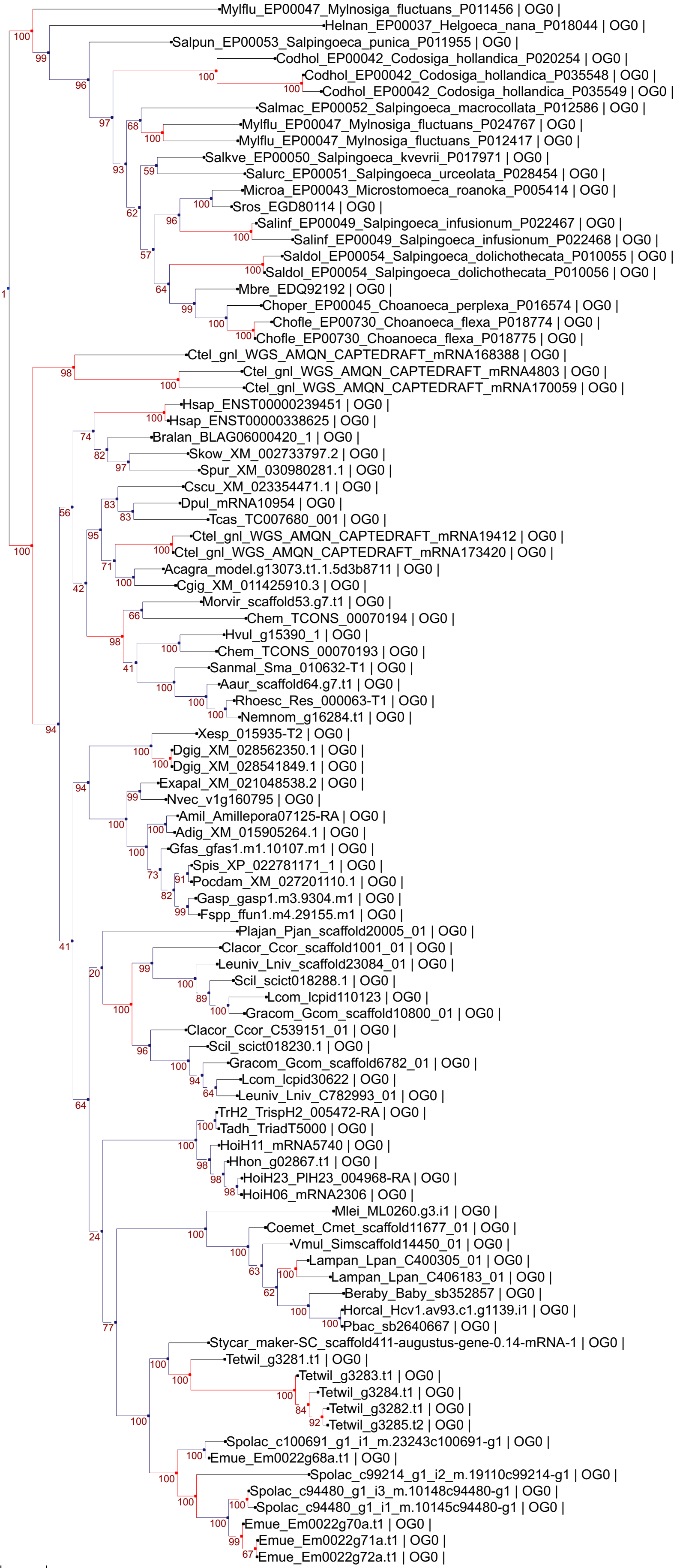
Mylflu_EP00047_Mylnosiga_fluctuans_P011456 | OG0 |
Helnan_EP00037_Helgoeca_nana_P018044 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P011955 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P020254 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P035548 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P035549 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P012586 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P024767 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P012417 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P017971 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P028454 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P005414 | OG0 |
Sros_EGD80114 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P022467 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P022468 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P010055 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P010056 | OG0 |
Mbre_EDQ92192 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P016574 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P018774 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P018775 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA168388 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA4803 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA170059 | OG0 |
Hsap_ENST00000239451 | OG0 |
Hsap_ENST00000338625 | OG0 |
Bralan_BLAG06000420_1 | OG0 |
Skow_XM_002733797.2 | OG0 |
Spur_XM_030980281.1 | OG0 |
Cscu_XM_023354471.1 | OG0 |
Dpul_mRNA10954 | OG0 |
Tcas_TC007680_001 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA19412 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA173420 | OG0 |
Acagra_model.g13073.t1.1.5d3b8711 | OG0 |
Cgig_XM_011425910.3 | OG0 |
Morvir_scaffold53.g7.t1 | OG0 |
Chem_TCONS_00070194 | OG0 |
Hvul_g15390_1 | OG0 |
Chem_TCONS_00070193 | OG0 |
Sanmal_Sma_010632-T1 | OG0 |
Aaur_scaffold64.g7.t1 | OG0 |
Rhoesc_Res_000063-T1 | OG0 |
Nemnom_g16284.t1 | OG0 |
Xesp_015935-T2 | OG0 |
Dgig_XM_028562350.1 | OG0 |
Dgig_XM_028541849.1 | OG0 |
Exapal_XM_021048538.2 | OG0 |
Nvec_v1g160795 | OG0 |
Amil_Amillepora07125-RA | OG0 |
Adig_XM_015905264.1 | OG0 |
Gfas_gfas1.m1.10107.m1 | OG0 |
Spis_XP_022781171_1 | OG0 |
Pocdam_XM_027201110.1 | OG0 |
Gasp_gasp1.m3.9304.m1 | OG0 |
Fspp_ffun1.m4.29155.m1 | OG0 |
Plajan_Pjan_scaffold20005_01 | OG0 |
Clacor_Ccor_scaffold1001_01 | OG0 |
Leuniv_Lniv_scaffold23084_01 | OG0 |
Scil_scict018288.1 | OG0 |
Lcom_lcpid110123 | OG0 |
Gracom_Gcom_scaffold10800_01 | OG0 |
Clacor_Ccor_C539151_01 | OG0 |
Scil_scict018230.1 | OG0 |
Gracom_Gcom_scaffold6782_01 | OG0 |
Lcom_lcpid30622 | OG0 |
Leuniv_Lniv_C782993_01 | OG0 |
TrH2_TrispH2_005472-RA | OG0 |
Tadh_TriadT5000 | OG0 |
HoiH11_mRNA5740 | OG0 |
Hhon_g02867.t1 | OG0 |
HoiH23_PlH23_004968-RA | OG0 |
HoiH06_mRNA2306 | OG0 |
Mlei_ML0260.g3.i1 | OG0 |
Coemet_Cmet_scaffold11677_01 | OG0 |
Vmul_Simscaffold14450_01 | OG0 |
Lampan_Lpan_C400305_01 | OG0 |
Lampan_Lpan_C406183_01 | OG0 |
Beraby_Baby_sb352857 | OG0 |
Horcal_Hcv1.av93.c1.g1139.i1 | OG0 |
Pbac_sb2640667 | OG0 |
Stycar_maker-SC_scaffold411-augustus-gene-0.14-mRNA-1 | OG0 |
Tetwil_g3281.t1 | OG0 |
Tetwil_g3283.t1 | OG0 |
Tetwil_g3284.t1 | OG0 |
Tetwil_g3282.t1 | OG0 |
Tetwil_g3285.t2 | OG0 |
Spolac_c100691_g1_i1_m.23243c100691-g1 | OG0 |
Emue_Em0022g68a.t1 | OG0 |
Spolac_c99214_g1_i2_m.19110c99214-g1 | OG0 |
Spolac_c94480_g1_i3_m.10148c94480-g1 | OG0 |
Spolac_c94480_g1_i1_m.10145c94480-g1 | OG0 |
Emue_Em0022g70a.t1 | OG0 |
Emue_Em0022g71a.t1 | OG0 |
Emue_Em0022g72a.t1 | OG0 |
0.416667