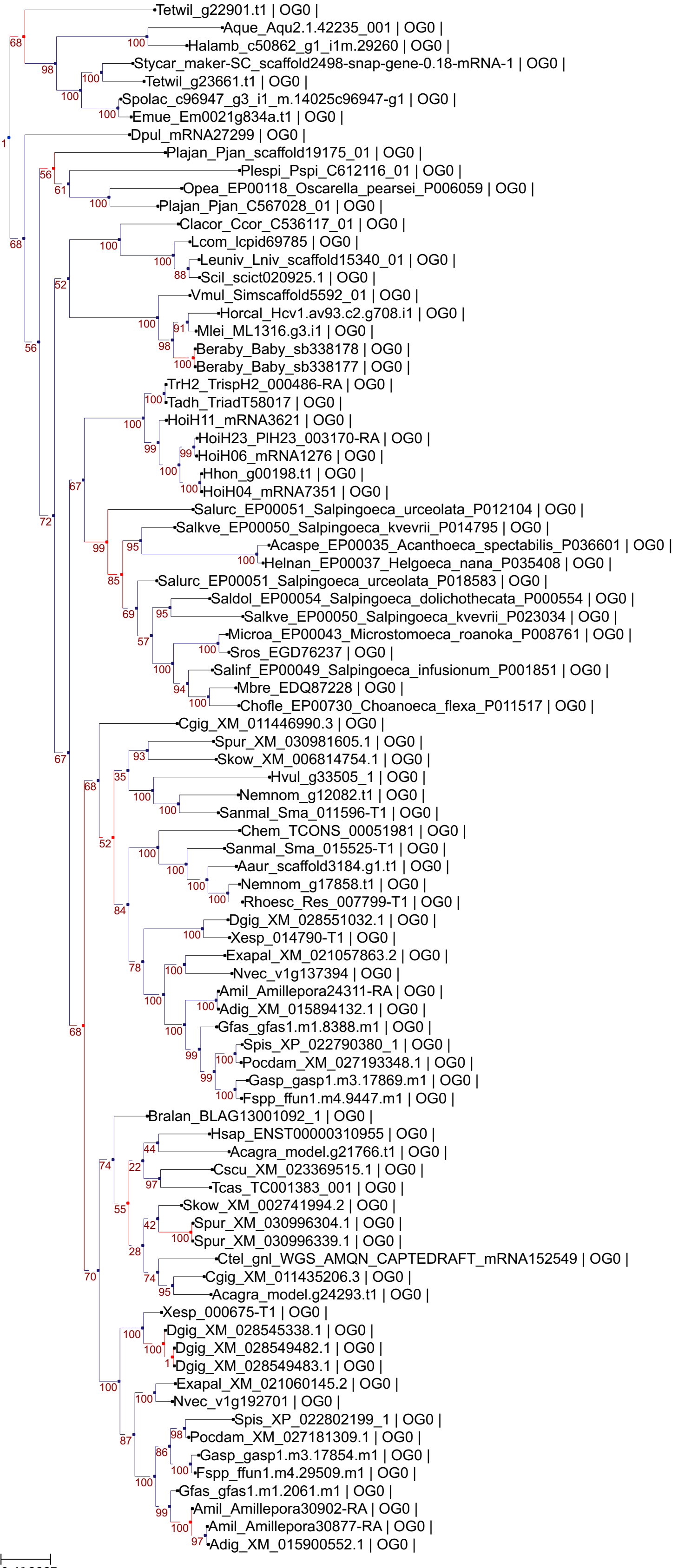
Tetwil_g22901.t1 | OG0 |
Aque_Aqu2.1.42235_001 | OG0 |
Halamb_c50862_g1_i1m.29260 | OG0 |
Stycar_maker-SC_scaffold2498-snap-gene-0.18-mRNA-1 | OG0 |
Tetwil_g23661.t1 | OG0 |
Spolac_c96947_g3_i1_m.14025c96947-g1 | OG0 |
Emue_Em0021g834a.t1 | OG0 |
Dpul_mRNA27299 | OG0 |
Plajan_Pjan_scaffold19175_01 | OG0 |
Plespi_Pspi_C612116_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P006059 | OG0 |
Plajan_Pjan_C567028_01 | OG0 |
Clacor_Ccor_C536117_01 | OG0 |
Lcom_lcpid69785 | OG0 |
Leuniv_Lniv_scaffold15340_01 | OG0 |
Scil_scict020925.1 | OG0 |
Vmul_Simscaffold5592_01 | OG0 |
Horcal_Hcv1.av93.c2.g708.i1 | OG0 |
Mlei_ML1316.g3.i1 | OG0 |
Beraby_Baby_sb338178 | OG0 |
Beraby_Baby_sb338177 | OG0 |
TrH2_TrispH2_000486-RA | OG0 |
Tadh_TriadT58017 | OG0 |
HoiH11_mRNA3621 | OG0 |
HoiH23_PlH23_003170-RA | OG0 |
HoiH06_mRNA1276 | OG0 |
Hhon_g00198.t1 | OG0 |
HoiH04_mRNA7351 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P012104 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P014795 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P036601 | OG0 |
Helnan_EP00037_Helgoeca_nana_P035408 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P018583 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P000554 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P023034 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P008761 | OG0 |
Sros_EGD76237 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P001851 | OG0 |
Mbre_EDQ87228 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P011517 | OG0 |
Cgig_XM_011446990.3 | OG0 |
Spur_XM_030981605.1 | OG0 |
Skow_XM_006814754.1 | OG0 |
Hvul_g33505_1 | OG0 |
Nemnom_g12082.t1 | OG0 |
Sanmal_Sma_011596-T1 | OG0 |
Chem_TCONS_00051981 | OG0 |
Sanmal_Sma_015525-T1 | OG0 |
Aaur_scaffold3184.g1.t1 | OG0 |
Nemnom_g17858.t1 | OG0 |
Rhoesc_Res_007799-T1 | OG0 |
Dgig_XM_028551032.1 | OG0 |
Xesp_014790-T1 | OG0 |
Exapal_XM_021057863.2 | OG0 |
Nvec_v1g137394 | OG0 |
Amil_Amillepora24311-RA | OG0 |
Adig_XM_015894132.1 | OG0 |
Gfas_gfas1.m1.8388.m1 | OG0 |
Spis_XP_022790380_1 | OG0 |
Pocdam_XM_027193348.1 | OG0 |
Gasp_gasp1.m3.17869.m1 | OG0 |
Fspp_ffun1.m4.9447.m1 | OG0 |
Bralan_BLAG13001092_1 | OG0 |
Hsap_ENST00000310955 | OG0 |
Acagra_model.g21766.t1 | OG0 |
Cscu_XM_023369515.1 | OG0 |
Tcas_TC001383_001 | OG0 |
Skow_XM_002741994.2 | OG0 |
Spur_XM_030996304.1 | OG0 |
Spur_XM_030996339.1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA152549 | OG0 |
Cgig_XM_011435206.3 | OG0 |
Acagra_model.g24293.t1 | OG0 |
Xesp_000675-T1 | OG0 |
Dgig_XM_028545338.1 | OG0 |
Dgig_XM_028549482.1 | OG0 |
Dgig_XM_028549483.1 | OG0 |
Exapal_XM_021060145.2 | OG0 |
Nvec_v1g192701 | OG0 |
Spis_XP_022802199_1 | OG0 |
Pocdam_XM_027181309.1 | OG0 |
Gasp_gasp1.m3.17854.m1 | OG0 |
Fspp_ffun1.m4.29509.m1 | OG0 |
Gfas_gfas1.m1.2061.m1 | OG0 |
Amil_Amillepora30902-RA | OG0 |
Amil_Amillepora30877-RA | OG0 |
Adig_XM_015900552.1 | OG0 |
0.416667