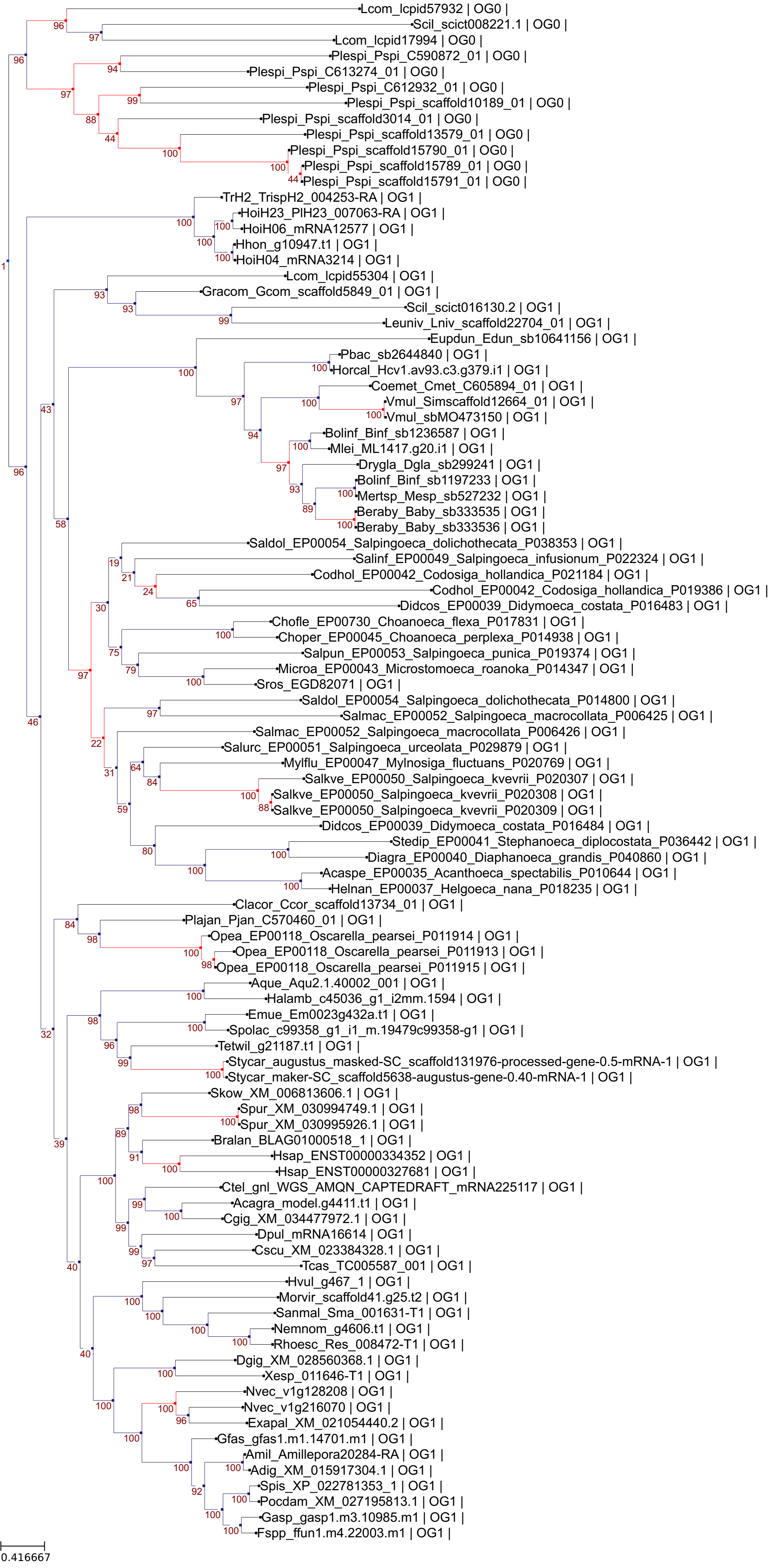
Lcom_lcpid57932 | OG0 |
Scil_scict008221.1 | OG0 |
Lcom_lcpid17994 | OG0 |
Plespi_Pspi_C590872_01 | OG0 |
Plespi_Pspi_C613274_01 | OG0 |
Plespi_Pspi_C612932_01 | OG0 |
Plespi_Pspi_scaffold10189_01 | OG0 |
Plespi_Pspi_scaffold3014_01 | OG0 |
Plespi_Pspi_scaffold13579_01 | OG0 |
Plespi_Pspi_scaffold15790_01 | OG0 |
Plespi_Pspi_scaffold15789_01 | OG0 |
Plespi_Pspi_scaffold15791_01 | OG0 |
TrH2_TrispH2_004253-RA | OG1 |
HoiH23_PlH23_007063-RA | OG1 |
HoiH06_mRNA12577 | OG1 |
Hhon_g10947.t1 | OG1 |
HoiH04_mRNA3214 | OG1 |
Lcom_lcpid55304 | OG1 |
Gracom_Gcom_scaffold5849_01 | OG1 |
Scil_scict016130.2 | OG1 |
Leuniv_Lniv_scaffold22704_01 | OG1 |
Eupdun_Edun_sb10641156 | OG1 |
Pbac_sb2644840 | OG1 |
Horcal_Hcv1.av93.c3.g379.i1 | OG1 |
Coemet_Cmet_C605894_01 | OG1 |
Vmul_Simscaffold12664_01 | OG1 |
Vmul_sbMO473150 | OG1 |
Bolinf_Binf_sb1236587 | OG1 |
Mlei_ML1417.g20.i1 | OG1 |
Drygla_Dgla_sb299241 | OG1 |
Bolinf_Binf_sb1197233 | OG1 |
Mertsp_Mesp_sb527232 | OG1 |
Beraby_Baby_sb333535 | OG1 |
Beraby_Baby_sb333536 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P038353 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P022324 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P021184 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P019386 | OG1 |
Didcos_EP00039_Didymoeca_costata_P016483 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P017831 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P014938 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P019374 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P014347 | OG1 |
Sros_EGD82071 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P014800 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P006425 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P006426 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P029879 | OG1 |
Mylflu_EP00047_Mylnosiga_fluctuans_P020769 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P020307 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P020308 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P020309 | OG1 |
Didcos_EP00039_Didymoeca_costata_P016484 | OG1 |
Stedip_EP00041_Stephanoeca_diplocostata_P036442 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P040860 | OG1 |
Acaspe_EP00035_Acanthoeca_spectabilis_P010644 | OG1 |
Helnan_EP00037_Helgoeca_nana_P018235 | OG1 |
Clacor_Ccor_scaffold13734_01 | OG1 |
Plajan_Pjan_C570460_01 | OG1 |
Opea_EP00118_Oscarella_pearsei_P011914 | OG1 |
Opea_EP00118_Oscarella_pearsei_P011913 | OG1 |
Opea_EP00118_Oscarella_pearsei_P011915 | OG1 |
Aque_Aqu2.1.40002_001 | OG1 |
Halamb_c45036_g1_i2mm.1594 | OG1 |
Emue_Em0023g432a.t1 | OG1 |
Spolac_c99358_g1_i1_m.19479c99358-g1 | OG1 |
Tetwil_g21187.t1 | OG1 |
Stycar_augustus_masked-SC_scaffold131976-processed-gene-0.5-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold5638-augustus-gene-0.40-mRNA-1 | OG1 |
Skow_XM_006813606.1 | OG1 |
Spur_XM_030994749.1 | OG1 |
Spur_XM_030995926.1 | OG1 |
Bralan_BLAG01000518_1 | OG1 |
Hsap_ENST00000334352 | OG1 |
Hsap_ENST00000327681 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA225117 | OG1 |
Acagra_model.g4411.t1 | OG1 |
Cgig_XM_034477972.1 | OG1 |
Dpul_mRNA16614 | OG1 |
Cscu_XM_023384328.1 | OG1 |
Tcas_TC005587_001 | OG1 |
Hvul_g467_1 | OG1 |
Morvir_scaffold41.g25.t2 | OG1 |
Sanmal_Sma_001631-T1 | OG1 |
Nemnom_g4606.t1 | OG1 |
Rhoesc_Res_008472-T1 | OG1 |
Dgig_XM_028560368.1 | OG1 |
Xesp_011646-T1 | OG1 |
Nvec_v1g128208 | OG1 |
Nvec_v1g216070 | OG1 |
Exapal_XM_021054440.2 | OG1 |
Gfas_gfas1.m1.14701.m1 | OG1 |
Amil_Amillepora20284-RA | OG1 |
Adig_XM_015917304.1 | OG1 |
Spis_XP_022781353_1 | OG1 |
Pocdam_XM_027195813.1 | OG1 |
Gasp_gasp1.m3.10985.m1 | OG1 |
Fspp_ffun1.m4.22003.m1 | OG1 |
0.416667