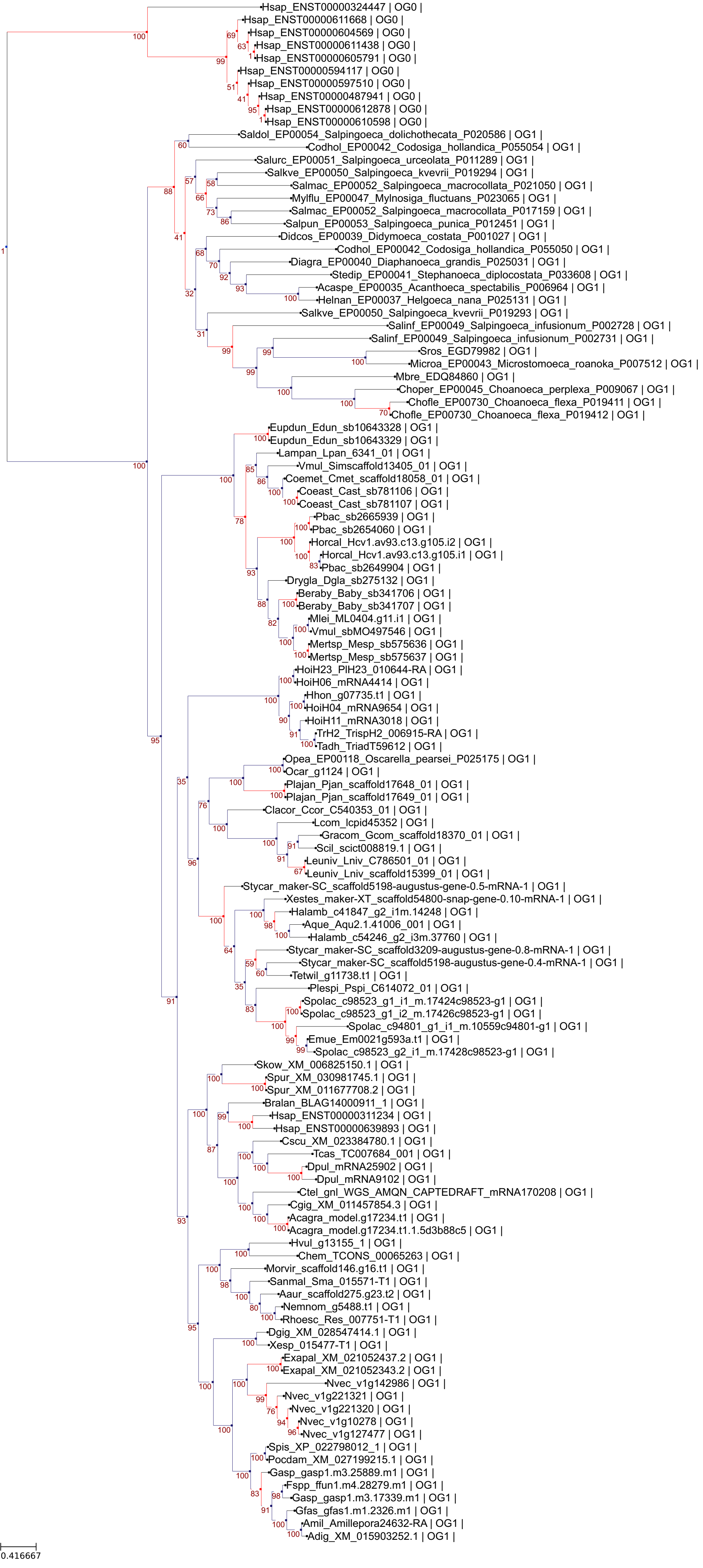
Hsap_ENST00000324447 | OG0 |
Hsap_ENST00000611668 | OG0 |
Hsap_ENST00000604569 | OG0 |
Hsap_ENST00000611438 | OG0 |
Hsap_ENST00000605791 | OG0 |
Hsap_ENST00000594117 | OG0 |
Hsap_ENST00000597510 | OG0 |
Hsap_ENST00000487941 | OG0 |
Hsap_ENST00000612878 | OG0 |
Hsap_ENST00000610598 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P020586 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P055054 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P011289 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P019294 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P021050 | OG1 |
Mylflu_EP00047_Mylnosiga_fluctuans_P023065 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P017159 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P012451 | OG1 |
Didcos_EP00039_Didymoeca_costata_P001027 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P055050 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P025031 | OG1 |
Stedip_EP00041_Stephanoeca_diplocostata_P033608 | OG1 |
Acaspe_EP00035_Acanthoeca_spectabilis_P006964 | OG1 |
Helnan_EP00037_Helgoeca_nana_P025131 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P019293 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P002728 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P002731 | OG1 |
Sros_EGD79982 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P007512 | OG1 |
Mbre_EDQ84860 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P009067 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P019411 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P019412 | OG1 |
Eupdun_Edun_sb10643328 | OG1 |
Eupdun_Edun_sb10643329 | OG1 |
Lampan_Lpan_6341_01 | OG1 |
Vmul_Simscaffold13405_01 | OG1 |
Coemet_Cmet_scaffold18058_01 | OG1 |
Coeast_Cast_sb781106 | OG1 |
Coeast_Cast_sb781107 | OG1 |
Pbac_sb2665939 | OG1 |
Pbac_sb2654060 | OG1 |
Horcal_Hcv1.av93.c13.g105.i2 | OG1 |
Horcal_Hcv1.av93.c13.g105.i1 | OG1 |
Pbac_sb2649904 | OG1 |
Drygla_Dgla_sb275132 | OG1 |
Beraby_Baby_sb341706 | OG1 |
Beraby_Baby_sb341707 | OG1 |
Mlei_ML0404.g11.i1 | OG1 |
Vmul_sbMO497546 | OG1 |
Mertsp_Mesp_sb575636 | OG1 |
Mertsp_Mesp_sb575637 | OG1 |
HoiH23_PlH23_010644-RA | OG1 |
HoiH06_mRNA4414 | OG1 |
Hhon_g07735.t1 | OG1 |
HoiH04_mRNA9654 | OG1 |
HoiH11_mRNA3018 | OG1 |
TrH2_TrispH2_006915-RA | OG1 |
Tadh_TriadT59612 | OG1 |
Opea_EP00118_Oscarella_pearsei_P025175 | OG1 |
Ocar_g1124 | OG1 |
Plajan_Pjan_scaffold17648_01 | OG1 |
Plajan_Pjan_scaffold17649_01 | OG1 |
Clacor_Ccor_C540353_01 | OG1 |
Lcom_lcpid45352 | OG1 |
Gracom_Gcom_scaffold18370_01 | OG1 |
Scil_scict008819.1 | OG1 |
Leuniv_Lniv_C786501_01 | OG1 |
Leuniv_Lniv_scaffold15399_01 | OG1 |
Stycar_maker-SC_scaffold5198-augustus-gene-0.5-mRNA-1 | OG1 |
Xestes_maker-XT_scaffold54800-snap-gene-0.10-mRNA-1 | OG1 |
Halamb_c41847_g2_i1m.14248 | OG1 |
Aque_Aqu2.1.41006_001 | OG1 |
Halamb_c54246_g2_i3m.37760 | OG1 |
Stycar_maker-SC_scaffold3209-augustus-gene-0.8-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold5198-augustus-gene-0.4-mRNA-1 | OG1 |
Tetwil_g11738.t1 | OG1 |
Plespi_Pspi_C614072_01 | OG1 |
Spolac_c98523_g1_i1_m.17424c98523-g1 | OG1 |
Spolac_c98523_g1_i2_m.17426c98523-g1 | OG1 |
Spolac_c94801_g1_i1_m.10559c94801-g1 | OG1 |
Emue_Em0021g593a.t1 | OG1 |
Spolac_c98523_g2_i1_m.17428c98523-g1 | OG1 |
Skow_XM_006825150.1 | OG1 |
Spur_XM_030981745.1 | OG1 |
Spur_XM_011677708.2 | OG1 |
Bralan_BLAG14000911_1 | OG1 |
Hsap_ENST00000311234 | OG1 |
Hsap_ENST00000639893 | OG1 |
Cscu_XM_023384780.1 | OG1 |
Tcas_TC007684_001 | OG1 |
Dpul_mRNA25902 | OG1 |
Dpul_mRNA9102 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA170208 | OG1 |
Cgig_XM_011457854.3 | OG1 |
Acagra_model.g17234.t1 | OG1 |
Acagra_model.g17234.t1.1.5d3b88c5 | OG1 |
Hvul_g13155_1 | OG1 |
Chem_TCONS_00065263 | OG1 |
Morvir_scaffold146.g16.t1 | OG1 |
Sanmal_Sma_015571-T1 | OG1 |
Aaur_scaffold275.g23.t2 | OG1 |
Nemnom_g5488.t1 | OG1 |
Rhoesc_Res_007751-T1 | OG1 |
Dgig_XM_028547414.1 | OG1 |
Xesp_015477-T1 | OG1 |
Exapal_XM_021052437.2 | OG1 |
Exapal_XM_021052343.2 | OG1 |
Nvec_v1g142986 | OG1 |
Nvec_v1g221321 | OG1 |
Nvec_v1g221320 | OG1 |
Nvec_v1g10278 | OG1 |
Nvec_v1g127477 | OG1 |
Spis_XP_022798012_1 | OG1 |
Pocdam_XM_027199215.1 | OG1 |
Gasp_gasp1.m3.25889.m1 | OG1 |
Fspp_ffun1.m4.28279.m1 | OG1 |
Gasp_gasp1.m3.17339.m1 | OG1 |
Gfas_gfas1.m1.2326.m1 | OG1 |
Amil_Amillepora24632-RA | OG1 |
Adig_XM_015903252.1 | OG1 |
0.416667