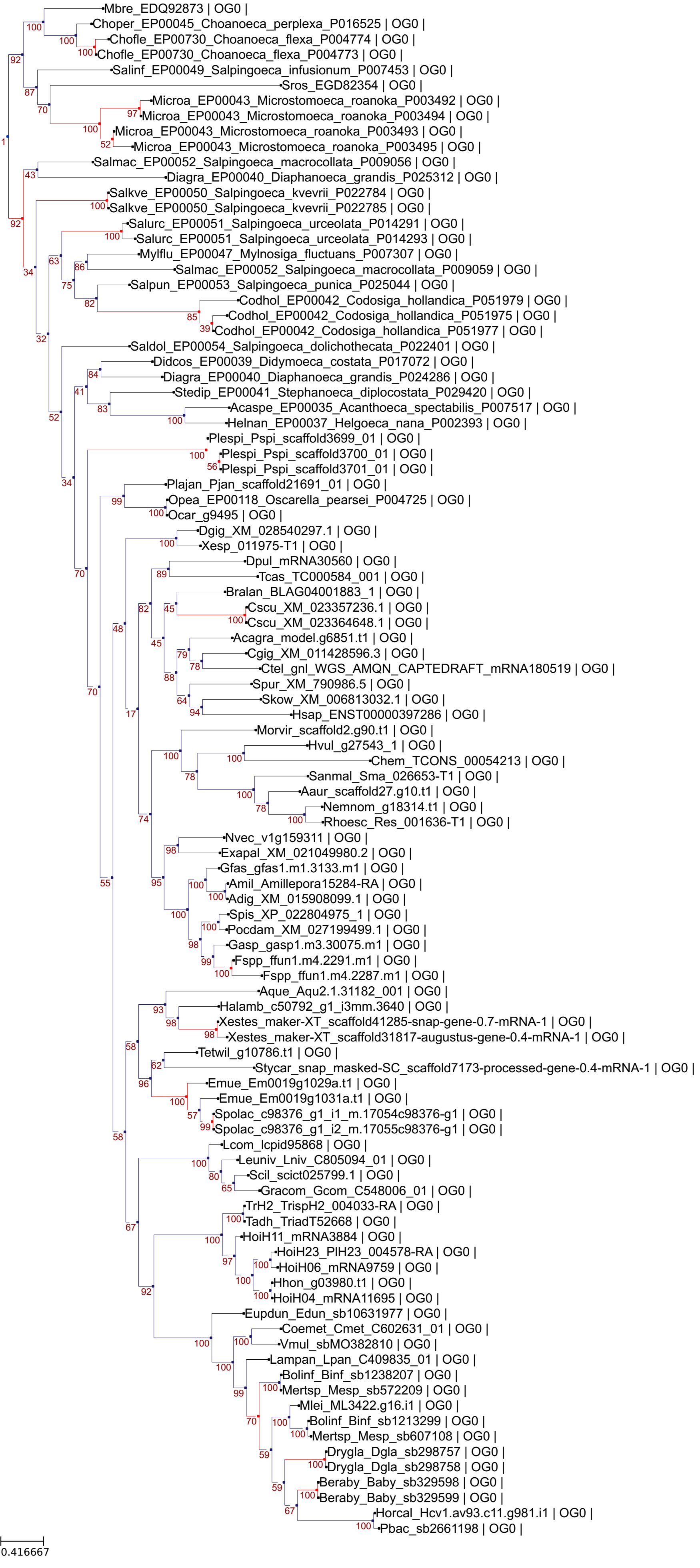
Mbre_EDQ92873 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P016525 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P004774 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P004773 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P007453 | OG0 |
Sros_EGD82354 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P003492 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P003494 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P003493 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P003495 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P009056 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P025312 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P022784 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P022785 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P014291 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P014293 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P007307 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P009059 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P025044 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P051979 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P051975 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P051977 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P022401 | OG0 |
Didcos_EP00039_Didymoeca_costata_P017072 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P024286 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P029420 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P007517 | OG0 |
Helnan_EP00037_Helgoeca_nana_P002393 | OG0 |
Plespi_Pspi_scaffold3699_01 | OG0 |
Plespi_Pspi_scaffold3700_01 | OG0 |
Plespi_Pspi_scaffold3701_01 | OG0 |
Plajan_Pjan_scaffold21691_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P004725 | OG0 |
Ocar_g9495 | OG0 |
Dgig_XM_028540297.1 | OG0 |
Xesp_011975-T1 | OG0 |
Dpul_mRNA30560 | OG0 |
Tcas_TC000584_001 | OG0 |
Bralan_BLAG04001883_1 | OG0 |
Cscu_XM_023357236.1 | OG0 |
Cscu_XM_023364648.1 | OG0 |
Acagra_model.g6851.t1 | OG0 |
Cgig_XM_011428596.3 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA180519 | OG0 |
Spur_XM_790986.5 | OG0 |
Skow_XM_006813032.1 | OG0 |
Hsap_ENST00000397286 | OG0 |
Morvir_scaffold2.g90.t1 | OG0 |
Hvul_g27543_1 | OG0 |
Chem_TCONS_00054213 | OG0 |
Sanmal_Sma_026653-T1 | OG0 |
Aaur_scaffold27.g10.t1 | OG0 |
Nemnom_g18314.t1 | OG0 |
Rhoesc_Res_001636-T1 | OG0 |
Nvec_v1g159311 | OG0 |
Exapal_XM_021049980.2 | OG0 |
Gfas_gfas1.m1.3133.m1 | OG0 |
Amil_Amillepora15284-RA | OG0 |
Adig_XM_015908099.1 | OG0 |
Spis_XP_022804975_1 | OG0 |
Pocdam_XM_027199499.1 | OG0 |
Gasp_gasp1.m3.30075.m1 | OG0 |
Fspp_ffun1.m4.2291.m1 | OG0 |
Fspp_ffun1.m4.2287.m1 | OG0 |
Aque_Aqu2.1.31182_001 | OG0 |
Halamb_c50792_g1_i3mm.3640 | OG0 |
Xestes_maker-XT_scaffold41285-snap-gene-0.7-mRNA-1 | OG0 |
Xestes_maker-XT_scaffold31817-augustus-gene-0.4-mRNA-1 | OG0 |
Tetwil_g10786.t1 | OG0 |
Stycar_snap_masked-SC_scaffold7173-processed-gene-0.4-mRNA-1 | OG0 |
Emue_Em0019g1029a.t1 | OG0 |
Emue_Em0019g1031a.t1 | OG0 |
Spolac_c98376_g1_i1_m.17054c98376-g1 | OG0 |
Spolac_c98376_g1_i2_m.17055c98376-g1 | OG0 |
Lcom_lcpid95868 | OG0 |
Leuniv_Lniv_C805094_01 | OG0 |
Scil_scict025799.1 | OG0 |
Gracom_Gcom_C548006_01 | OG0 |
TrH2_TrispH2_004033-RA | OG0 |
Tadh_TriadT52668 | OG0 |
HoiH11_mRNA3884 | OG0 |
HoiH23_PlH23_004578-RA | OG0 |
HoiH06_mRNA9759 | OG0 |
Hhon_g03980.t1 | OG0 |
HoiH04_mRNA11695 | OG0 |
Eupdun_Edun_sb10631977 | OG0 |
Coemet_Cmet_C602631_01 | OG0 |
Vmul_sbMO382810 | OG0 |
Lampan_Lpan_C409835_01 | OG0 |
Bolinf_Binf_sb1238207 | OG0 |
Mertsp_Mesp_sb572209 | OG0 |
Mlei_ML3422.g16.i1 | OG0 |
Bolinf_Binf_sb1213299 | OG0 |
Mertsp_Mesp_sb607108 | OG0 |
Drygla_Dgla_sb298757 | OG0 |
Drygla_Dgla_sb298758 | OG0 |
Beraby_Baby_sb329598 | OG0 |
Beraby_Baby_sb329599 | OG0 |
Horcal_Hcv1.av93.c11.g981.i1 | OG0 |
Pbac_sb2661198 | OG0 |
0.416667