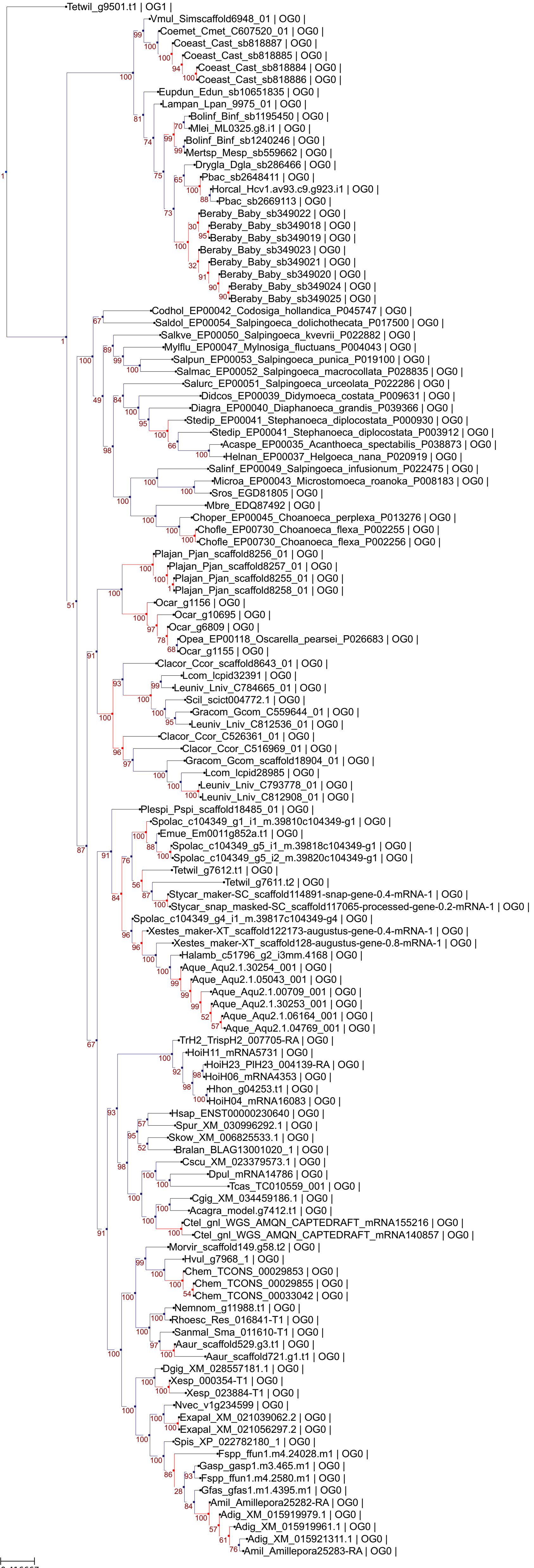
Tetwil_g9501.t1 | OG1 |
Vmul_Simscaffold6948_01 | OG0 |
Coemet_Cmet_C607520_01 | OG0 |
Coeast_Cast_sb818887 | OG0 |
Coeast_Cast_sb818885 | OG0 |
Coeast_Cast_sb818884 | OG0 |
Coeast_Cast_sb818886 | OG0 |
Eupdun_Edun_sb10651835 | OG0 |
Lampan_Lpan_9975_01 | OG0 |
Bolinf_Binf_sb1195450 | OG0 |
Mlei_ML0325.g8.i1 | OG0 |
Bolinf_Binf_sb1240246 | OG0 |
Mertsp_Mesp_sb559662 | OG0 |
Drygla_Dgla_sb286466 | OG0 |
Pbac_sb2648411 | OG0 |
Horcal_Hcv1.av93.c9.g923.i1 | OG0 |
Pbac_sb2669113 | OG0 |
Beraby_Baby_sb349022 | OG0 |
Beraby_Baby_sb349018 | OG0 |
Beraby_Baby_sb349019 | OG0 |
Beraby_Baby_sb349023 | OG0 |
Beraby_Baby_sb349021 | OG0 |
Beraby_Baby_sb349020 | OG0 |
Beraby_Baby_sb349024 | OG0 |
Beraby_Baby_sb349025 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P045747 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P017500 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P022882 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P004043 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P019100 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P028835 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P022286 | OG0 |
Didcos_EP00039_Didymoeca_costata_P009631 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P039366 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P000930 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P003912 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P038873 | OG0 |
Helnan_EP00037_Helgoeca_nana_P020919 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P022475 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P008183 | OG0 |
Sros_EGD81805 | OG0 |
Mbre_EDQ87492 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P013276 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P002255 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P002256 | OG0 |
Plajan_Pjan_scaffold8256_01 | OG0 |
Plajan_Pjan_scaffold8257_01 | OG0 |
Plajan_Pjan_scaffold8255_01 | OG0 |
Plajan_Pjan_scaffold8258_01 | OG0 |
Ocar_g1156 | OG0 |
Ocar_g10695 | OG0 |
Ocar_g6809 | OG0 |
Opea_EP00118_Oscarella_pearsei_P026683 | OG0 |
Ocar_g1155 | OG0 |
Clacor_Ccor_scaffold8643_01 | OG0 |
Lcom_lcpid32391 | OG0 |
Leuniv_Lniv_C784665_01 | OG0 |
Scil_scict004772.1 | OG0 |
Gracom_Gcom_C559644_01 | OG0 |
Leuniv_Lniv_C812536_01 | OG0 |
Clacor_Ccor_C526361_01 | OG0 |
Clacor_Ccor_C516969_01 | OG0 |
Gracom_Gcom_scaffold18904_01 | OG0 |
Lcom_lcpid28985 | OG0 |
Leuniv_Lniv_C793778_01 | OG0 |
Leuniv_Lniv_C812908_01 | OG0 |
Plespi_Pspi_scaffold18485_01 | OG0 |
Spolac_c104349_g1_i1_m.39810c104349-g1 | OG0 |
Emue_Em0011g852a.t1 | OG0 |
Spolac_c104349_g5_i1_m.39818c104349-g1 | OG0 |
Spolac_c104349_g5_i2_m.39820c104349-g1 | OG0 |
Tetwil_g7612.t1 | OG0 |
Tetwil_g7611.t2 | OG0 |
Stycar_maker-SC_scaffold114891-snap-gene-0.4-mRNA-1 | OG0 |
Stycar_snap_masked-SC_scaffold117065-processed-gene-0.2-mRNA-1 | OG0 |
Spolac_c104349_g4_i1_m.39817c104349-g4 | OG0 |
Xestes_maker-XT_scaffold122173-augustus-gene-0.4-mRNA-1 | OG0 |
Xestes_maker-XT_scaffold128-augustus-gene-0.8-mRNA-1 | OG0 |
Halamb_c51796_g2_i3mm.4168 | OG0 |
Aque_Aqu2.1.30254_001 | OG0 |
Aque_Aqu2.1.05043_001 | OG0 |
Aque_Aqu2.1.00709_001 | OG0 |
Aque_Aqu2.1.30253_001 | OG0 |
Aque_Aqu2.1.06164_001 | OG0 |
Aque_Aqu2.1.04769_001 | OG0 |
TrH2_TrispH2_007705-RA | OG0 |
HoiH11_mRNA5731 | OG0 |
HoiH23_PlH23_004139-RA | OG0 |
HoiH06_mRNA4353 | OG0 |
Hhon_g04253.t1 | OG0 |
HoiH04_mRNA16083 | OG0 |
Hsap_ENST00000230640 | OG0 |
Spur_XM_030996292.1 | OG0 |
Skow_XM_006825533.1 | OG0 |
Bralan_BLAG13001020_1 | OG0 |
Cscu_XM_023379573.1 | OG0 |
Dpul_mRNA14786 | OG0 |
Tcas_TC010559_001 | OG0 |
Cgig_XM_034459186.1 | OG0 |
Acagra_model.g7412.t1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA155216 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA140857 | OG0 |
Morvir_scaffold149.g58.t2 | OG0 |
Hvul_g7968_1 | OG0 |
Chem_TCONS_00029853 | OG0 |
Chem_TCONS_00029855 | OG0 |
Chem_TCONS_00033042 | OG0 |
Nemnom_g11988.t1 | OG0 |
Rhoesc_Res_016841-T1 | OG0 |
Sanmal_Sma_011610-T1 | OG0 |
Aaur_scaffold529.g3.t1 | OG0 |
Aaur_scaffold721.g1.t1 | OG0 |
Dgig_XM_028557181.1 | OG0 |
Xesp_000354-T1 | OG0 |
Xesp_023884-T1 | OG0 |
Nvec_v1g234599 | OG0 |
Exapal_XM_021039062.2 | OG0 |
Exapal_XM_021056297.2 | OG0 |
Spis_XP_022782180_1 | OG0 |
Fspp_ffun1.m4.24028.m1 | OG0 |
Gasp_gasp1.m3.465.m1 | OG0 |
Fspp_ffun1.m4.2580.m1 | OG0 |
Gfas_gfas1.m1.4395.m1 | OG0 |
Amil_Amillepora25282-RA | OG0 |
Adig_XM_015919979.1 | OG0 |
Adig_XM_015919961.1 | OG0 |
Adig_XM_015921311.1 | OG0 |
Amil_Amillepora25283-RA | OG0 |
0.416667