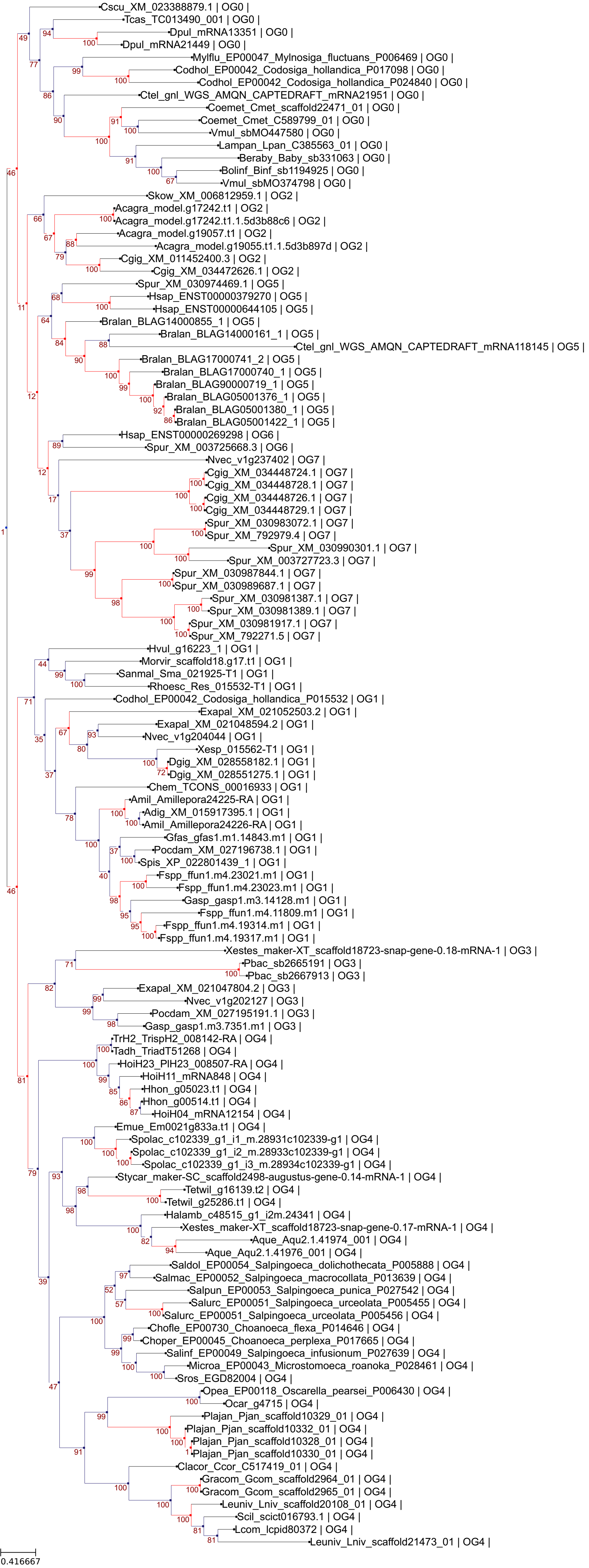
Cscu_XM_023388879.1 | OG0 |
Tcas_TC013490_001 | OG0 |
Dpul_mRNA13351 | OG0 |
Dpul_mRNA21449 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P006469 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P017098 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P024840 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA21951 | OG0 |
Coemet_Cmet_scaffold22471_01 | OG0 |
Coemet_Cmet_C589799_01 | OG0 |
Vmul_sbMO447580 | OG0 |
Lampan_Lpan_C385563_01 | OG0 |
Beraby_Baby_sb331063 | OG0 |
Bolinf_Binf_sb1194925 | OG0 |
Vmul_sbMO374798 | OG0 |
Skow_XM_006812959.1 | OG2 |
Acagra_model.g17242.t1 | OG2 |
Acagra_model.g17242.t1.1.5d3b88c6 | OG2 |
Acagra_model.g19057.t1 | OG2 |
Acagra_model.g19055.t1.1.5d3b897d | OG2 |
Cgig_XM_011452400.3 | OG2 |
Cgig_XM_034472626.1 | OG2 |
Spur_XM_030974469.1 | OG5 |
Hsap_ENST00000379270 | OG5 |
Hsap_ENST00000644105 | OG5 |
Bralan_BLAG14000855_1 | OG5 |
Bralan_BLAG14000161_1 | OG5 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA118145 | OG5 |
Bralan_BLAG17000741_2 | OG5 |
Bralan_BLAG17000740_1 | OG5 |
Bralan_BLAG90000719_1 | OG5 |
Bralan_BLAG05001376_1 | OG5 |
Bralan_BLAG05001380_1 | OG5 |
Bralan_BLAG05001422_1 | OG5 |
Hsap_ENST00000269298 | OG6 |
Spur_XM_003725668.3 | OG6 |
Nvec_v1g237402 | OG7 |
Cgig_XM_034448724.1 | OG7 |
Cgig_XM_034448728.1 | OG7 |
Cgig_XM_034448726.1 | OG7 |
Cgig_XM_034448729.1 | OG7 |
Spur_XM_030983072.1 | OG7 |
Spur_XM_792979.4 | OG7 |
Spur_XM_030990301.1 | OG7 |
Spur_XM_003727723.3 | OG7 |
Spur_XM_030987844.1 | OG7 |
Spur_XM_030989687.1 | OG7 |
Spur_XM_030981387.1 | OG7 |
Spur_XM_030981389.1 | OG7 |
Spur_XM_030981917.1 | OG7 |
Spur_XM_792271.5 | OG7 |
Hvul_g16223_1 | OG1 |
Morvir_scaffold18.g17.t1 | OG1 |
Sanmal_Sma_021925-T1 | OG1 |
Rhoesc_Res_015532-T1 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P015532 | OG1 |
Exapal_XM_021052503.2 | OG1 |
Exapal_XM_021048594.2 | OG1 |
Nvec_v1g204044 | OG1 |
Xesp_015562-T1 | OG1 |
Dgig_XM_028558182.1 | OG1 |
Dgig_XM_028551275.1 | OG1 |
Chem_TCONS_00016933 | OG1 |
Amil_Amillepora24225-RA | OG1 |
Adig_XM_015917395.1 | OG1 |
Amil_Amillepora24226-RA | OG1 |
Gfas_gfas1.m1.14843.m1 | OG1 |
Pocdam_XM_027196738.1 | OG1 |
Spis_XP_022801439_1 | OG1 |
Fspp_ffun1.m4.23021.m1 | OG1 |
Fspp_ffun1.m4.23023.m1 | OG1 |
Gasp_gasp1.m3.14128.m1 | OG1 |
Fspp_ffun1.m4.11809.m1 | OG1 |
Fspp_ffun1.m4.19314.m1 | OG1 |
Fspp_ffun1.m4.19317.m1 | OG1 |
Xestes_maker-XT_scaffold18723-snap-gene-0.18-mRNA-1 | OG3 |
Pbac_sb2665191 | OG3 |
Pbac_sb2667913 | OG3 |
Exapal_XM_021047804.2 | OG3 |
Nvec_v1g202127 | OG3 |
Pocdam_XM_027195191.1 | OG3 |
Gasp_gasp1.m3.7351.m1 | OG3 |
TrH2_TrispH2_008142-RA | OG4 |
Tadh_TriadT51268 | OG4 |
HoiH23_PlH23_008507-RA | OG4 |
HoiH11_mRNA848 | OG4 |
Hhon_g05023.t1 | OG4 |
Hhon_g00514.t1 | OG4 |
HoiH04_mRNA12154 | OG4 |
Emue_Em0021g833a.t1 | OG4 |
Spolac_c102339_g1_i1_m.28931c102339-g1 | OG4 |
Spolac_c102339_g1_i2_m.28933c102339-g1 | OG4 |
Spolac_c102339_g1_i3_m.28934c102339-g1 | OG4 |
Stycar_maker-SC_scaffold2498-augustus-gene-0.14-mRNA-1 | OG4 |
Tetwil_g16139.t2 | OG4 |
Tetwil_g25286.t1 | OG4 |
Halamb_c48515_g1_i2m.24341 | OG4 |
Xestes_maker-XT_scaffold18723-snap-gene-0.17-mRNA-1 | OG4 |
Aque_Aqu2.1.41974_001 | OG4 |
Aque_Aqu2.1.41976_001 | OG4 |
Saldol_EP00054_Salpingoeca_dolichothecata_P005888 | OG4 |
Salmac_EP00052_Salpingoeca_macrocollata_P013639 | OG4 |
Salpun_EP00053_Salpingoeca_punica_P027542 | OG4 |
Salurc_EP00051_Salpingoeca_urceolata_P005455 | OG4 |
Salurc_EP00051_Salpingoeca_urceolata_P005456 | OG4 |
Chofle_EP00730_Choanoeca_flexa_P014646 | OG4 |
Choper_EP00045_Choanoeca_perplexa_P017665 | OG4 |
Salinf_EP00049_Salpingoeca_infusionum_P027639 | OG4 |
Microa_EP00043_Microstomoeca_roanoka_P028461 | OG4 |
Sros_EGD82004 | OG4 |
Opea_EP00118_Oscarella_pearsei_P006430 | OG4 |
Ocar_g4715 | OG4 |
Plajan_Pjan_scaffold10329_01 | OG4 |
Plajan_Pjan_scaffold10332_01 | OG4 |
Plajan_Pjan_scaffold10328_01 | OG4 |
Plajan_Pjan_scaffold10330_01 | OG4 |
Clacor_Ccor_C517419_01 | OG4 |
Gracom_Gcom_scaffold2964_01 | OG4 |
Gracom_Gcom_scaffold2965_01 | OG4 |
Leuniv_Lniv_scaffold20108_01 | OG4 |
Scil_scict016793.1 | OG4 |
Lcom_lcpid80372 | OG4 |
Leuniv_Lniv_scaffold21473_01 | OG4 |
0.416667