A phylogenetic tree diagram. The scale bar at the bottom left reads 0.416667.

Leaf labels (top to bottom):

Xestes_maker-XT_scaffold4896-augustus-gene-0.37-mRNA-1 | OG0 |
Aque_Aqu2.1.32109_001 | OG0 |
Halamb_c37078_g1_i1mm.473 | OG0 |
Tetwil_g17623.t2 | OG0 |
Stycar_snap_masked-SC_scaffold1907-processed-gene-0.2-mRNA-1 | OG0 |
Emue_Em0021g248a.t1 | OG0 |
Spolac_c99292_g1_i1_m.19270c99292-g1 | OG0 |
Stycar_maker-SC_scaffold140749-augustus-gene-0.3-mRNA-1 | OG0 |
Aque_Aqu2.1.32002_001 | OG0 |
Halamb_c52026_g1_i1m.31960 | OG0 |
Tetwil_g15800.t1 | OG0 |
Spolac_c94516_g1_i1_m.10183c104494-g1 | OG0 |
Spolac_c104494_g1_i2_m.40934c104494-g1 | OG0 |
Spolac_c104494_g1_i1_m.40931c104494-g1 | OG0 |
Emue_Em0011g995a.t1 | OG0 |
Emue_Em0011g1001a.t1 | OG0 |
Emue_Em0011g1010a.t1 | OG0 |
Emue_Em0011g997a.t1 | OG0 |
Emue_Em0011g1009a.t1 | OG0 |
Emue_Em0011g992a.t1 | OG0 |
Emue_Em0011g1003a.t1 | OG0 |
Nvec_v1g168581 | OG0 |
Exapal_XM_021046420.2 | OG0 |
Adig_XM_015918121.1 | OG0 |
Amil_Amillepora06398-RA | OG0 |
Adig_XM_015918122.1 | OG0 |
Gfas_gfas1.m1.1424.m1 | OG0 |
Fspp_ffun1.m4.14734.m1 | OG0 |
Gasp_gasp1.m3.30224.m1 | OG0 |
Spis_XP_022794023_1 | OG0 |
Pocdam_XM_027204607.1 | OG0 |
Xesp_015005-T1 | OG0 |
Nvec_v1g106062 | OG0 |
Nvec_v1g106211 | OG0 |
Exapal_XM_028656633.1 | OG0 |
Nvec_v1g106104 | OG0 |
Gfas_gfas1.m1.20328.m1 | OG0 |
Gasp_gasp1.m3.30226.m1 | OG0 |
Fspp_ffun1.m4.21558.m1 | OG0 |
Fspp_ffun1.m4.21559.m1 | OG0 |
Gfas_gfas1.m1.1425.m1 | OG0 |
Amil_Amillepora06399-RA | OG0 |
Adig_XM_015918114.1 | OG0 |
Spis_XP_022794036_1 | OG0 |
Pocdam_XM_027179837.1 | OG0 |
Fspp_ffun1.m4.14735.m1 | OG0 |
Gasp_gasp1.m3.30225.m1 | OG0 |
Dpul_mRNA19352 | OG2 |
Cgig_XM_034473439.1 | OG2 |
Spur_XM_030987746.1 | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA226189 | OG2 |
Acagra_model.g4403.t1 | OG2 |
Acagra_model.g4404.t1 | OG2 |
Nvec_v1g230851 | OG3 |
Nvec_v1g194898 | OG3 |
Gfas_gfas1.m1.18858.m1 | OG3 |
Adig_XM_015905200.1 | OG3 |
Amil_Amillepora05036-RA | OG3 |
Fspp_ffun1.m4.13031.m1 | OG3 |
Gasp_gasp1.m3.763.m1 | OG3 |
Spis_XP_022798906_1 | OG3 |
Pocdam_XM_027200321.1 | OG3 |
Spur_XM_780780.5 | OG3 |
Bralan_BLAG18000070_1 | OG3 |
Hsap_ENST00000008527 | OG3 |
Hsap_ENST00000443527 | OG3 |
Cgig_XM_011456825.3 | OG3 |
Acagra_model.g25645.t1 | OG3 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA178510 | OG3 |
Cscu_XM_023386415.1 | OG3 |
Dpul_mRNA9179 | OG3 |
Tcas_TC010454_001 | OG3 |
Bralan_BLAG10000392_1 | OG1 |
Bralan_BLAG18000072_1 | OG1 |
Nvec_v1g203127 | OG1 |
Exapal_XM_021059408.2 | OG1 |
Xesp_015804-T1 | OG1 |
Adig_XM_015903146.1 | OG1 |
Amil_Amillepora00156-RA | OG1 |
Gfas_gfas1.m1.4328.m1 | OG1 |
Fspp_ffun1.m4.31073.m1 | OG1 |
Gasp_gasp1.m3.1042.m1 | OG1 |
Spis_XP_022808227_1 | OG1 |
Pocdam_XM_027187917.1 | OG1 |
Spur_XM_030997502.1 | OG1 |
Horcal_Hcv1.av93.c2.g829.i1 | OG1 |
Lampan_Lpan_14604_01 | OG1 |
Mlei_ML0530.g4.i1 | OG1 |
Drygla_Dgla_sb299915 | OG1 |
Beraby_Baby_sb339147 | OG1 |
Beraby_Baby_sb339145 | OG1 |
Cgig_XM_011416394.3 | OG1 |
Acagra_model.g443.t1 | OG1 |
Acagra_model.g443.t1.1.5d3b8234 | OG1 |
Dpul_mRNA874 | OG1 |
Dpul_mRNA26007 | OG1 |
Dpul_mRNA14300 | OG1 |
Dpul_mRNA14296 | OG1 |
Nemnom_g4554.t1 | OG1 |
Rhoesc_Res_008527-T1 | OG1 |
Chem_TCONS_00062708 | OG1 |
Coeast_Cast_sb751164 | OG1 |
Coemet_Cmet_C600285_01 | OG1 |
Coeast_Cast_sb773050 | OG1 |

0.416667