | OG3 |
Nvec_v1g235840 | OG3 |
Gasp_gasp1.m3.9709.m1 | OG3 |
Pocdam_XM_027185409.1 | OG3 |
Spis_XP_022794982_1 | OG3 |
Gasp_gasp1.m3.9707.m1 | OG3 |
Gfas_gfas1.m1.16501.m1 | OG3 |
Gasp_gasp1.m3.31934.m1 | OG3 |
Fspp_ffun1.m4.28202.m1 | OG3 |
Amil_Amillepora20577-RA | OG3 |
Adig_XM_015925184.1 | OG3 |
Amil_Amillepora20568-RA | OG3 |
Spur_XM_030974717.1 | OG3 |
Skow_XM_006820969.1 | OG3 |
Bralan_BLAG16000250_1 | OG3 |
Hsap_ENST00000368361 | OG3 |
Hsap_ENST00000395066 | OG3 |
Hsap_ENST00000434813 | OG3 |
Hsap_ENST00000316308 | OG3 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA164184 | OG3 |
Cscu_XM_023367965.1 | OG3 |
Dpul_mRNA21871 | OG3 |
Tcas_TC001529_001 | OG3 |
Cgig_XM_020071416.2 | OG3 |
Lcom_lcpid244591 | OG3 |
Acagra_model.g18718.t1_model.g18719.t1.2.5d400dbd | OG3 |
Acagra_model.g18718.t1_model.g18719.t1 | OG3 |
Opea_EP00118_Oscarella_pearsei_P005252 | OG3 |
Ocar_g138 | OG3 |
Opea_EP00118_Oscarella_pearsei_P005251 | OG3 |
Plespi_Pspi_scaffold9212_01 | OG3 |
Plespi_Pspi_scaffold2986_01 | OG3 |
Plespi_Pspi_scaffold2984_01 | OG3 |
Tetwil_g8598.t1 | OG3 |
Stycar_snap_masked-SC_scaffold5693-processed-gene-0.4-mRNA-1 | OG3 |
Spolac_c97043_g1_i1_m.14186c97043-g1 | OG3 |
Spolac_c97043_g1_i2_m.14189c97043-g1 | OG3 |
Spolac_c97043_g1_i3_m.14192c97043-g1 | OG3 |
Halamb_c6058_g1_i1m.958 | OG3 |
Halamb_c16481_g3_i1m.2806 | OG3 |
Aque_Aqu2.1.33934_001 | OG3 |
Halamb_c36241_g1_i3mm.427 | OG3 |
Halamb_c34612_g1_i1mm.342 | OG3 |
0.416667