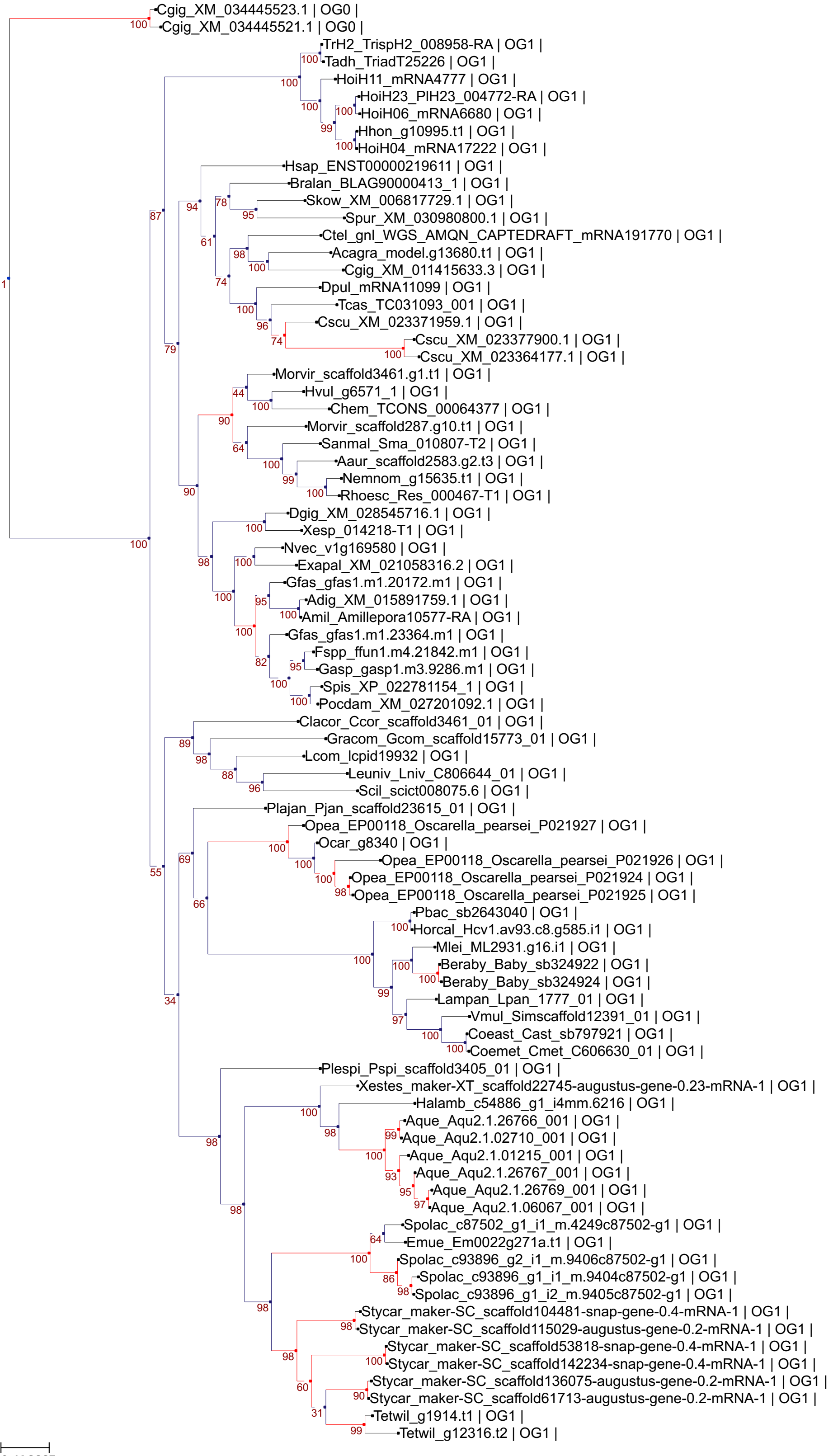
Cgig_XM_034445523.1 | OG0 |
Cgig_XM_034445521.1 | OG0 |
TrH2_TrispH2_008958-RA | OG1 |
Tadh_TriadT25226 | OG1 |
HoiH11_mRNA4777 | OG1 |
HoiH23_PlH23_004772-RA | OG1 |
HoiH06_mRNA6680 | OG1 |
Hhon_g10995.t1 | OG1 |
HoiH04_mRNA17222 | OG1 |
Hsap_ENST00000219611 | OG1 |
Bralan_BLAG90000413_1 | OG1 |
Skow_XM_006817729.1 | OG1 |
Spur_XM_030980800.1 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA191770 | OG1 |
Acagra_model.g13680.t1 | OG1 |
Cgig_XM_011415633.3 | OG1 |
Dpul_mRNA11099 | OG1 |
Tcas_TC031093_001 | OG1 |
Cscu_XM_023371959.1 | OG1 |
Cscu_XM_023377900.1 | OG1 |
Cscu_XM_023364177.1 | OG1 |
Morvir_scaffold3461.g1.t1 | OG1 |
Hvul_g6571_1 | OG1 |
Chem_TCONS_00064377 | OG1 |
Morvir_scaffold287.g10.t1 | OG1 |
Sanmal_Sma_010807-T2 | OG1 |
Aaur_scaffold2583.g2.t3 | OG1 |
Nemnom_g15635.t1 | OG1 |
Rhoesc_Res_000467-T1 | OG1 |
Dgig_XM_028545716.1 | OG1 |
Xesp_014218-T1 | OG1 |
Nvec_v1g169580 | OG1 |
Exapal_XM_021058316.2 | OG1 |
Gfas_gfas1.m1.20172.m1 | OG1 |
Adig_XM_015891759.1 | OG1 |
Amil_Amillepora10577-RA | OG1 |
Gfas_gfas1.m1.23364.m1 | OG1 |
Fspp_ffun1.m4.21842.m1 | OG1 |
Gasp_gasp1.m3.9286.m1 | OG1 |
Spis_XP_022781154_1 | OG1 |
Pocdam_XM_027201092.1 | OG1 |
Clacor_Ccor_scaffold3461_01 | OG1 |
Gracom_Gcom_scaffold15773_01 | OG1 |
Lcom_lcpid19932 | OG1 |
Leuniv_Lniv_C806644_01 | OG1 |
Scil_scict008075.6 | OG1 |
Plajan_Pjan_scaffold23615_01 | OG1 |
Opea_EP00118_Oscarella_pearsei_P021927 | OG1 |
Ocar_g8340 | OG1 |
Opea_EP00118_Oscarella_pearsei_P021926 | OG1 |
Opea_EP00118_Oscarella_pearsei_P021924 | OG1 |
Opea_EP00118_Oscarella_pearsei_P021925 | OG1 |
Pbac_sb2643040 | OG1 |
Horcal_Hcv1.av93.c8.g585.i1 | OG1 |
Mlei_ML2931.g16.i1 | OG1 |
Beraby_Baby_sb324922 | OG1 |
Beraby_Baby_sb324924 | OG1 |
Lampan_Lpan_1777_01 | OG1 |
Vmul_Simscaffold12391_01 | OG1 |
Coeast_Cast_sb797921 | OG1 |
Coemet_Cmet_C606630_01 | OG1 |
Plespi_Pspi_scaffold3405_01 | OG1 |
Xestes_maker-XT_scaffold22745-augustus-gene-0.23-mRNA-1 | OG1 |
Halamb_c54886_g1_i4mm.6216 | OG1 |
Aque_Aqu2.1.26766_001 | OG1 |
Aque_Aqu2.1.02710_001 | OG1 |
Aque_Aqu2.1.01215_001 | OG1 |
Aque_Aqu2.1.26767_001 | OG1 |
Aque_Aqu2.1.26769_001 | OG1 |
Aque_Aqu2.1.06067_001 | OG1 |
Spolac_c87502_g1_i1_m.4249c87502-g1 | OG1 |
Emue_Em0022g271a.t1 | OG1 |
Spolac_c93896_g2_i1_m.9406c87502-g1 | OG1 |
Spolac_c93896_g1_i1_m.9404c87502-g1 | OG1 |
Spolac_c93896_g1_i2_m.9405c87502-g1 | OG1 |
Stycar_maker-SC_scaffold104481-snap-gene-0.4-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold115029-augustus-gene-0.2-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold53818-snap-gene-0.4-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold142234-snap-gene-0.4-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold136075-augustus-gene-0.2-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold61713-augustus-gene-0.2-mRNA-1 | OG1 |
Tetwil_g1914.t1 | OG1 |
Tetwil_g12316.t2 | OG1 |
0.416667