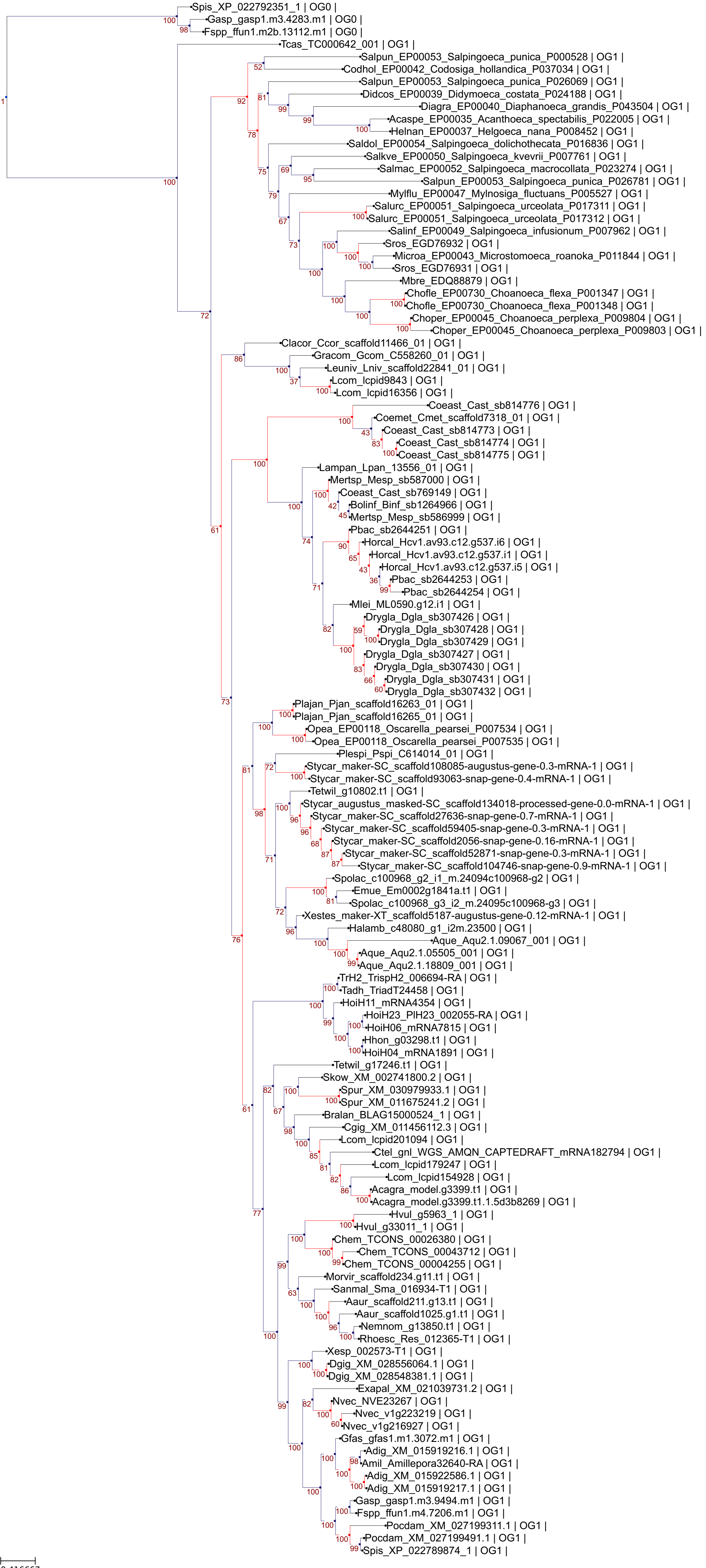
Spis_XP_022792351_1 | OG0 |
Gasp_gasp1.m3.4283.m1 | OG0 |
Fspp_ffun1.m2b.13112.m1 | OG0 |
Tcas_TC000642_001 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P000528 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P037034 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P026069 | OG1 |
Didcos_EP00039_Didymoeca_costata_P024188 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P043504 | OG1 |
Acaspe_EP00035_Acanthoeca_spectabilis_P022005 | OG1 |
Helnan_EP00037_Helgoeca_nana_P008452 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P016836 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P007761 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P023274 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P026781 | OG1 |
Mylflu_EP00047_Mylnosiga_fluctuans_P005527 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P017311 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P017312 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P007962 | OG1 |
Sros_EGD76932 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P011844 | OG1 |
Sros_EGD76931 | OG1 |
Mbre_EDQ88879 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P001347 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P001348 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P009804 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P009803 | OG1 |
Clacor_Ccor_scaffold11466_01 | OG1 |
Gracom_Gcom_C558260_01 | OG1 |
Leuniv_Lniv_scaffold22841_01 | OG1 |
Lcom_lcpid9843 | OG1 |
Lcom_lcpid16356 | OG1 |
Coeast_Cast_sb814776 | OG1 |
Coemet_Cmet_scaffold7318_01 | OG1 |
Coeast_Cast_sb814773 | OG1 |
Coeast_Cast_sb814774 | OG1 |
Coeast_Cast_sb814775 | OG1 |
Lampan_Lpan_13556_01 | OG1 |
Mertsp_Mesp_sb587000 | OG1 |
Coeast_Cast_sb769149 | OG1 |
Bolinf_Binf_sb1264966 | OG1 |
Mertsp_Mesp_sb586999 | OG1 |
Pbac_sb2644251 | OG1 |
Horcal_Hcv1.av93.c12.g537.i6 | OG1 |
Horcal_Hcv1.av93.c12.g537.i1 | OG1 |
Horcal_Hcv1.av93.c12.g537.i5 | OG1 |
Pbac_sb2644253 | OG1 |
Pbac_sb2644254 | OG1 |
Mlei_ML0590.g12.i1 | OG1 |
Drygla_Dgla_sb307426 | OG1 |
Drygla_Dgla_sb307428 | OG1 |
Drygla_Dgla_sb307429 | OG1 |
Drygla_Dgla_sb307427 | OG1 |
Drygla_Dgla_sb307430 | OG1 |
Drygla_Dgla_sb307431 | OG1 |
Drygla_Dgla_sb307432 | OG1 |
Plajan_Pjan_scaffold16263_01 | OG1 |
Plajan_Pjan_scaffold16265_01 | OG1 |
Opea_EP00118_Oscarella_pearsei_P007534 | OG1 |
Opea_EP00118_Oscarella_pearsei_P007535 | OG1 |
Plespi_Pspi_C614014_01 | OG1 |
Stycar_maker-SC_scaffold108085-augustus-gene-0.3-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold93063-snap-gene-0.4-mRNA-1 | OG1 |
Tetwil_g10802.t1 | OG1 |
Stycar_augustus_masked-SC_scaffold134018-processed-gene-0.0-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold27636-snap-gene-0.7-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold59405-snap-gene-0.3-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold2056-snap-gene-0.16-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold52871-snap-gene-0.3-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold104746-snap-gene-0.9-mRNA-1 | OG1 |
Spolac_c100968_g2_i1_m.24094c100968-g2 | OG1 |
Emue_Em0002g1841a.t1 | OG1 |
Spolac_c100968_g3_i2_m.24095c100968-g3 | OG1 |
Xestes_maker-XT_scaffold5187-augustus-gene-0.12-mRNA-1 | OG1 |
Halamb_c48080_g1_i2m.23500 | OG1 |
Aque_Aqu2.1.09067_001 | OG1 |
Aque_Aqu2.1.05505_001 | OG1 |
Aque_Aqu2.1.18809_001 | OG1 |
TrH2_TrispH2_006694-RA | OG1 |
Tadh_TriadT24458 | OG1 |
HoiH11_mRNA4354 | OG1 |
HoiH23_PlH23_002055-RA | OG1 |
HoiH06_mRNA7815 | OG1 |
Hhon_g03298.t1 | OG1 |
HoiH04_mRNA1891 | OG1 |
Tetwil_g17246.t1 | OG1 |
Skow_XM_002741800.2 | OG1 |
Spur_XM_030979933.1 | OG1 |
Spur_XM_011675241.2 | OG1 |
Bralan_BLAG15000524_1 | OG1 |
Cgig_XM_011456112.3 | OG1 |
Lcom_lcpid201094 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA182794 | OG1 |
Lcom_lcpid179247 | OG1 |
Lcom_lcpid154928 | OG1 |
Acagra_model.g3399.t1 | OG1 |
Acagra_model.g3399.t1.1.5d3b8269 | OG1 |
Hvul_g5963_1 | OG1 |
Hvul_g33011_1 | OG1 |
Chem_TCONS_00026380 | OG1 |
Chem_TCONS_00043712 | OG1 |
Chem_TCONS_00004255 | OG1 |
Morvir_scaffold234.g11.t1 | OG1 |
Sanmal_Sma_016934-T1 | OG1 |
Aaur_scaffold211.g13.t1 | OG1 |
Aaur_scaffold1025.g1.t1 | OG1 |
Nemnom_g13850.t1 | OG1 |
Rhoesc_Res_012365-T1 | OG1 |
Xesp_002573-T1 | OG1 |
Dgig_XM_028556064.1 | OG1 |
Dgig_XM_028548381.1 | OG1 |
Exapal_XM_021039731.2 | OG1 |
Nvec_NVE23267 | OG1 |
Nvec_v1g223219 | OG1 |
Nvec_v1g216927 | OG1 |
Gfas_gfas1.m1.3072.m1 | OG1 |
Adig_XM_015919216.1 | OG1 |
Amil_Amillepora32640-RA | OG1 |
Adig_XM_015922586.1 | OG1 |
Adig_XM_015919217.1 | OG1 |
Gasp_gasp1.m3.9494.m1 | OG1 |
Fspp_ffun1.m4.7206.m1 | OG1 |
Pocdam_XM_027199311.1 | OG1 |
Pocdam_XM_027199491.1 | OG1 |
Spis_XP_022789874_1 | OG1 |
0.416667