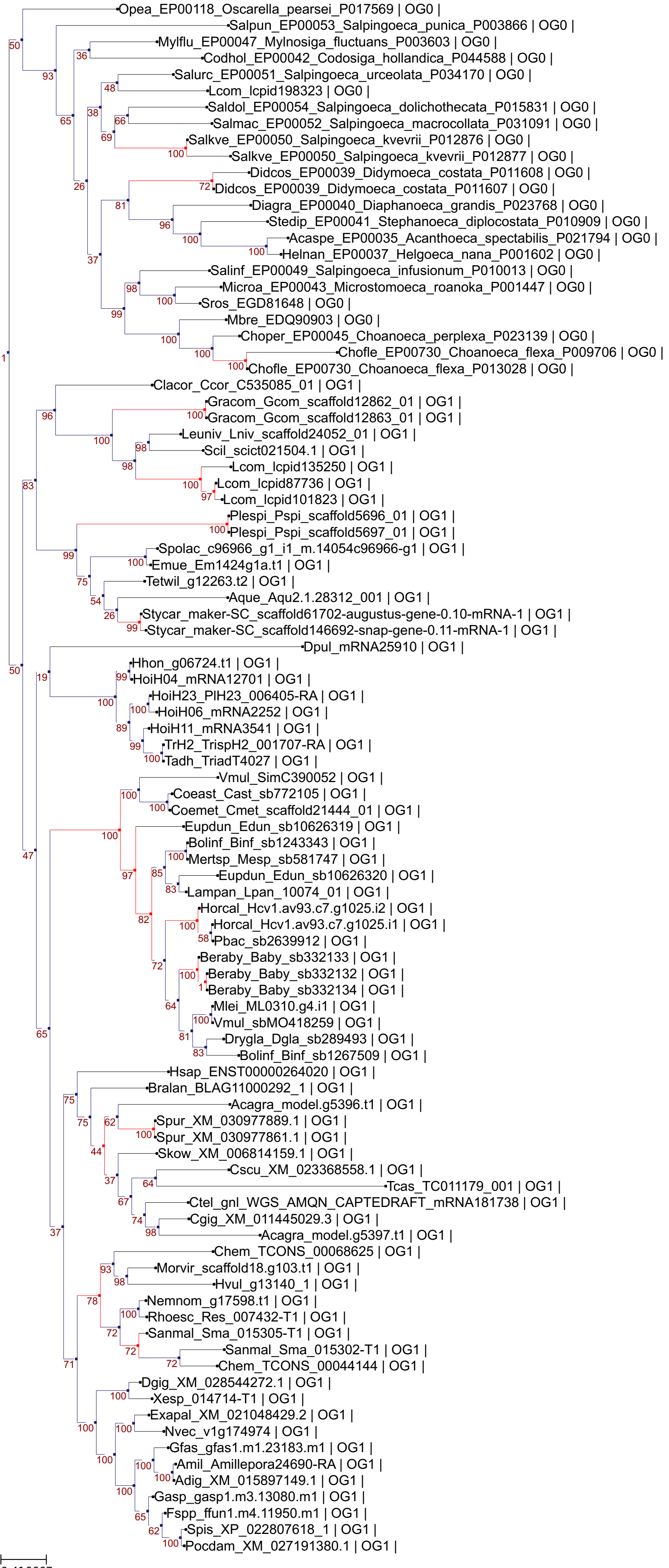
Opea_EP00118_Oscarella_pearsei_P017569 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P003866 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P003603 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P044588 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P034170 | OG0 |
Lcom_lcpid198323 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P015831 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P031091 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P012876 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P012877 | OG0 |
Didcos_EP00039_Didymoeca_costata_P011608 | OG0 |
Didcos_EP00039_Didymoeca_costata_P011607 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P023768 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P010909 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P021794 | OG0 |
Helnan_EP00037_Helgoeca_nana_P001602 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P010013 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P001447 | OG0 |
Sros_EGD81648 | OG0 |
Mbre_EDQ90903 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P023139 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P009706 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P013028 | OG0 |
Clacor_Ccor_C535085_01 | OG1 |
Gracom_Gcom_scaffold12862_01 | OG1 |
Gracom_Gcom_scaffold12863_01 | OG1 |
Leuniv_Lniv_scaffold24052_01 | OG1 |
Scil_scict021504.1 | OG1 |
Lcom_lcpid135250 | OG1 |
Lcom_lcpid87736 | OG1 |
Lcom_lcpid101823 | OG1 |
Plespi_Pspi_scaffold5696_01 | OG1 |
Plespi_Pspi_scaffold5697_01 | OG1 |
Spolac_c96966_g1_i1_m.14054c96966-g1 | OG1 |
Emue_Em1424g1a.t1 | OG1 |
Tetwil_g12263.t2 | OG1 |
Aque_Aqu2.1.28312_001 | OG1 |
Stycar_maker-SC_scaffold61702-augustus-gene-0.10-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold146692-snap-gene-0.11-mRNA-1 | OG1 |
Dpul_mRNA25910 | OG1 |
Hhon_g06724.t1 | OG1 |
HoiH04_mRNA12701 | OG1 |
HoiH23_PlH23_006405-RA | OG1 |
HoiH06_mRNA2252 | OG1 |
HoiH11_mRNA3541 | OG1 |
TrH2_TrispH2_001707-RA | OG1 |
Tadh_TriadT4027 | OG1 |
Vmul_SimC390052 | OG1 |
Coeast_Cast_sb772105 | OG1 |
Coemet_Cmet_scaffold21444_01 | OG1 |
Eupdun_Edun_sb10626319 | OG1 |
Bolinf_Binf_sb1243343 | OG1 |
Mertsp_Mesp_sb581747 | OG1 |
Eupdun_Edun_sb10626320 | OG1 |
Lampan_Lpan_10074_01 | OG1 |
Horcal_Hcv1.av93.c7.g1025.i2 | OG1 |
Horcal_Hcv1.av93.c7.g1025.i1 | OG1 |
Pbac_sb2639912 | OG1 |
Beraby_Baby_sb332133 | OG1 |
Beraby_Baby_sb332132 | OG1 |
Beraby_Baby_sb332134 | OG1 |
Mlei_ML0310.g4.i1 | OG1 |
Vmul_sbMO418259 | OG1 |
Drygla_Dgla_sb289493 | OG1 |
Bolinf_Binf_sb1267509 | OG1 |
Hsap_ENST00000264020 | OG1 |
Bralan_BLAG11000292_1 | OG1 |
Acagra_model.g5396.t1 | OG1 |
Spur_XM_030977889.1 | OG1 |
Spur_XM_030977861.1 | OG1 |
Skow_XM_006814159.1 | OG1 |
Cscu_XM_023368558.1 | OG1 |
Tcas_TC011179_001 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA181738 | OG1 |
Cgig_XM_011445029.3 | OG1 |
Acagra_model.g5397.t1 | OG1 |
Chem_TCONS_00068625 | OG1 |
Morvir_scaffold18.g103.t1 | OG1 |
Hvul_g13140_1 | OG1 |
Nemnom_g17598.t1 | OG1 |
Rhoesc_Res_007432-T1 | OG1 |
Sanmal_Sma_015305-T1 | OG1 |
Sanmal_Sma_015302-T1 | OG1 |
Chem_TCONS_00044144 | OG1 |
Dgig_XM_028544272.1 | OG1 |
Xesp_014714-T1 | OG1 |
Exapal_XM_021048429.2 | OG1 |
Nvec_v1g174974 | OG1 |
Gfas_gfas1.m1.23183.m1 | OG1 |
Amil_Amillepora24690-RA | OG1 |
Adig_XM_015897149.1 | OG1 |
Gasp_gasp1.m3.13080.m1 | OG1 |
Fspp_ffun1.m4.11950.m1 | OG1 |
Spis_XP_022807618_1 | OG1 |
Pocdam_XM_027191380.1 | OG1 |
0.416667