A phylogenetic tree with the following taxa labels:

- Bralan_BLAG01001886_1 | OG1 |
- Hsap_ENST00000321367 | OG1 |
- Hsap_ENST00000455843 | OG1 |
- Hsap_ENST00000672673 | OG1 |
- Skow_XM_002730912.2 | OG2 |
- Bralan_BLAG01001883_1 | OG2 |
- Spur_XM_030992455.1 | OG2 |
- Cscu_XM_023359703.1 | OG2 |
- Tcas_TC006627_001 | OG2 |
- Dpul_mRNA21304 | OG2 |
- Cgig_XM_034481240.1 | OG2 |
- Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA173215 | OG2 |
- Acagra_model.g23686.t1 | OG2 |
- TrH2_TrispH2_006913-RA | OG0 |
- Tadh_TriadT50746 | OG0 |
- HoiH23_PlH23_010642-RA | OG0 |
- HoiH06_mRNA8293 | OG0 |
- HoiH11_mRNA9630 | OG0 |
- Hhon_g07733.t1 | OG0 |
- HoiH04_mRNA9650 | OG0 |
- Eupdun_Edun_sb10680495 | OG0 |
- Horcal_Hcv1.av93.c12.g169.i1 | OG0 |
- Pbac_sb2666182 | OG0 |
- Pbac_sb2644867 | OG0 |
- Pbac_sb2641095 | OG0 |
- Vmul_Simscaffold5649_01 | OG0 |
- Coemet_Cmet_scaffold7245_01 | OG0 |
- Drygla_Dgla_sb297541 | OG0 |
- Drygla_Dgla_sb297537 | OG0 |
- Lampan_Lpan_1938_01 | OG0 |
- Beraby_Baby_sb326522 | OG0 |
- Bolinf_Binf_sb1248731 | OG0 |
- Mlei_ML1288.g3.i1 | OG0 |
- Chem_TCONS_00059040 | OG0 |
- Hvul_g8709_1 | OG0 |
- Morvir_scaffold180.g34.t1 | OG0 |
- Sanmal_Sma_019223-T3 | OG0 |
- Aaur_scaffold401.g12.t3 | OG0 |
- Aaur_scaffold401.g11.t2 | OG0 |
- Rhoesc_Res_006572-T4 | OG0 |
- Nemnom_g15057.t1 | OG0 |
- Dgig_XM_028544621.1 | OG0 |
- Nvec_v1g239234 | OG0 |
- Exapal_XM_021038281.2 | OG0 |
- Fspp_ffun1.m4.25132.m1 | OG0 |
- Pocdam_XM_027183161.1 | OG0 |
- Spis_XP_022780489_1 | OG0 |
- Spis_XP_022780487_1 | OG0 |
- Gasp_gasp1.m3.17475.m1 | OG0 |
- Gfas_gfas1.m1.624.m1 | OG0 |
- Adig_XM_015916845.1 | OG0 |
- Amil_Amillepora22517-RA | OG0 |
- Lcom_lcpid142212 | OG0 |
- Plajan_Pjan_C556817_01 | OG0 |
- Opea_EP00118_Oscarella_pearsei_P028805 | OG0 |
- Ocar_g1788 | OG0 |
- Clacor_Ccor_scaffold9789_01 | OG0 |
- Gracom_Gcom_C553542_01 | OG0 |
- Lcom_lcpid57592 | OG0 |
- Leuniv_Lniv_scaffold8550_01 | OG0 |
- Clacor_Ccor_scaffold2870_01 | OG0 |
- Leuniv_Lniv_scaffold20180_01 | OG0 |
- Gracom_Gcom_scaffold12546_01 | OG0 |
- Lcom_lcpid166049 | OG0 |
- Scil_scict022571.1 | OG0 |
- Plespi_Pspi_scaffold603_01 | OG0 |
- Emue_Em0015g1221a.t1 | OG0 |
- Spolac_c87907_g1_i1_m.4592c87907-g1 | OG0 |
- Stycar_maker-SC_scaffold971-snap-gene-0.13-mRNA-1 | OG0 |
- Tetwil_g4291.t2 | OG0 |
- Stycar_maker-SC_scaffold2524-augustus-gene-0.1-mRNA-1 | OG0 |
- Halamb_c53526_g2_i2m.35779 | OG0 |
- Aque_Aqu2.1.23238_001 | OG0 |
- Xestes_maker-XT_scaffold73001-augustus-gene-0.12-mRNA-1 | OG0 |
- Halamb_c2085_g1_i1m.320 | OG0 |
- Halamb_c33994_g1_i1m.7915 | OG0 |
- Halamb_c37132_g1_i1m.9795 | OG0 |
- Aque_Aqu2.1.02905_001 | OG0 |
- Aque_Aqu2.1.23239_001 | OG0 |
- Tcas_TC014984_001 | OG0 |
- Dpul_mRNA6749 | OG0 |
- Cscu_XM_023359705.1 | OG0 |
- Cscu_XM_023386057.1 | OG0 |
- Cscu_XM_023362767.1 | OG0 |
- Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA177047 | OG0 |
- Cgig_XM_011419765.3 | OG0 |
- Acagra_model.g23683.t1.1.5d3b8ae3 | OG0 |
- Spur_XM_030992282.1 | OG0 |
- Skow_XM_002738866.2 | OG0 |
- Bralan_BLAG01001881_1 | OG0 |
- Hsap_ENST00000616972 | OG0 |
- Xesp_004637-T1 | OG0 |
- Dgig_XM_028544876.1 | OG0 |
- Morvir_scaffold229.g24.t2 | OG0 |
- Sanmal_Sma_018993-T1 | OG0 |
- Rhoesc_Res_006071-T1 | OG0 |
- Aaur_scaffold818.g10.t1 | OG0 |
- Chem_TCONS_00057235 | OG0 |
- Hvul_g14507_1 | OG0 |
- Hvul_g14508_1 | OG0 |
- Exapal_XM_021053467.2 | OG0 |
- Nvec_v1g223708 | OG0 |
- Nvec_v1g130720 | OG0 |
- Fspp_ffun1.m4.6744.m1 | OG0 |
- Gasp_gasp1.m3.6909.m1 | OG0 |
- Spis_XP_022792071_1 | OG0 |
- Pocdam_XM_027193431.1 | OG0 |
- Gfas_gfas1.m1.14173.m1 | OG0 |
- Amil_Amillepora22151-RA | OG0 |
- Adig_XM_015906788.1 | OG0 |

Scale bar: 0.416667