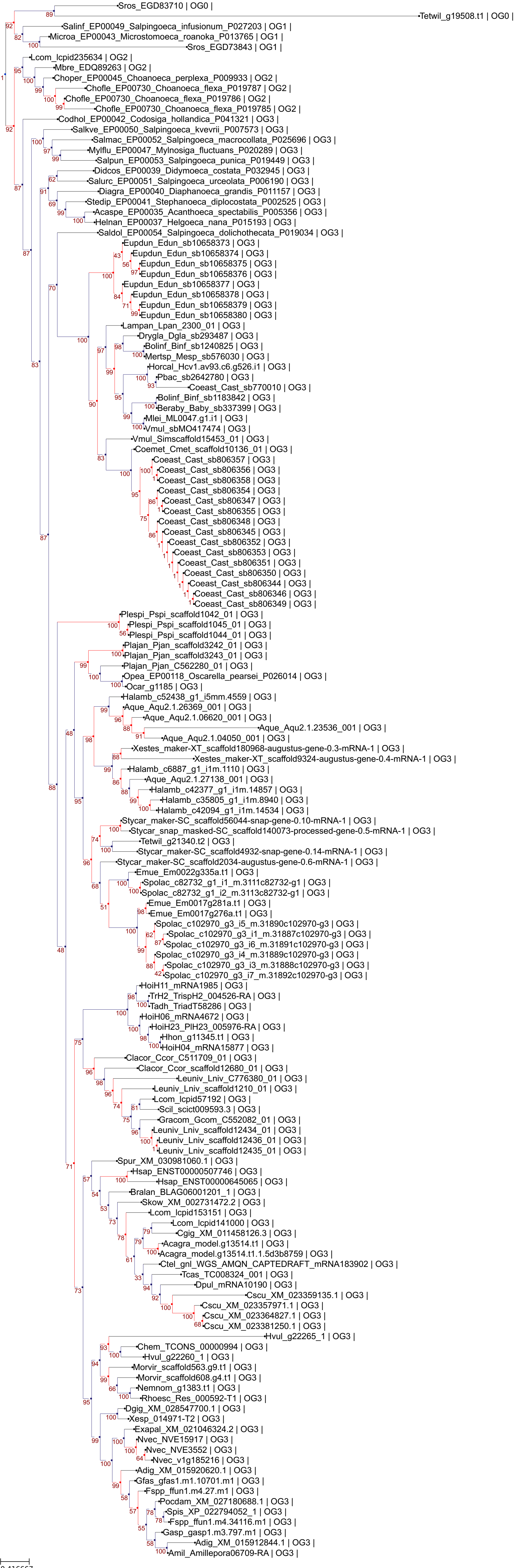
Sros_EGD83710 | OG0 |
Tetwil_g19508.t1 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P027203 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P013765 | OG1 |
Sros_EGD73843 | OG1 |
Lcom_lcpid235634 | OG2 |
Mbre_EDQ89263 | OG2 |
Choper_EP00045_Choanoeca_perplexa_P009933 | OG2 |
Chofle_EP00730_Choanoeca_flexa_P019787 | OG2 |
Chofle_EP00730_Choanoeca_flexa_P019786 | OG2 |
Chofle_EP00730_Choanoeca_flexa_P019785 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P041321 | OG3 |
Salkve_EP00050_Salpingoeca_kvevrii_P007573 | OG3 |
Salmac_EP00052_Salpingoeca_macrocollata_P025696 | OG3 |
Mylflu_EP00047_Mylnosiga_fluctuans_P020289 | OG3 |
Salpun_EP00053_Salpingoeca_punica_P019449 | OG3 |
Didcos_EP00039_Didymoeca_costata_P032945 | OG3 |
Salurc_EP00051_Salpingoeca_urceolata_P006190 | OG3 |
Diagra_EP00040_Diaphanoeca_grandis_P011157 | OG3 |
Stedip_EP00041_Stephanoeca_diplocostata_P002525 | OG3 |
Acaspe_EP00035_Acanthoeca_spectabilis_P005356 | OG3 |
Helnan_EP00037_Helgoeca_nana_P015193 | OG3 |
Saldol_EP00054_Salpingoeca_dolichothecata_P019034 | OG3 |
Eupdun_Edun_sb10658373 | OG3 |
Eupdun_Edun_sb10658374 | OG3 |
Eupdun_Edun_sb10658375 | OG3 |
Eupdun_Edun_sb10658376 | OG3 |
Eupdun_Edun_sb10658377 | OG3 |
Eupdun_Edun_sb10658378 | OG3 |
Eupdun_Edun_sb10658379 | OG3 |
Eupdun_Edun_sb10658380 | OG3 |
Lampan_Lpan_2300_01 | OG3 |
Drygla_Dgla_sb293487 | OG3 |
Bolinf_Binf_sb1240825 | OG3 |
Mertsp_Mesp_sb576030 | OG3 |
Horcal_Hcv1.av93.c6.g526.i1 | OG3 |
Pbac_sb2642780 | OG3 |
Coeast_Cast_sb770010 | OG3 |
Bolinf_Binf_sb1183842 | OG3 |
Beraby_Baby_sb337399 | OG3 |
Mlei_ML0047.g1.i1 | OG3 |
Vmul_sbMO417474 | OG3 |
Vmul_Simscaffold15453_01 | OG3 |
Coemet_Cmet_scaffold10136_01 | OG3 |
Coeast_Cast_sb806357 | OG3 |
Coeast_Cast_sb806356 | OG3 |
Coeast_Cast_sb806358 | OG3 |
Coeast_Cast_sb806354 | OG3 |
Coeast_Cast_sb806347 | OG3 |
Coeast_Cast_sb806355 | OG3 |
Coeast_Cast_sb806348 | OG3 |
Coeast_Cast_sb806345 | OG3 |
Coeast_Cast_sb806352 | OG3 |
Coeast_Cast_sb806353 | OG3 |
Coeast_Cast_sb806351 | OG3 |
Coeast_Cast_sb806350 | OG3 |
Coeast_Cast_sb806344 | OG3 |
Coeast_Cast_sb806346 | OG3 |
Coeast_Cast_sb806349 | OG3 |
Plespi_Pspi_scaffold1042_01 | OG3 |
Plespi_Pspi_scaffold1045_01 | OG3 |
Plespi_Pspi_scaffold1044_01 | OG3 |
Plajan_Pjan_scaffold3242_01 | OG3 |
Plajan_Pjan_scaffold3243_01 | OG3 |
Plajan_Pjan_C562280_01 | OG3 |
Opea_EP00118_Oscarella_pearsei_P026014 | OG3 |
Ocar_g1185 | OG3 |
Halamb_c52438_g1_i5mm.4559 | OG3 |
Aque_Aqu2.1.26369_001 | OG3 |
Aque_Aqu2.1.06620_001 | OG3 |
Aque_Aqu2.1.23536_001 | OG3 |
Aque_Aqu2.1.04050_001 | OG3 |
Xestes_maker-XT_scaffold180968-augustus-gene-0.3-mRNA-1 | OG3 |
Xestes_maker-XT_scaffold9324-augustus-gene-0.4-mRNA-1 | OG3 |
Halamb_c6887_g1_i1m.1110 | OG3 |
Aque_Aqu2.1.27138_001 | OG3 |
Halamb_c42377_g1_i1m.14857 | OG3 |
Halamb_c35805_g1_i1m.8940 | OG3 |
Halamb_c42094_g1_i1m.14534 | OG3 |
Stycar_maker-SC_scaffold56044-snap-gene-0.10-mRNA-1 | OG3 |
Stycar_snap_masked-SC_scaffold140073-processed-gene-0.5-mRNA-1 | OG3 |
Tetwil_g21340.t2 | OG3 |
Stycar_maker-SC_scaffold4932-snap-gene-0.14-mRNA-1 | OG3 |
Stycar_maker-SC_scaffold2034-augustus-gene-0.6-mRNA-1 | OG3 |
Emue_Em0022g335a.t1 | OG3 |
Spolac_c82732_g1_i1_m.3111c82732-g1 | OG3 |
Spolac_c82732_g1_i2_m.3113c82732-g1 | OG3 |
Emue_Em0017g281a.t1 | OG3 |
Emue_Em0017g276a.t1 | OG3 |
Spolac_c102970_g3_i5_m.31890c102970-g3 | OG3 |
Spolac_c102970_g3_i1_m.31887c102970-g3 | OG3 |
Spolac_c102970_g3_i6_m.31891c102970-g3 | OG3 |
Spolac_c102970_g3_i4_m.31889c102970-g3 | OG3 |
Spolac_c102970_g3_i3_m.31888c102970-g3 | OG3 |
Spolac_c102970_g3_i7_m.31892c102970-g3 | OG3 |
HoiH11_mRNA1985 | OG3 |
TrH2_TrispH2_004526-RA | OG3 |
Tadh_TriadT58286 | OG3 |
HoiH06_mRNA4672 | OG3 |
HoiH23_PlH23_005976-RA | OG3 |
Hhon_g11345.t1 | OG3 |
HoiH04_mRNA15877 | OG3 |
Clacor_Ccor_C511709_01 | OG3 |
Clacor_Ccor_scaffold12680_01 | OG3 |
Leuniv_Lniv_C776380_01 | OG3 |
Leuniv_Lniv_scaffold1210_01 | OG3 |
Lcom_lcpid57192 | OG3 |
Scil_scict009593.3 | OG3 |
Gracom_Gcom_C552082_01 | OG3 |
Leuniv_Lniv_scaffold12434_01 | OG3 |
Leuniv_Lniv_scaffold12436_01 | OG3 |
Leuniv_Lniv_scaffold12435_01 | OG3 |
Spur_XM_030981060.1 | OG3 |
Hsap_ENST00000507746 | OG3 |
Hsap_ENST00000645065 | OG3 |
Bralan_BLAG06001201_1 | OG3 |
Skow_XM_002731472.2 | OG3 |
Lcom_lcpid153151 | OG3 |
Lcom_lcpid141000 | OG3 |
Cgig_XM_011458126.3 | OG3 |
Acagra_model.g13514.t1 | OG3 |
Acagra_model.g13514.t1.1.5d3b8759 | OG3 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA183902 | OG3 |
Tcas_TC008324_001 | OG3 |
Dpul_mRNA10190 | OG3 |
Cscu_XM_023359135.1 | OG3 |
Cscu_XM_023357971.1 | OG3 |
Cscu_XM_023364827.1 | OG3 |
Cscu_XM_023381250.1 | OG3 |
Hvul_g22265_1 | OG3 |
Chem_TCONS_00000994 | OG3 |
Hvul_g22260_1 | OG3 |
Morvir_scaffold563.g9.t1 | OG3 |
Morvir_scaffold608.g4.t1 | OG3 |
Nemnom_g1383.t1 | OG3 |
Rhoesc_Res_000592-T1 | OG3 |
Dgig_XM_028547700.1 | OG3 |
Xesp_014971-T2 | OG3 |
Exapal_XM_021046324.2 | OG3 |
Nvec_NVE15917 | OG3 |
Nvec_NVE3552 | OG3 |
Nvec_v1g185216 | OG3 |
Adig_XM_015920620.1 | OG3 |
Gfas_gfas1.m1.10701.m1 | OG3 |
Fspp_ffun1.m4.27.m1 | OG3 |
Pocdam_XM_027180688.1 | OG3 |
Spis_XP_022794052_1 | OG3 |
Fspp_ffun1.m4.34116.m1 | OG3 |
Gasp_gasp1.m3.797.m1 | OG3 |
Adig_XM_015912844.1 | OG3 |
Amil_Amillepora06709-RA | OG3 |
0.416667