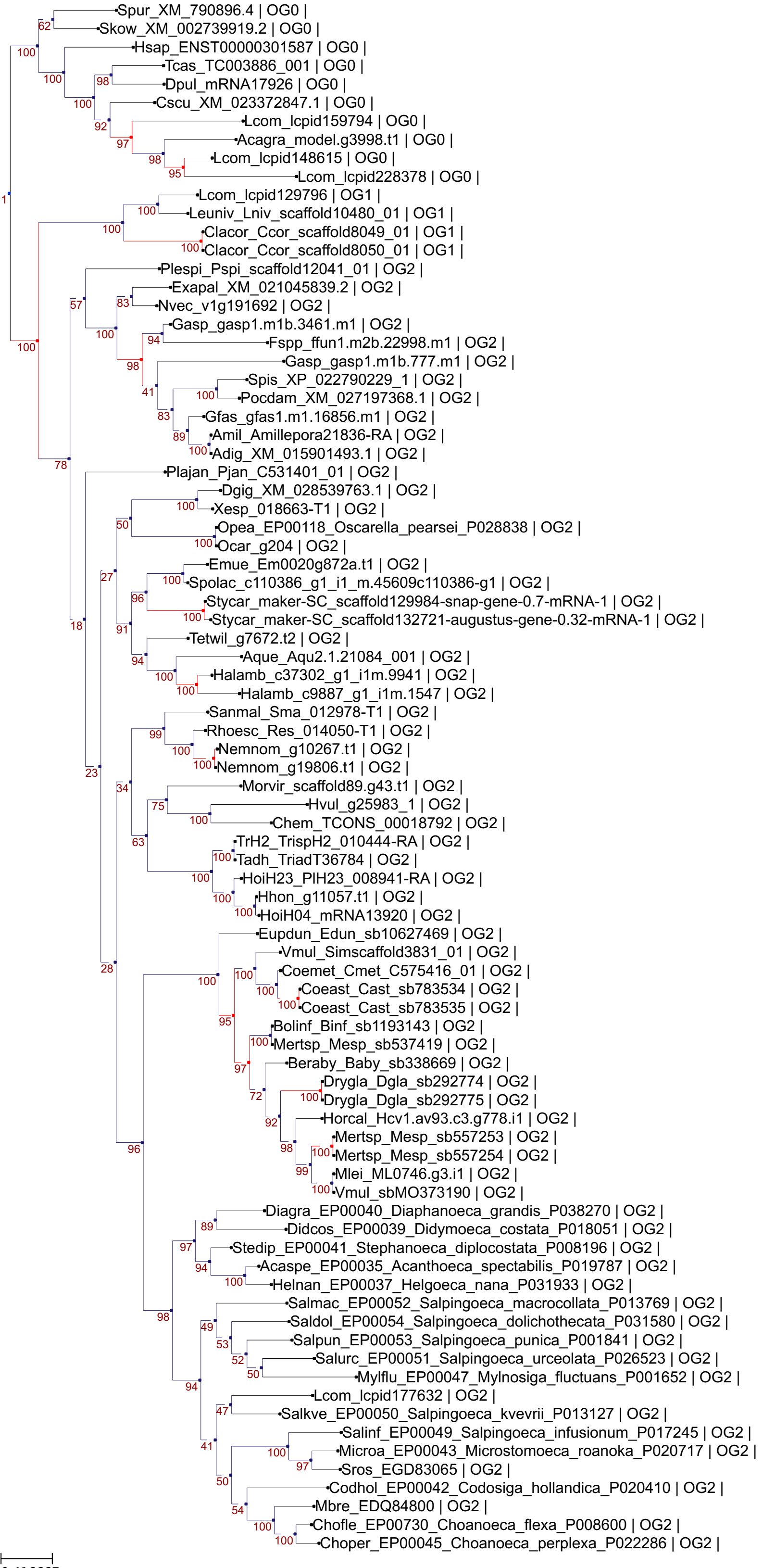
Spur_XM_790896.4 | OG0 |
Skow_XM_002739919.2 | OG0 |
Hsap_ENST00000301587 | OG0 |
Tcas_TC003886_001 | OG0 |
Dpul_mRNA17926 | OG0 |
Cscu_XM_023372847.1 | OG0 |
Lcom_lcpid159794 | OG0 |
Acagra_model.g3998.t1 | OG0 |
Lcom_lcpid148615 | OG0 |
Lcom_lcpid228378 | OG0 |
Lcom_lcpid129796 | OG1 |
Leuniv_Lniv_scaffold10480_01 | OG1 |
Clacor_Ccor_scaffold8049_01 | OG1 |
Clacor_Ccor_scaffold8050_01 | OG1 |
Plespi_Pspi_scaffold12041_01 | OG2 |
Exapal_XM_021045839.2 | OG2 |
Nvec_v1g191692 | OG2 |
Gasp_gasp1.m1b.3461.m1 | OG2 |
Fspp_ffun1.m2b.22998.m1 | OG2 |
Gasp_gasp1.m1b.777.m1 | OG2 |
Spis_XP_022790229_1 | OG2 |
Pocdam_XM_027197368.1 | OG2 |
Gfas_gfas1.m1.16856.m1 | OG2 |
Amil_Amillepora21836-RA | OG2 |
Adig_XM_015901493.1 | OG2 |
Plajan_Pjan_C531401_01 | OG2 |
Dgig_XM_028539763.1 | OG2 |
Xesp_018663-T1 | OG2 |
Opea_EP00118_Oscarella_pearsei_P028838 | OG2 |
Ocar_g204 | OG2 |
Emue_Em0020g872a.t1 | OG2 |
Spolac_c110386_g1_i1_m.45609c110386-g1 | OG2 |
Stycar_maker-SC_scaffold129984-snap-gene-0.7-mRNA-1 | OG2 |
Stycar_maker-SC_scaffold132721-augustus-gene-0.32-mRNA-1 | OG2 |
Tetwil_g7672.t2 | OG2 |
Aque_Aqu2.1.21084_001 | OG2 |
Halamb_c37302_g1_i1m.9941 | OG2 |
Halamb_c9887_g1_i1m.1547 | OG2 |
Sanmal_Sma_012978-T1 | OG2 |
Rhoesc_Res_014050-T1 | OG2 |
Nemnom_g10267.t1 | OG2 |
Nemnom_g19806.t1 | OG2 |
Morvir_scaffold89.g43.t1 | OG2 |
Hvul_g25983_1 | OG2 |
Chem_TCONS_00018792 | OG2 |
TrH2_TrispH2_010444-RA | OG2 |
Tadh_TriadT36784 | OG2 |
HoiH23_PlH23_008941-RA | OG2 |
Hhon_g11057.t1 | OG2 |
HoiH04_mRNA13920 | OG2 |
Eupdun_Edun_sb10627469 | OG2 |
Vmul_Simscaffold3831_01 | OG2 |
Coemet_Cmet_C575416_01 | OG2 |
Coeast_Cast_sb783534 | OG2 |
Coeast_Cast_sb783535 | OG2 |
Bolinf_Binf_sb1193143 | OG2 |
Mertsp_Mesp_sb537419 | OG2 |
Beraby_Baby_sb338669 | OG2 |
Drygla_Dgla_sb292774 | OG2 |
Drygla_Dgla_sb292775 | OG2 |
Horcal_Hcv1.av93.c3.g778.i1 | OG2 |
Mertsp_Mesp_sb557253 | OG2 |
Mertsp_Mesp_sb557254 | OG2 |
Mlei_ML0746.g3.i1 | OG2 |
Vmul_sbMO373190 | OG2 |
Diagra_EP00040_Diaphanoeca_grandis_P038270 | OG2 |
Didcos_EP00039_Didymoeca_costata_P018051 | OG2 |
Stedip_EP00041_Stephanoeca_diplocostata_P008196 | OG2 |
Acaspe_EP00035_Acanthoeca_spectabilis_P019787 | OG2 |
Helnan_EP00037_Helgoeca_nana_P031933 | OG2 |
Salmac_EP00052_Salpingoeca_macrocollata_P013769 | OG2 |
Saldol_EP00054_Salpingoeca_dolichothecata_P031580 | OG2 |
Salpun_EP00053_Salpingoeca_punica_P001841 | OG2 |
Salurc_EP00051_Salpingoeca_urceolata_P026523 | OG2 |
Mylflu_EP00047_Mylnosiga_fluctuans_P001652 | OG2 |
Lcom_lcpid177632 | OG2 |
Salkve_EP00050_Salpingoeca_kvevrii_P013127 | OG2 |
Salinf_EP00049_Salpingoeca_infusionum_P017245 | OG2 |
Microa_EP00043_Microstomoeca_roanoka_P020717 | OG2 |
Sros_EGD83065 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P020410 | OG2 |
Mbre_EDQ84800 | OG2 |
Chofle_EP00730_Choanoeca_flexa_P008600 | OG2 |
Choper_EP00045_Choanoeca_perplexa_P022286 | OG2 |
0.416667