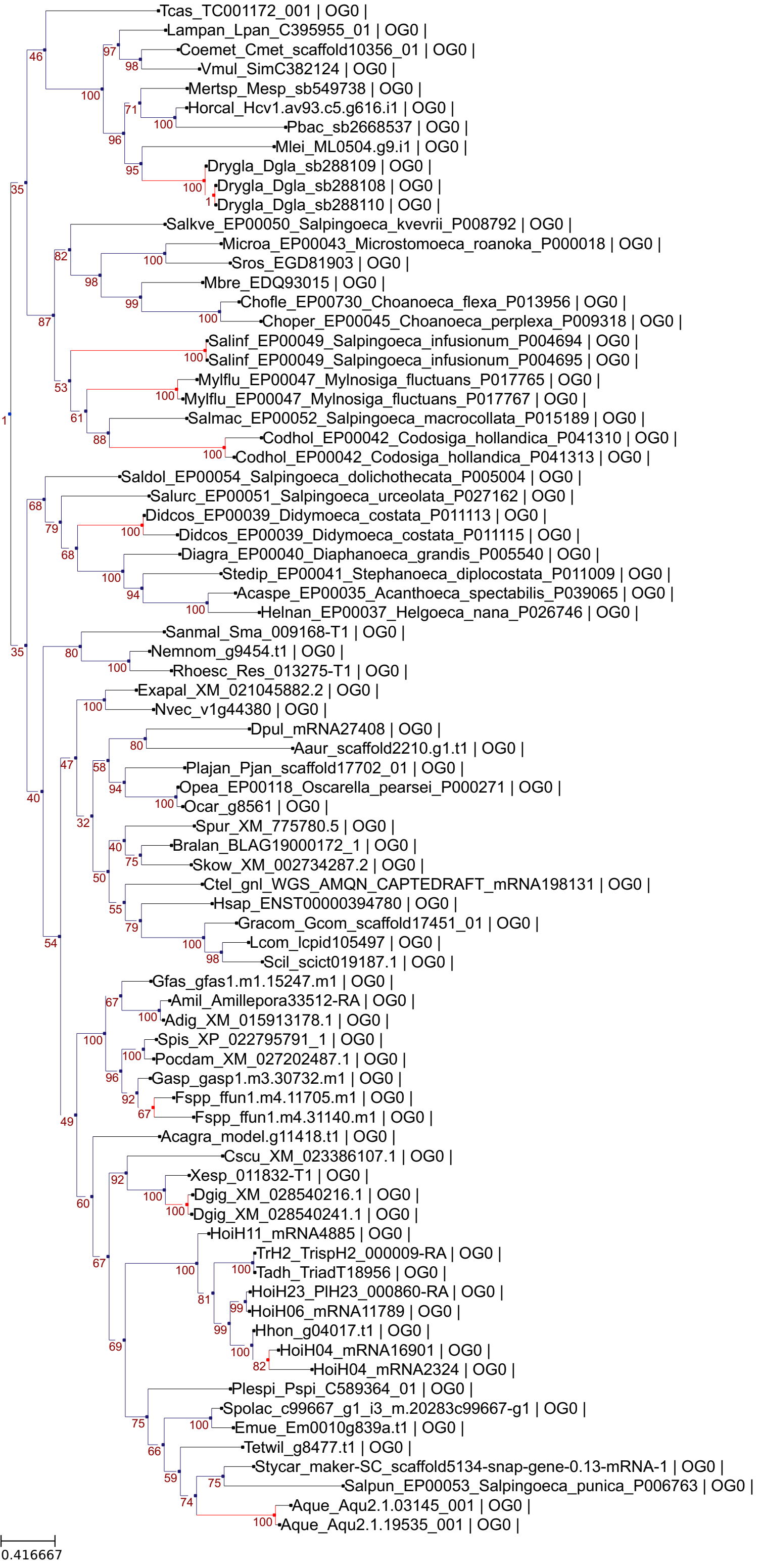
Tcas_TC001172_001 | OG0 |
Lampan_Lpan_C395955_01 | OG0 |
Coemet_Cmet_scaffold10356_01 | OG0 |
Vmul_SimC382124 | OG0 |
Mertsp_Mesp_sb549738 | OG0 |
Horcal_Hcv1.av93.c5.g616.i1 | OG0 |
Pbac_sb2668537 | OG0 |
Mlei_ML0504.g9.i1 | OG0 |
Drygla_Dgla_sb288109 | OG0 |
Drygla_Dgla_sb288108 | OG0 |
Drygla_Dgla_sb288110 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P008792 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P000018 | OG0 |
Sros_EGD81903 | OG0 |
Mbre_EDQ93015 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P013956 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P009318 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P004694 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P004695 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P017765 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P017767 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P015189 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P041310 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P041313 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P005004 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P027162 | OG0 |
Didcos_EP00039_Didymoeca_costata_P011113 | OG0 |
Didcos_EP00039_Didymoeca_costata_P011115 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P005540 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P011009 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P039065 | OG0 |
Helnan_EP00037_Helgoeca_nana_P026746 | OG0 |
Sanmal_Sma_009168-T1 | OG0 |
Nemnom_g9454.t1 | OG0 |
Rhoesc_Res_013275-T1 | OG0 |
Exapal_XM_021045882.2 | OG0 |
Nvec_v1g44380 | OG0 |
Dpul_mRNA27408 | OG0 |
Aaur_scaffold2210.g1.t1 | OG0 |
Plajan_Pjan_scaffold17702_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P000271 | OG0 |
Ocar_g8561 | OG0 |
Spur_XM_775780.5 | OG0 |
Bralan_BLAG19000172_1 | OG0 |
Skow_XM_002734287.2 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA198131 | OG0 |
Hsap_ENST00000394780 | OG0 |
Gracom_Gcom_scaffold17451_01 | OG0 |
Lcom_lcpid105497 | OG0 |
Scil_scict019187.1 | OG0 |
Gfas_gfas1.m1.15247.m1 | OG0 |
Amil_Amillepora33512-RA | OG0 |
Adig_XM_015913178.1 | OG0 |
Spis_XP_022795791_1 | OG0 |
Pocdam_XM_027202487.1 | OG0 |
Gasp_gasp1.m3.30732.m1 | OG0 |
Fspp_ffun1.m4.11705.m1 | OG0 |
Fspp_ffun1.m4.31140.m1 | OG0 |
Acagra_model.g11418.t1 | OG0 |
Cscu_XM_023386107.1 | OG0 |
Xesp_011832-T1 | OG0 |
Dgig_XM_028540216.1 | OG0 |
Dgig_XM_028540241.1 | OG0 |
HoiH11_mRNA4885 | OG0 |
TrH2_TrispH2_000009-RA | OG0 |
Tadh_TriadT18956 | OG0 |
HoiH23_PlH23_000860-RA | OG0 |
HoiH06_mRNA11789 | OG0 |
Hhon_g04017.t1 | OG0 |
HoiH04_mRNA16901 | OG0 |
HoiH04_mRNA2324 | OG0 |
Plespi_Pspi_C589364_01 | OG0 |
Spolac_c99667_g1_i3_m.20283c99667-g1 | OG0 |
Emue_Em0010g839a.t1 | OG0 |
Tetwil_g8477.t1 | OG0 |
Stycar_maker-SC_scaffold5134-snap-gene-0.13-mRNA-1 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P006763 | OG0 |
Aque_Aqu2.1.03145_001 | OG0 |
Aque_Aqu2.1.19535_001 | OG0 |
0.416667