A phylogenetic tree with labelled leaf nodes. Branch support values and node labels are shown along the tree. The following taxon labels appear (top to bottom):

Emue_Em0013g714a.t1 | OG0
Stycar_snap_masked-SC_scaffold5160-processed-gene-0.25-mRNA-1 | OG0
Tetwil_g23238.t2 | OG0
Aque_Aqu2.1.40621_001 | OG0
Halamb_c42115_g1_i1m.14560 | OG0
Xestes_augustus_masked-XT_scaffold105391-processed-gene-0.1-mRNA-1 | OG0
Xestes_maker-XT_scaffold872-augustus-gene-0.18-mRNA-1 | OG0
Opea_EP00118_Oscarella_pearsei_P004882 | OG0
Plajan_Pjan_C551473_01 | OG0
Clacor_Ccor_scaffold8216_01 | OG0
Clacor_Ccor_scaffold9012_01 | OG0
Clacor_Ccor_scaffold9013_01 | OG0
Gracom_Gcom_scaffold4416_01 | OG0
Gracom_Gcom_scaffold4226_01 | OG0
Scil_scict017101.1 | OG0
Leuniv_Lniv_scaffold23565_01 | OG0
Scil_scict001601.2 | OG0
Lcom_lcpid64678 | OG0
Lcom_lcpid73800 | OG0
Lcom_lcpid71534 | OG0
Lcom_lcpid88611 | OG0
Lcom_lcpid100716 | OG0
Lcom_lcpid97573 | OG0
Hsap_ENST00000380871 | OG0
Morvir_scaffold477.g8.t1 | OG0
Sanmal_Sma_026504-T1 | OG0
Nemnom_g2105.t1 | OG0
Rhoesc_Res_001049-T1 | OG0
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA79432 | OG0
Tcas_TC012743_001 | OG0
Acagra_model.g1596.t1 | OG0
Cscu_XM_023361107.1 | OG0
Cgig_XM_011414579.3 | OG0
Bralan_BLAG03001042_1 | OG0
Hsap_ENST00000382438 | OG0
Skow_NM_001165004.1 | OG0
Spur_XM_779642.5 | OG0
Dgig_XM_028547334.1 | OG0
Xesp_013038-T1 | OG0
Exapal_XM_021044213.2 | OG0
Nvec_v1g208034 | OG0
Amil_Amillepora06232-RA | OG0
Adig_XM_015915185.1 | OG0
Spis_XP_022782914_1 | OG0
Pocdam_XM_027185775.1 | OG0
Fspp_ffun1.m4.4461.m1 | OG0
Gasp_gasp1.m3.3680.m1 | OG0
Hvul_g31954_1 | OG1
Chem_TCONS_00016211 | OG1
Morvir_scaffold11.g115.t1 | OG1
Nemnom_g13472.t1 | OG1
Sanmal_Sma_008525-T1 | OG1
Rhoesc_Res_003043-T1 | OG1
Xesp_013044-T1 | OG1
Dgig_XM_028547544.1 | OG1
Exapal_XM_021044212.2 | OG1
Nvec_v1g208035 | OG1
Gasp_gasp1.m3.679.m1 | OG1
Spis_XP_022782912_1 | OG1
Pocdam_XM_027185774.1 | OG1
Fspp_ffun1.m4.4460.m1 | OG1
Gfas_gfas1.m1.19176.m1 | OG1
Amil_Amillepora06234-RA | OG1
Adig_XM_015915184.1 | OG1
Dpul_mRNA26167 | OG2
Tcas_TC011745_001 | OG2
Spur_XM_778030.4 | OG2
Skow_XM_006815396.1 | OG2
Bralan_BLAG03001041_2 | OG2
Hsap_ENST00000325017 | OG2
Hsap_ENST00000329198 | OG2
Hsap_ENST00000344586 | OG2
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA170007 | OG2
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA220898 | OG2
Cscu_XM_023361114.1 | OG2
Acagra_model.g1144.t1 | OG2
Cgig_XM_011456691.3 | OG2
HoiH11_mRNA4950 | OG3
Tadh_TriadT9507 | OG3
TrH2_TrispH2_003382-RA | OG3
HoiH06_mRNA11553 | OG3
Hhon_g07292.t1 | OG3
HoiH23_PlH23_010042-RA | OG3
HoiH04_mRNA8177 | OG3
Cscu_XM_023361927.1 | OG3
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA172338 | OG3
Cgig_XM_011453501.3 | OG3
Acagra_model.g2292.t1 | OG3
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA40416 | OG3
Bralan_BLAG04001574_1 | OG3
Spur_NM_214653.1 | OG3
Skow_NM_001164910.1 | OG3
Hsap_ENST00000354822 | OG3
Hsap_ENST00000351817 | OG3
Tcas_TC008996_001 | OG3
Cscu_XM_023376262.1 | OG3
Dpul_mRNA2105 | OG3
Dpul_mRNA6100 | OG4
Tcas_TC007014_001 | OG4
Cscu_XM_023368090.1 | OG4
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA221803 | OG4
Cgig_XM_011439089.3 | OG4
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA70572 | OG4
Acagra_model.g2291.t1 | OG4
Spur_NM_001129811.2 | OG4
Skow_NM_001164932.1 | OG4
Bralan_BLAG04001575_1 | OG4
Hsap_ENST00000377142 | OG4
Hsap_ENST00000258829 | OG4
HoiH06_mRNA3929 | OG4
HoiH04_mRNA13330 | OG4
HoiH23_PlH23_010043-RA | OG4
TrH2_TrispH2_003383-RA | OG4
Tadh_TriadT9561 | OG4
Dgig_XM_028537572.1 | OG4
Xesp_021545-T1 | OG4
Dgig_XM_028553282.1 | OG4
Xesp_021495-T1 | OG4
Morvir_scaffold273.g2.t1 | OG4
Morvir_scaffold83.g2.t1 | OG4
Nemnom_g19277.t1 | OG4
Sanmal_Sma_023401-T1 | OG4
Chem_TCONS_00013206 | OG4
Lcom_lcpid91511 | OG4
Chem_TCONS_00069083 | OG4
Morvir_scaffold46.g33.t1 | OG4
Chem_TCONS_00070249 | OG4
Scar_scaffold2956.g1.t1 | OG4
Nemnom_g19279.t1 | OG4
Rhoesc_Res_004687-T1 | OG4
Nvec_v1g68616 | OG4
Exapal_XM_021051233.2 | OG4
Xesp_022051-T1 | OG4
Amil_Amillepora26578-RA | OG4
Adig_XM_015905817.1 | OG4
Gfas_gfas1.m1.776.m1 | OG4
Spis_XP_022794616_1 | OG4
Pocdam_XM_027183109.1 | OG4
Fspp_ffun1.m4.14105.m1 | OG4
Gasp_gasp1.m3.4129.m1 | OG4
Dgig_XM_028537344.1 | OG4
Xesp_016908-T1 | OG4
Xesp_021558-T1 | OG4
Nvec_v1g192469 | OG4
Exapal_XM_021051226.2 | OG4
Spis_XP_022794643_1 | OG4
Pocdam_XM_027183114.1 | OG4
Amil_Amillepora26577-RA | OG4
Adig_XM_015905819.1 | OG4
Exapal_XM_021051225.2 | OG4
Nvec_v1g127828 | OG4
Amil_Amillepora26581-RA | OG4
Adig_XM_015905823.1 | OG4
Gfas_gfas1.m1.775.m1 | OG4
Fspp_ffun1.m4.14112.m1 | OG4
Gasp_gasp1.m3.4134.m1 | OG4
Spis_XP_022794603_1 | OG4
Pocdam_XM_027183103.1 | OG4
Nvec_v1g127891 | OG4
Exapal_XM_021051229.2 | OG4
Gasp_gasp1.m3.4132.m1 | OG4
Spis_XP_022794604_1 | OG4
Pocdam_XM_027183104.1 | OG4
Gfas_gfas1.m1.773.m1 | OG4
Amil_Amillepora26580-RA | OG4
Adig_XM_015905827.1 | OG4
Nvec_v1g48313 | OG4
Exapal_XM_021051231.2 | OG4
Gfas_gfas1.m1.777.m1 | OG4
Amil_Amillepora26579-RA | OG4
Adig_XM_015905829.1 | OG4
Fspp_ffun1.m4.14106.m1 | OG4
Gasp_gasp1.m3.4130.m1 | OG4
Pocdam_XM_027183102.1 | OG4
Spis_XP_022794618_1 | OG4

Scale bar: 0.416667