A phylogenetic tree diagram with the following taxa labels (read top to bottom):

- Dpul_mRNA10988 | OG0 |
- Tcas_TC033579_002 | OG0 |
- Pbac_sb2667786 | OG0 |
- Horcal_Hcv1.av93.c6.g993.i1 | OG0 |
- Mlei_ML1855.g23.i1 | OG0 |
- Drygla_Dgla_sb310710 | OG0 |
- Coemet_Cmet_scaffold5241_01 | OG0 |
- Morvir_scaffold73.g36.t1 | OG0 |
- Hvul_g17719_1 | OG0 |
- Aaur_scaffold172.g3.t1 | OG0 |
- Sanmal_Sma_024675-T1 | OG0 |
- Aaur_scaffold5941.g1.t1 | OG0 |
- Nemnom_g15924.t1 | OG0 |
- Rhoesc_Res_010739-T1 | OG0 |
- TrH2_TrispH2_000715-RA | OG0 |
- Tadh_TriadT50816 | OG0 |
- Hhon_g05017.t1 | OG0 |
- HoiH04_mRNA6113 | OG0 |
- HoiH11_mRNA7198 | OG0 |
- HoiH23_PlH23_008034-RA | OG0 |
- HoiH06_mRNA11184 | OG0 |
- Clacor_Ccor_C523987_01 | OG0 |
- Opea_EP00118_Oscarella_pearsei_P010335 | OG0 |
- Ocar_g8950 | OG0 |
- Emue_Em0018g1164a.t1 | OG0 |
- Tetwil_g9204.t1 | OG0 |
- Aque_Aqu2.1.37172_001 | OG0 |
- Xestes_maker-XT_scaffold25661-augustus-gene-0.8-mRNA-1 | OG0 |
- Cscu_XM_023368334.1 | OG0 |
- Bralan_BLAG07001622_1 | OG0 |
- Hsap_ENST00000565504 | OG0 |
- Spur_XM_794025.5 | OG0 |
- Skow_XM_002736220.2 | OG0 |
- Cgig_XM_011447368.3 | OG0 |
- Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA174336 | OG0 |
- Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA111660 | OG0 |
- Dgig_XM_028536611.1 | OG0 |
- Xesp_005363-T1 | OG0 |
- Exapal_XM_021047436.2 | OG0 |
- Gasp_gasp1.m3.3359.m1 | OG0 |
- Spis_XP_022785340_1 | OG0 |
- Pocdam_XM_027181579.1 | OG0 |
- Nvec_NVE7649 | OG0 |
- Adig_XM_015892433.1 | OG0 |
- Gfas_gfas1.m1.14932.m1 | OG0 |
- Amil_Amillepora10803-RA | OG0 |
- Adig_XM_015894168.1 | OG0 |

Branch support values: 74, 28, 100, 100, 90, 95, 54, 61, 74, 98, 46, 86, 78, 100, 97, 85, 87, 99, 70, 100, 99, 99, 91, 100, 1, 35, 26, 72, 84, 87, 47, 89, 100, 17, 95, 99, 89, 53, 100, 52, 92, 99, 43, 69, 28, 26

Scale bar: 0.416667