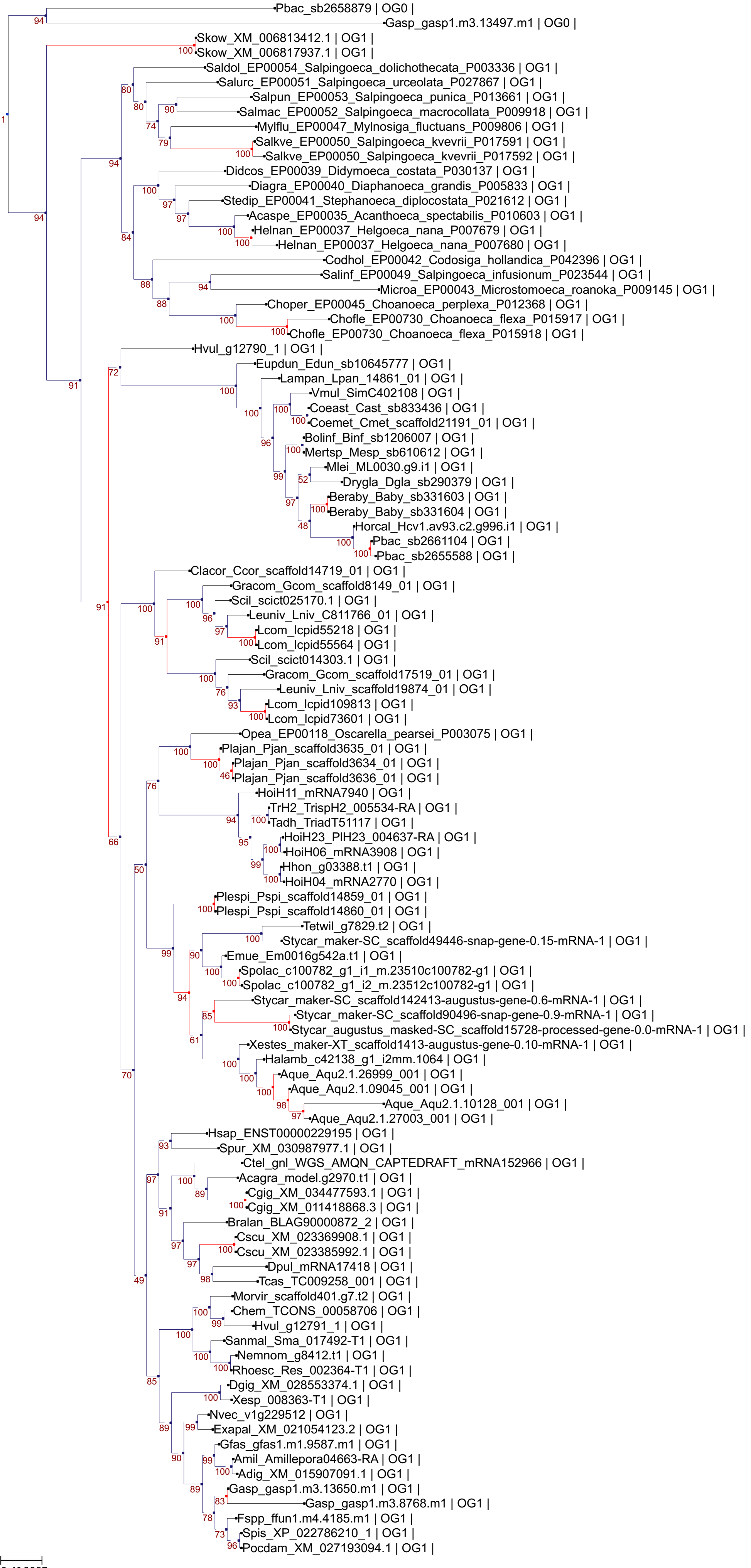
Pbac_sb2658879 | OG0 |
Gasp_gasp1.m3.13497.m1 | OG0 |
Skow_XM_006813412.1 | OG1 |
Skow_XM_006817937.1 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P003336 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P027867 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P013661 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P009918 | OG1 |
Mylflu_EP00047_Mylnosiga_fluctuans_P009806 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P017591 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P017592 | OG1 |
Didcos_EP00039_Didymoeca_costata_P030137 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P005833 | OG1 |
Stedip_EP00041_Stephanoeca_diplocostata_P021612 | OG1 |
Acaspe_EP00035_Acanthoeca_spectabilis_P010603 | OG1 |
Helnan_EP00037_Helgoeca_nana_P007679 | OG1 |
Helnan_EP00037_Helgoeca_nana_P007680 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P042396 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P023544 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P009145 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P012368 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P015917 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P015918 | OG1 |
Hvul_g12790_1 | OG1 |
Eupdun_Edun_sb10645777 | OG1 |
Lampan_Lpan_14861_01 | OG1 |
Vmul_SimC402108 | OG1 |
Coeast_Cast_sb833436 | OG1 |
Coemet_Cmet_scaffold21191_01 | OG1 |
Bolinf_Binf_sb1206007 | OG1 |
Mertsp_Mesp_sb610612 | OG1 |
Mlei_ML0030.g9.i1 | OG1 |
Drygla_Dgla_sb290379 | OG1 |
Beraby_Baby_sb331603 | OG1 |
Beraby_Baby_sb331604 | OG1 |
Horcal_Hcv1.av93.c2.g996.i1 | OG1 |
Pbac_sb2661104 | OG1 |
Pbac_sb2655588 | OG1 |
Clacor_Ccor_scaffold14719_01 | OG1 |
Gracom_Gcom_scaffold8149_01 | OG1 |
Scil_scict025170.1 | OG1 |
Leuniv_Lniv_C811766_01 | OG1 |
Lcom_lcpid55218 | OG1 |
Lcom_lcpid55564 | OG1 |
Scil_scict014303.1 | OG1 |
Gracom_Gcom_scaffold17519_01 | OG1 |
Leuniv_Lniv_scaffold19874_01 | OG1 |
Lcom_lcpid109813 | OG1 |
Lcom_lcpid73601 | OG1 |
Opea_EP00118_Oscarella_pearsei_P003075 | OG1 |
Plajan_Pjan_scaffold3635_01 | OG1 |
Plajan_Pjan_scaffold3634_01 | OG1 |
Plajan_Pjan_scaffold3636_01 | OG1 |
HoiH11_mRNA7940 | OG1 |
TrH2_TrispH2_005534-RA | OG1 |
Tadh_TriadT51117 | OG1 |
HoiH23_PlH23_004637-RA | OG1 |
HoiH06_mRNA3908 | OG1 |
Hhon_g03388.t1 | OG1 |
HoiH04_mRNA2770 | OG1 |
Plespi_Pspi_scaffold14859_01 | OG1 |
Plespi_Pspi_scaffold14860_01 | OG1 |
Tetwil_g7829.t2 | OG1 |
Stycar_maker-SC_scaffold49446-snap-gene-0.15-mRNA-1 | OG1 |
Emue_Em0016g542a.t1 | OG1 |
Spolac_c100782_g1_i1_m.23510c100782-g1 | OG1 |
Spolac_c100782_g1_i2_m.23512c100782-g1 | OG1 |
Stycar_maker-SC_scaffold142413-augustus-gene-0.6-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold90496-snap-gene-0.9-mRNA-1 | OG1 |
Stycar_augustus_masked-SC_scaffold15728-processed-gene-0.0-mRNA-1 | OG1 |
Xestes_maker-XT_scaffold1413-augustus-gene-0.10-mRNA-1 | OG1 |
Halamb_c42138_g1_i2mm.1064 | OG1 |
Aque_Aqu2.1.26999_001 | OG1 |
Aque_Aqu2.1.09045_001 | OG1 |
Aque_Aqu2.1.10128_001 | OG1 |
Aque_Aqu2.1.27003_001 | OG1 |
Hsap_ENST00000229195 | OG1 |
Spur_XM_030987977.1 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA152966 | OG1 |
Acagra_model.g2970.t1 | OG1 |
Cgig_XM_034477593.1 | OG1 |
Cgig_XM_011418868.3 | OG1 |
Bralan_BLAG90000872_2 | OG1 |
Cscu_XM_023369908.1 | OG1 |
Cscu_XM_023385992.1 | OG1 |
Dpul_mRNA17418 | OG1 |
Tcas_TC009258_001 | OG1 |
Morvir_scaffold401.g7.t2 | OG1 |
Chem_TCONS_00058706 | OG1 |
Hvul_g12791_1 | OG1 |
Sanmal_Sma_017492-T1 | OG1 |
Nemnom_g8412.t1 | OG1 |
Rhoesc_Res_002364-T1 | OG1 |
Dgig_XM_028553374.1 | OG1 |
Xesp_008363-T1 | OG1 |
Nvec_v1g229512 | OG1 |
Exapal_XM_021054123.2 | OG1 |
Gfas_gfas1.m1.9587.m1 | OG1 |
Amil_Amillepora04663-RA | OG1 |
Adig_XM_015907091.1 | OG1 |
Gasp_gasp1.m3.13650.m1 | OG1 |
Gasp_gasp1.m3.8768.m1 | OG1 |
Fspp_ffun1.m4.4185.m1 | OG1 |
Spis_XP_022786210_1 | OG1 |
Pocdam_XM_027193094.1 | OG1 |
0.416667