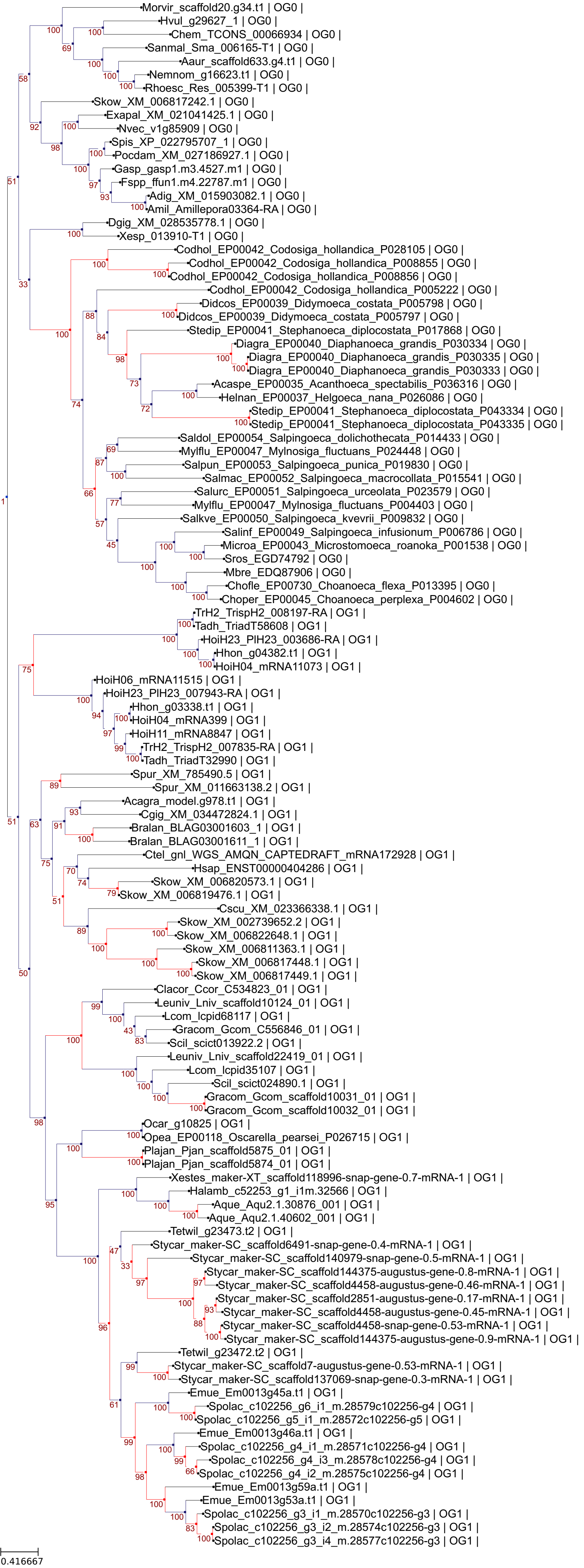
Morvir_scaffold20.g34.t1 | OG0 |
Hvul_g29627_1 | OG0 |
Chem_TCONS_00066934 | OG0 |
Sanmal_Sma_006165-T1 | OG0 |
Aaur_scaffold633.g4.t1 | OG0 |
Nemnom_g16623.t1 | OG0 |
Rhoesc_Res_005399-T1 | OG0 |
Skow_XM_006817242.1 | OG0 |
Exapal_XM_021041425.1 | OG0 |
Nvec_v1g85909 | OG0 |
Spis_XP_022795707_1 | OG0 |
Pocdam_XM_027186927.1 | OG0 |
Gasp_gasp1.m3.4527.m1 | OG0 |
Fspp_ffun1.m4.22787.m1 | OG0 |
Adig_XM_015903082.1 | OG0 |
Amil_Amillepora03364-RA | OG0 |
Dgig_XM_028535778.1 | OG0 |
Xesp_013910-T1 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P028105 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P008855 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P008856 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P005222 | OG0 |
Didcos_EP00039_Didymoeca_costata_P005798 | OG0 |
Didcos_EP00039_Didymoeca_costata_P005797 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P017868 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P030334 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P030335 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P030333 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P036316 | OG0 |
Helnan_EP00037_Helgoeca_nana_P026086 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P043334 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P043335 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P014433 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P024448 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P019830 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P015541 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P023579 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P004403 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P009832 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P006786 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P001538 | OG0 |
Sros_EGD74792 | OG0 |
Mbre_EDQ87906 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P013395 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P004602 | OG0 |
TrH2_TrispH2_008197-RA | OG1 |
Tadh_TriadT58608 | OG1 |
HoiH23_PlH23_003686-RA | OG1 |
Hhon_g04382.t1 | OG1 |
HoiH04_mRNA11073 | OG1 |
HoiH06_mRNA11515 | OG1 |
HoiH23_PlH23_007943-RA | OG1 |
Hhon_g03338.t1 | OG1 |
HoiH04_mRNA399 | OG1 |
HoiH11_mRNA8847 | OG1 |
TrH2_TrispH2_007835-RA | OG1 |
Tadh_TriadT32990 | OG1 |
Spur_XM_785490.5 | OG1 |
Spur_XM_011663138.2 | OG1 |
Acagra_model.g978.t1 | OG1 |
Cgig_XM_034472824.1 | OG1 |
Bralan_BLAG03001603_1 | OG1 |
Bralan_BLAG03001611_1 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA172928 | OG1 |
Hsap_ENST00000404286 | OG1 |
Skow_XM_006820573.1 | OG1 |
Skow_XM_006819476.1 | OG1 |
Cscu_XM_023366338.1 | OG1 |
Skow_XM_002739652.2 | OG1 |
Skow_XM_006822648.1 | OG1 |
Skow_XM_006811363.1 | OG1 |
Skow_XM_006817448.1 | OG1 |
Skow_XM_006817449.1 | OG1 |
Clacor_Ccor_C534823_01 | OG1 |
Leuniv_Lniv_scaffold10124_01 | OG1 |
Lcom_lcpid68117 | OG1 |
Gracom_Gcom_C556846_01 | OG1 |
Scil_scict013922.2 | OG1 |
Leuniv_Lniv_scaffold22419_01 | OG1 |
Lcom_lcpid35107 | OG1 |
Scil_scict024890.1 | OG1 |
Gracom_Gcom_scaffold10031_01 | OG1 |
Gracom_Gcom_scaffold10032_01 | OG1 |
Ocar_g10825 | OG1 |
Opea_EP00118_Oscarella_pearsei_P026715 | OG1 |
Plajan_Pjan_scaffold5875_01 | OG1 |
Plajan_Pjan_scaffold5874_01 | OG1 |
Xestes_maker-XT_scaffold118996-snap-gene-0.7-mRNA-1 | OG1 |
Halamb_c52253_g1_i1m.32566 | OG1 |
Aque_Aqu2.1.30876_001 | OG1 |
Aque_Aqu2.1.40602_001 | OG1 |
Tetwil_g23473.t2 | OG1 |
Stycar_maker-SC_scaffold6491-snap-gene-0.4-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold140979-snap-gene-0.5-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold144375-augustus-gene-0.8-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold4458-augustus-gene-0.46-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold2851-augustus-gene-0.17-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold4458-augustus-gene-0.45-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold4458-snap-gene-0.53-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold144375-augustus-gene-0.9-mRNA-1 | OG1 |
Tetwil_g23472.t2 | OG1 |
Stycar_maker-SC_scaffold7-augustus-gene-0.53-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold137069-snap-gene-0.3-mRNA-1 | OG1 |
Emue_Em0013g45a.t1 | OG1 |
Spolac_c102256_g6_i1_m.28579c102256-g4 | OG1 |
Spolac_c102256_g5_i1_m.28572c102256-g5 | OG1 |
Emue_Em0013g46a.t1 | OG1 |
Spolac_c102256_g4_i1_m.28571c102256-g4 | OG1 |
Spolac_c102256_g4_i3_m.28578c102256-g4 | OG1 |
Spolac_c102256_g4_i2_m.28575c102256-g4 | OG1 |
Emue_Em0013g59a.t1 | OG1 |
Emue_Em0013g53a.t1 | OG1 |
Spolac_c102256_g3_i1_m.28570c102256-g3 | OG1 |
Spolac_c102256_g3_i2_m.28574c102256-g3 | OG1 |
Spolac_c102256_g3_i4_m.28577c102256-g3 | OG1 |
0.416667