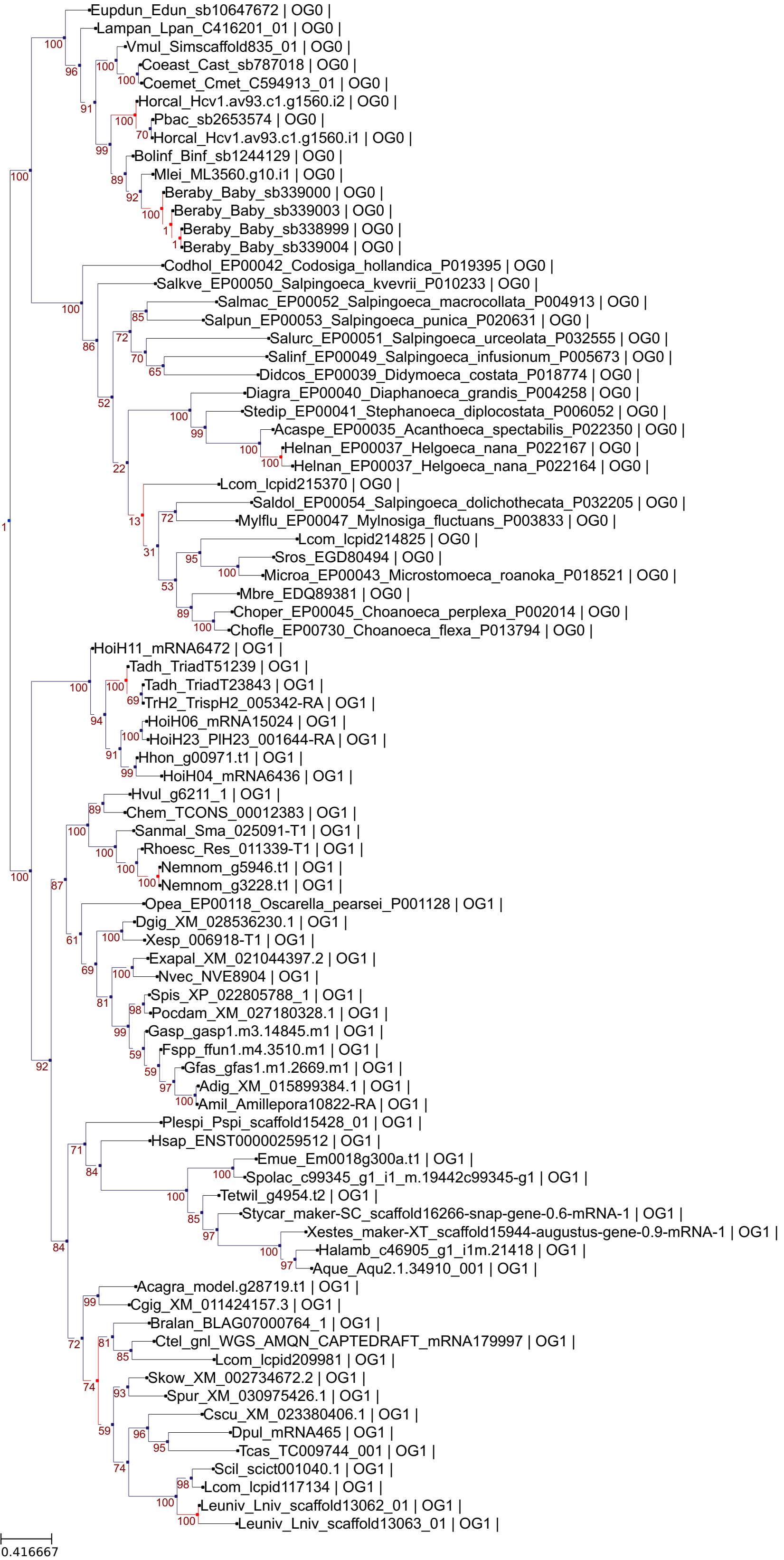
Eupdun_Edun_sb10647672 | OG0 |
Lampan_Lpan_C416201_01 | OG0 |
Vmul_Simscaffold835_01 | OG0 |
Coeast_Cast_sb787018 | OG0 |
Coemet_Cmet_C594913_01 | OG0 |
Horcal_Hcv1.av93.c1.g1560.i2 | OG0 |
Pbac_sb2653574 | OG0 |
Horcal_Hcv1.av93.c1.g1560.i1 | OG0 |
Bolinf_Binf_sb1244129 | OG0 |
Mlei_ML3560.g10.i1 | OG0 |
Beraby_Baby_sb339000 | OG0 |
Beraby_Baby_sb339003 | OG0 |
Beraby_Baby_sb338999 | OG0 |
Beraby_Baby_sb339004 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P019395 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P010233 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P004913 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P020631 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P032555 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P005673 | OG0 |
Didcos_EP00039_Didymoeca_costata_P018774 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P004258 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P006052 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P022350 | OG0 |
Helnan_EP00037_Helgoeca_nana_P022167 | OG0 |
Helnan_EP00037_Helgoeca_nana_P022164 | OG0 |
Lcom_lcpid215370 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P032205 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P003833 | OG0 |
Lcom_lcpid214825 | OG0 |
Sros_EGD80494 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P018521 | OG0 |
Mbre_EDQ89381 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P002014 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P013794 | OG0 |
HoiH11_mRNA6472 | OG1 |
Tadh_TriadT51239 | OG1 |
Tadh_TriadT23843 | OG1 |
TrH2_TrispH2_005342-RA | OG1 |
HoiH06_mRNA15024 | OG1 |
HoiH23_PlH23_001644-RA | OG1 |
Hhon_g00971.t1 | OG1 |
HoiH04_mRNA6436 | OG1 |
Hvul_g6211_1 | OG1 |
Chem_TCONS_00012383 | OG1 |
Sanmal_Sma_025091-T1 | OG1 |
Rhoesc_Res_011339-T1 | OG1 |
Nemnom_g5946.t1 | OG1 |
Nemnom_g3228.t1 | OG1 |
Opea_EP00118_Oscarella_pearsei_P001128 | OG1 |
Dgig_XM_028536230.1 | OG1 |
Xesp_006918-T1 | OG1 |
Exapal_XM_021044397.2 | OG1 |
Nvec_NVE8904 | OG1 |
Spis_XP_022805788_1 | OG1 |
Pocdam_XM_027180328.1 | OG1 |
Gasp_gasp1.m3.14845.m1 | OG1 |
Fspp_ffun1.m4.3510.m1 | OG1 |
Gfas_gfas1.m1.2669.m1 | OG1 |
Adig_XM_015899384.1 | OG1 |
Amil_Amillepora10822-RA | OG1 |
Plespi_Pspi_scaffold15428_01 | OG1 |
Hsap_ENST00000259512 | OG1 |
Emue_Em0018g300a.t1 | OG1 |
Spolac_c99345_g1_i1_m.19442c99345-g1 | OG1 |
Tetwil_g4954.t2 | OG1 |
Stycar_maker-SC_scaffold16266-snap-gene-0.6-mRNA-1 | OG1 |
Xestes_maker-XT_scaffold15944-augustus-gene-0.9-mRNA-1 | OG1 |
Halamb_c46905_g1_i1m.21418 | OG1 |
Aque_Aqu2.1.34910_001 | OG1 |
Acagra_model.g28719.t1 | OG1 |
Cgig_XM_011424157.3 | OG1 |
Bralan_BLAG07000764_1 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA179997 | OG1 |
Lcom_lcpid209981 | OG1 |
Skow_XM_002734672.2 | OG1 |
Spur_XM_030975426.1 | OG1 |
Cscu_XM_023380406.1 | OG1 |
Dpul_mRNA465 | OG1 |
Tcas_TC009744_001 | OG1 |
Scil_scict001040.1 | OG1 |
Lcom_lcpid117134 | OG1 |
Leuniv_Lniv_scaffold13062_01 | OG1 |
Leuniv_Lniv_scaffold13063_01 | OG1 |
0.416667