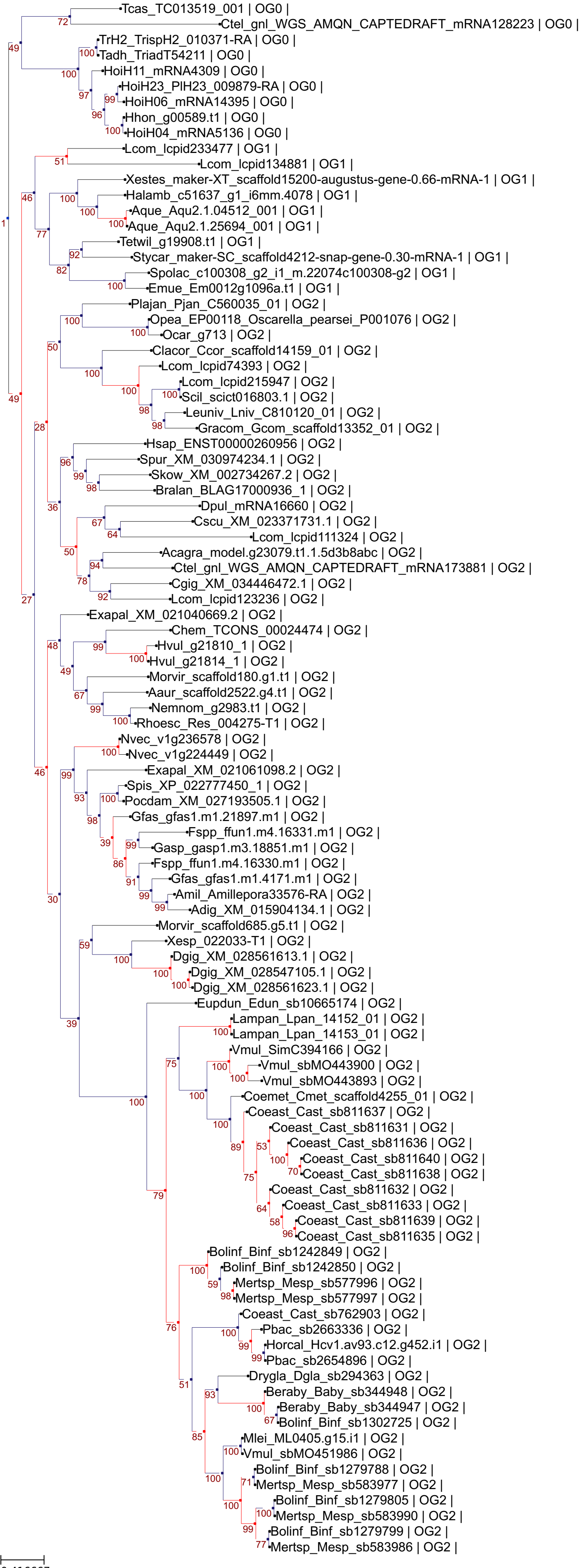
Tcas_TC013519_001 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA128223 | OG0 |
TrH2_TrispH2_010371-RA | OG0 |
Tadh_TriadT54211 | OG0 |
HoiH11_mRNA4309 | OG0 |
HoiH23_PlH23_009879-RA | OG0 |
HoiH06_mRNA14395 | OG0 |
Hhon_g00589.t1 | OG0 |
HoiH04_mRNA5136 | OG0 |
Lcom_lcpid233477 | OG1 |
Lcom_lcpid134881 | OG1 |
Xestes_maker-XT_scaffold15200-augustus-gene-0.66-mRNA-1 | OG1 |
Halamb_c51637_g1_i6mm.4078 | OG1 |
Aque_Aqu2.1.04512_001 | OG1 |
Aque_Aqu2.1.25694_001 | OG1 |
Tetwil_g19908.t1 | OG1 |
Stycar_maker-SC_scaffold4212-snap-gene-0.30-mRNA-1 | OG1 |
Spolac_c100308_g2_i1_m.22074c100308-g2 | OG1 |
Emue_Em0012g1096a.t1 | OG1 |
Plajan_Pjan_C560035_01 | OG2 |
Opea_EP00118_Oscarella_pearsei_P001076 | OG2 |
Ocar_g713 | OG2 |
Clacor_Ccor_scaffold14159_01 | OG2 |
Lcom_lcpid74393 | OG2 |
Lcom_lcpid215947 | OG2 |
Scil_scict016803.1 | OG2 |
Leuniv_Lniv_C810120_01 | OG2 |
Gracom_Gcom_scaffold13352_01 | OG2 |
Hsap_ENST00000260956 | OG2 |
Spur_XM_030974234.1 | OG2 |
Skow_XM_002734267.2 | OG2 |
Bralan_BLAG17000936_1 | OG2 |
Dpul_mRNA16660 | OG2 |
Cscu_XM_023371731.1 | OG2 |
Lcom_lcpid111324 | OG2 |
Acagra_model.g23079.t1.1.5d3b8abc | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA173881 | OG2 |
Cgig_XM_034446472.1 | OG2 |
Lcom_lcpid123236 | OG2 |
Exapal_XM_021040669.2 | OG2 |
Chem_TCONS_00024474 | OG2 |
Hvul_g21810_1 | OG2 |
Hvul_g21814_1 | OG2 |
Morvir_scaffold180.g1.t1 | OG2 |
Aaur_scaffold2522.g4.t1 | OG2 |
Nemnom_g2983.t1 | OG2 |
Rhoesc_Res_004275-T1 | OG2 |
Nvec_v1g236578 | OG2 |
Nvec_v1g224449 | OG2 |
Exapal_XM_021061098.2 | OG2 |
Spis_XP_022777450_1 | OG2 |
Pocdam_XM_027193505.1 | OG2 |
Gfas_gfas1.m1.21897.m1 | OG2 |
Fspp_ffun1.m4.16331.m1 | OG2 |
Gasp_gasp1.m3.18851.m1 | OG2 |
Fspp_ffun1.m4.16330.m1 | OG2 |
Gfas_gfas1.m1.4171.m1 | OG2 |
Amil_Amillepora33576-RA | OG2 |
Adig_XM_015904134.1 | OG2 |
Morvir_scaffold685.g5.t1 | OG2 |
Xesp_022033-T1 | OG2 |
Dgig_XM_028561613.1 | OG2 |
Dgig_XM_028547105.1 | OG2 |
Dgig_XM_028561623.1 | OG2 |
Eupdun_Edun_sb10665174 | OG2 |
Lampan_Lpan_14152_01 | OG2 |
Lampan_Lpan_14153_01 | OG2 |
Vmul_SimC394166 | OG2 |
Vmul_sbMO443900 | OG2 |
Vmul_sbMO443893 | OG2 |
Coemet_Cmet_scaffold4255_01 | OG2 |
Coeast_Cast_sb811637 | OG2 |
Coeast_Cast_sb811631 | OG2 |
Coeast_Cast_sb811636 | OG2 |
Coeast_Cast_sb811640 | OG2 |
Coeast_Cast_sb811638 | OG2 |
Coeast_Cast_sb811632 | OG2 |
Coeast_Cast_sb811633 | OG2 |
Coeast_Cast_sb811639 | OG2 |
Coeast_Cast_sb811635 | OG2 |
Bolinf_Binf_sb1242849 | OG2 |
Bolinf_Binf_sb1242850 | OG2 |
Mertsp_Mesp_sb577996 | OG2 |
Mertsp_Mesp_sb577997 | OG2 |
Coeast_Cast_sb762903 | OG2 |
Pbac_sb2663336 | OG2 |
Horcal_Hcv1.av93.c12.g452.i1 | OG2 |
Pbac_sb2654896 | OG2 |
Drygla_Dgla_sb294363 | OG2 |
Beraby_Baby_sb344948 | OG2 |
Beraby_Baby_sb344947 | OG2 |
Bolinf_Binf_sb1302725 | OG2 |
Mlei_ML0405.g15.i1 | OG2 |
Vmul_sbMO451986 | OG2 |
Bolinf_Binf_sb1279788 | OG2 |
Mertsp_Mesp_sb583977 | OG2 |
Bolinf_Binf_sb1279805 | OG2 |
Mertsp_Mesp_sb583990 | OG2 |
Bolinf_Binf_sb1279799 | OG2 |
Mertsp_Mesp_sb583986 | OG2 |
0.416667