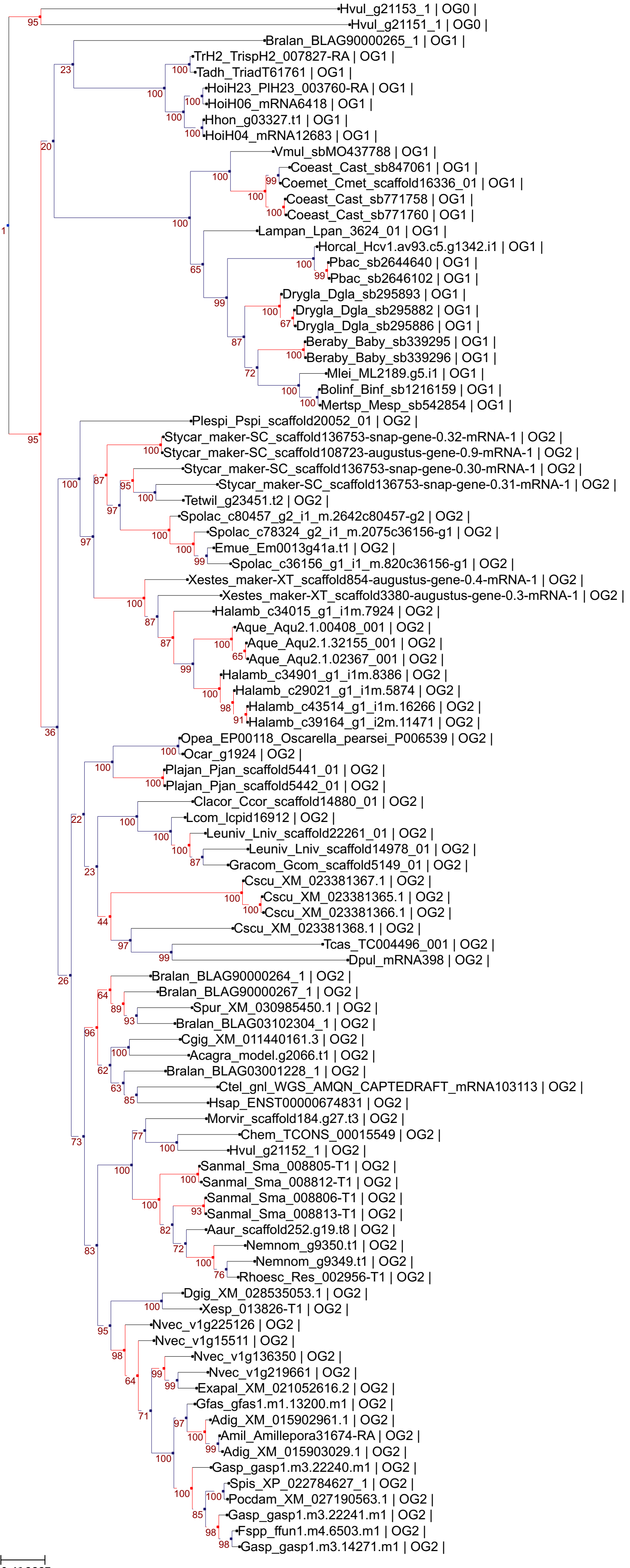
Hvul_g21153_1 | OG0 |
Hvul_g21151_1 | OG0 |
Bralan_BLAG90000265_1 | OG1 |
TrH2_TrispH2_007827-RA | OG1 |
Tadh_TriadT61761 | OG1 |
HoiH23_PIH23_003760-RA | OG1 |
HoiH06_mRNA6418 | OG1 |
Hhon_g03327.t1 | OG1 |
HoiH04_mRNA12683 | OG1 |
Vmul_sbMO437788 | OG1 |
Coeast_Cast_sb847061 | OG1 |
Coemet_Cmet_scaffold16336_01 | OG1 |
Coeast_Cast_sb771758 | OG1 |
Coeast_Cast_sb771760 | OG1 |
Lampan_Lpan_3624_01 | OG1 |
Horcal_Hcv1.av93.c5.g1342.i1 | OG1 |
Pbac_sb2644640 | OG1 |
Pbac_sb2646102 | OG1 |
Drygla_Dgla_sb295893 | OG1 |
Drygla_Dgla_sb295882 | OG1 |
Drygla_Dgla_sb295886 | OG1 |
Beraby_Baby_sb339295 | OG1 |
Beraby_Baby_sb339296 | OG1 |
Mlei_ML2189.g5.i1 | OG1 |
Bolinf_Binf_sb1216159 | OG1 |
Mertsp_Mesp_sb542854 | OG1 |
Plespi_Pspi_scaffold20052_01 | OG2 |
Stycar_maker-SC_scaffold136753-snap-gene-0.32-mRNA-1 | OG2 |
Stycar_maker-SC_scaffold108723-augustus-gene-0.9-mRNA-1 | OG2 |
Stycar_maker-SC_scaffold136753-snap-gene-0.30-mRNA-1 | OG2 |
Stycar_maker-SC_scaffold136753-snap-gene-0.31-mRNA-1 | OG2 |
Tetwil_g23451.t2 | OG2 |
Spolac_c80457_g2_i1_m.2642c80457-g2 | OG2 |
Spolac_c78324_g2_i1_m.2075c36156-g1 | OG2 |
Emue_Em0013g41a.t1 | OG2 |
Spolac_c36156_g1_i1_m.820c36156-g1 | OG2 |
Xestes_maker-XT_scaffold854-augustus-gene-0.4-mRNA-1 | OG2 |
Xestes_maker-XT_scaffold3380-augustus-gene-0.3-mRNA-1 | OG2 |
Halamb_c34015_g1_i1m.7924 | OG2 |
Aque_Aqu2.1.00408_001 | OG2 |
Aque_Aqu2.1.32155_001 | OG2 |
Aque_Aqu2.1.02367_001 | OG2 |
Halamb_c34901_g1_i1m.8386 | OG2 |
Halamb_c29021_g1_i1m.5874 | OG2 |
Halamb_c43514_g1_i1m.16266 | OG2 |
Halamb_c39164_g1_i2m.11471 | OG2 |
Opea_EP00118_Oscarella_pearsei_P006539 | OG2 |
Ocar_g1924 | OG2 |
Plajan_Pjan_scaffold5441_01 | OG2 |
Plajan_Pjan_scaffold5442_01 | OG2 |
Clacor_Ccor_scaffold14880_01 | OG2 |
Lcom_lcpid16912 | OG2 |
Leuniv_Lniv_scaffold22261_01 | OG2 |
Leuniv_Lniv_scaffold14978_01 | OG2 |
Gracom_Gcom_scaffold5149_01 | OG2 |
Cscu_XM_023381367.1 | OG2 |
Cscu_XM_023381365.1 | OG2 |
Cscu_XM_023381366.1 | OG2 |
Cscu_XM_023381368.1 | OG2 |
Tcas_TC004496_001 | OG2 |
Dpul_mRNA398 | OG2 |
Bralan_BLAG90000264_1 | OG2 |
Bralan_BLAG90000267_1 | OG2 |
Spur_XM_030985450.1 | OG2 |
Bralan_BLAG03102304_1 | OG2 |
Cgig_XM_011440161.3 | OG2 |
Acagra_model.g2066.t1 | OG2 |
Bralan_BLAG03001228_1 | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA103113 | OG2 |
Hsap_ENST00000674831 | OG2 |
Morvir_scaffold184.g27.t3 | OG2 |
Chem_TCONS_00015549 | OG2 |
Hvul_g21152_1 | OG2 |
Sanmal_Sma_008805-T1 | OG2 |
Sanmal_Sma_008812-T1 | OG2 |
Sanmal_Sma_008806-T1 | OG2 |
Sanmal_Sma_008813-T1 | OG2 |
Aaur_scaffold252.g19.t8 | OG2 |
Nemnom_g9350.t1 | OG2 |
Nemnom_g9349.t1 | OG2 |
Rhoesc_Res_002956-T1 | OG2 |
Dgig_XM_028535053.1 | OG2 |
Xesp_013826-T1 | OG2 |
Nvec_v1g225126 | OG2 |
Nvec_v1g15511 | OG2 |
Nvec_v1g136350 | OG2 |
Nvec_v1g219661 | OG2 |
Exapal_XM_021052616.2 | OG2 |
Gfas_gfas1.m1.13200.m1 | OG2 |
Adig_XM_015902961.1 | OG2 |
Amil_Amillepora31674-RA | OG2 |
Adig_XM_015903029.1 | OG2 |
Gasp_gasp1.m3.22240.m1 | OG2 |
Spis_XP_022784627_1 | OG2 |
Pocdam_XM_027190563.1 | OG2 |
Gasp_gasp1.m3.22241.m1 | OG2 |
Fspp_ffun1.m4.6503.m1 | OG2 |
Gasp_gasp1.m3.14271.m1 | OG2 |
0.416667