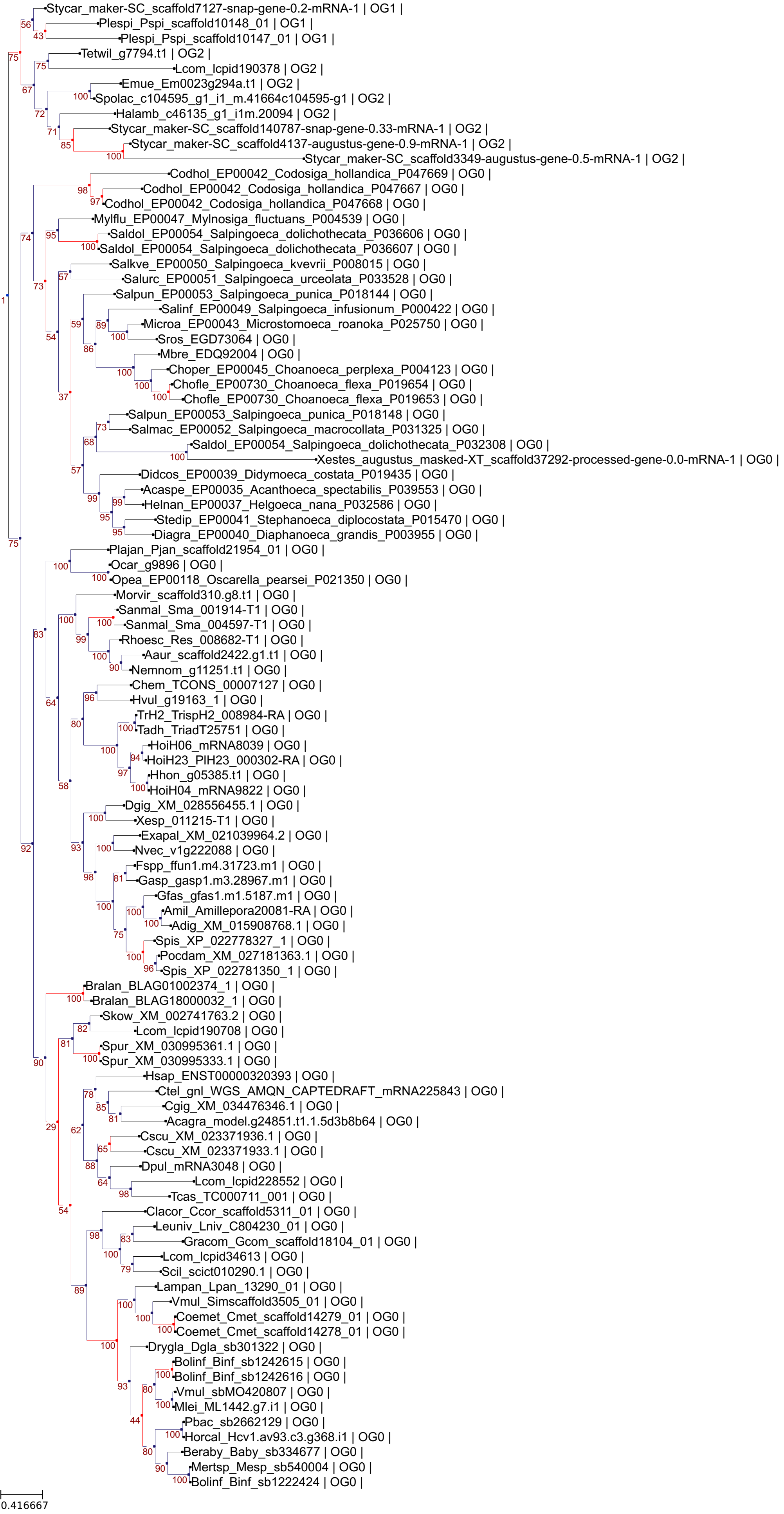
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Plespi_Pspi_scaffold10148_01 | OG1 |
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Tetwil_g7794.t1 | OG2 |
Lcom_lcpid190378 | OG2 |
Emue_Em0023g294a.t1 | OG2 |
Spolac_c104595_g1_i1_m.41664c104595-g1 | OG2 |
Halamb_c46135_g1_i1m.20094 | OG2 |
Stycar_maker-SC_scaffold140787-snap-gene-0.33-mRNA-1 | OG2 |
Stycar_maker-SC_scaffold4137-augustus-gene-0.9-mRNA-1 | OG2 |
Stycar_maker-SC_scaffold3349-augustus-gene-0.5-mRNA-1 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P047669 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P047667 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P047668 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P004539 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P036606 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P036607 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P008015 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P033528 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P018144 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P000422 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P025750 | OG0 |
Sros_EGD73064 | OG0 |
Mbre_EDQ92004 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P004123 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P019654 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P019653 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P018148 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P031325 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P032308 | OG0 |
Xestes_augustus_masked-XT_scaffold37292-processed-gene-0.0-mRNA-1 | OG0 |
Didcos_EP00039_Didymoeca_costata_P019435 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P039553 | OG0 |
Helnan_EP00037_Helgoeca_nana_P032586 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P015470 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P003955 | OG0 |
Plajan_Pjan_scaffold21954_01 | OG0 |
Ocar_g9896 | OG0 |
Opea_EP00118_Oscarella_pearsei_P021350 | OG0 |
Morvir_scaffold310.g8.t1 | OG0 |
Sanmal_Sma_001914-T1 | OG0 |
Sanmal_Sma_004597-T1 | OG0 |
Rhoesc_Res_008682-T1 | OG0 |
Aaur_scaffold2422.g1.t1 | OG0 |
Nemnom_g11251.t1 | OG0 |
Chem_TCONS_00007127 | OG0 |
Hvul_g19163_1 | OG0 |
TrH2_TrispH2_008984-RA | OG0 |
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HoiH06_mRNA8039 | OG0 |
HoiH23_PIH23_000302-RA | OG0 |
Hhon_g05385.t1 | OG0 |
HoiH04_mRNA9822 | OG0 |
Dgig_XM_028556455.1 | OG0 |
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Nvec_v1g222088 | OG0 |
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Amil_Amillepora20081-RA | OG0 |
Adig_XM_015908768.1 | OG0 |
Spis_XP_022778327_1 | OG0 |
Pocdam_XM_027181363.1 | OG0 |
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Tcas_TC000711_001 | OG0 |
Clacor_Ccor_scaffold5311_01 | OG0 |
Leuniv_Lniv_C804230_01 | OG0 |
Gracom_Gcom_scaffold18104_01 | OG0 |
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Scil_scict010290.1 | OG0 |
Lampan_Lpan_13290_01 | OG0 |
Vmul_Simscaffold3505_01 | OG0 |
Coemet_Cmet_scaffold14279_01 | OG0 |
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Drygla_Dgla_sb301322 | OG0 |
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Bolinf_Binf_sb1242616 | OG0 |
Vmul_sbMO420807 | OG0 |
Mlei_ML1442.g7.i1 | OG0 |
Pbac_sb2662129 | OG0 |
Horcal_Hcv1.av93.c3.g368.i1 | OG0 |
Beraby_Baby_sb334677 | OG0 |
Mertsp_Mesp_sb540004 | OG0 |
Bolinf_Binf_sb1222424 | OG0 |
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