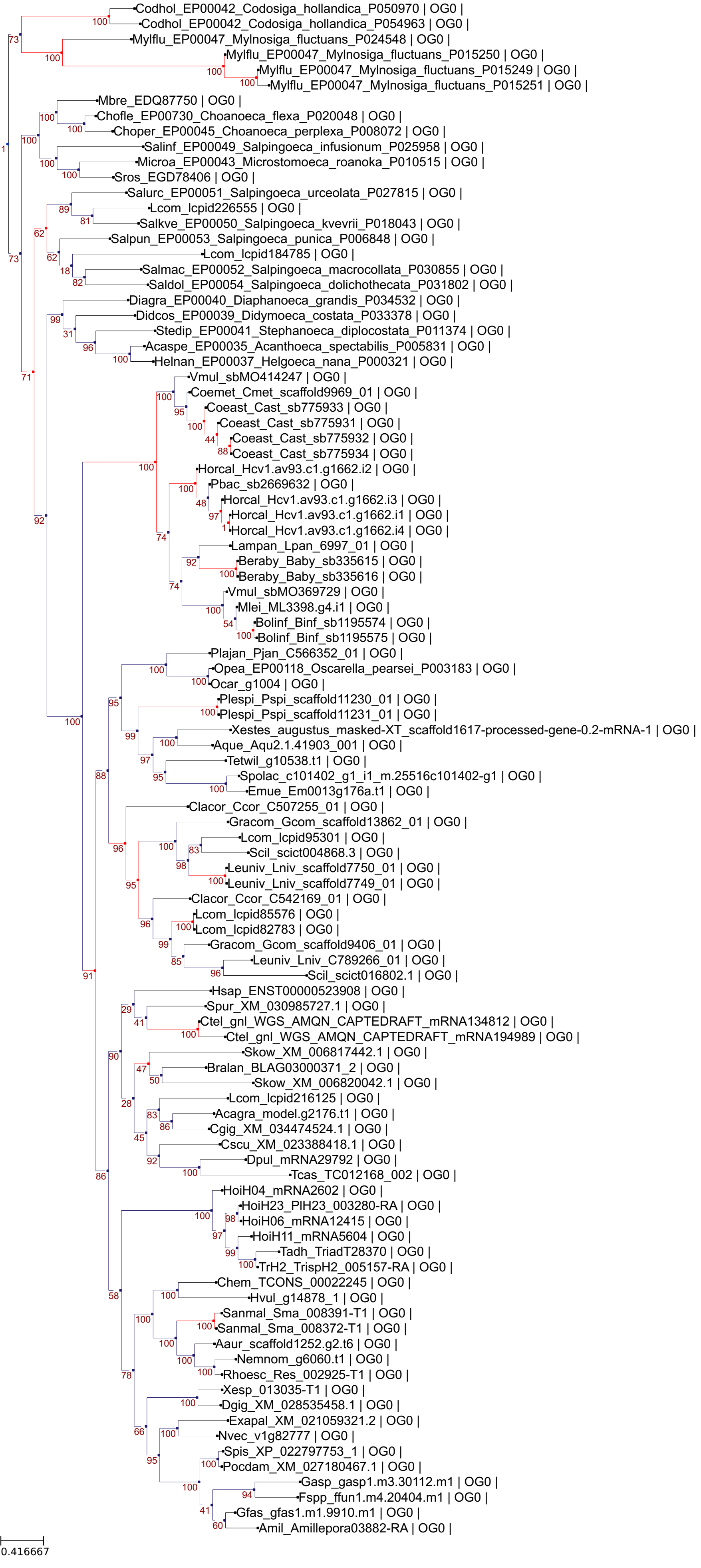
Codhol_EP00042_Codosiga_hollandica_P050970 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P054963 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P024548 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P015250 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P015249 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P015251 | OG0 |
Mbre_EDQ87750 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P020048 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P008072 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P025958 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P010515 | OG0 |
Sros_EGD78406 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P027815 | OG0 |
Lcom_lcpid226555 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P018043 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P006848 | OG0 |
Lcom_lcpid184785 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P030855 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P031802 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P034532 | OG0 |
Didcos_EP00039_Didymoeca_costata_P033378 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P011374 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P005831 | OG0 |
Helnan_EP00037_Helgoeca_nana_P000321 | OG0 |
Vmul_sbMO414247 | OG0 |
Coemet_Cmet_scaffold9969_01 | OG0 |
Coeast_Cast_sb775933 | OG0 |
Coeast_Cast_sb775931 | OG0 |
Coeast_Cast_sb775932 | OG0 |
Coeast_Cast_sb775934 | OG0 |
Horcal_Hcv1.av93.c1.g1662.i2 | OG0 |
Pbac_sb2669632 | OG0 |
Horcal_Hcv1.av93.c1.g1662.i3 | OG0 |
Horcal_Hcv1.av93.c1.g1662.i1 | OG0 |
Horcal_Hcv1.av93.c1.g1662.i4 | OG0 |
Lampan_Lpan_6997_01 | OG0 |
Beraby_Baby_sb335615 | OG0 |
Beraby_Baby_sb335616 | OG0 |
Vmul_sbMO369729 | OG0 |
Mlei_ML3398.g4.i1 | OG0 |
Bolinf_Binf_sb1195574 | OG0 |
Bolinf_Binf_sb1195575 | OG0 |
Plajan_Pjan_C566352_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P003183 | OG0 |
Ocar_g1004 | OG0 |
Plespi_Pspi_scaffold11230_01 | OG0 |
Plespi_Pspi_scaffold11231_01 | OG0 |
Xestes_augustus_masked-XT_scaffold1617-processed-gene-0.2-mRNA-1 | OG0 |
Aque_Aqu2.1.41903_001 | OG0 |
Tetwil_g10538.t1 | OG0 |
Spolac_c101402_g1_i1_m.25516c101402-g1 | OG0 |
Emue_Em0013g176a.t1 | OG0 |
Clacor_Ccor_C507255_01 | OG0 |
Gracom_Gcom_scaffold13862_01 | OG0 |
Lcom_lcpid95301 | OG0 |
Scil_scict004868.3 | OG0 |
Leuniv_Lniv_scaffold7750_01 | OG0 |
Leuniv_Lniv_scaffold7749_01 | OG0 |
Clacor_Ccor_C542169_01 | OG0 |
Lcom_lcpid85576 | OG0 |
Lcom_lcpid82783 | OG0 |
Gracom_Gcom_scaffold9406_01 | OG0 |
Leuniv_Lniv_C789266_01 | OG0 |
Scil_scict016802.1 | OG0 |
Hsap_ENST00000523908 | OG0 |
Spur_XM_030985727.1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA134812 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA194989 | OG0 |
Skow_XM_006817442.1 | OG0 |
Bralan_BLAG03000371_2 | OG0 |
Skow_XM_006820042.1 | OG0 |
Lcom_lcpid216125 | OG0 |
Acagra_model.g2176.t1 | OG0 |
Cgig_XM_034474524.1 | OG0 |
Cscu_XM_023388418.1 | OG0 |
Dpul_mRNA29792 | OG0 |
Tcas_TC012168_002 | OG0 |
HoiH04_mRNA2602 | OG0 |
HoiH23_PlH23_003280-RA | OG0 |
HoiH06_mRNA12415 | OG0 |
HoiH11_mRNA5604 | OG0 |
Tadh_TriadT28370 | OG0 |
TrH2_TrispH2_005157-RA | OG0 |
Chem_TCONS_00022245 | OG0 |
Hvul_g14878_1 | OG0 |
Sanmal_Sma_008391-T1 | OG0 |
Sanmal_Sma_008372-T1 | OG0 |
Aaur_scaffold1252.g2.t6 | OG0 |
Nemnom_g6060.t1 | OG0 |
Rhoesc_Res_002925-T1 | OG0 |
Xesp_013035-T1 | OG0 |
Dgig_XM_028535458.1 | OG0 |
Exapal_XM_021059321.2 | OG0 |
Nvec_v1g82777 | OG0 |
Spis_XP_022797753_1 | OG0 |
Pocdam_XM_027180467.1 | OG0 |
Gasp_gasp1.m3.30112.m1 | OG0 |
Fspp_ffun1.m4.20404.m1 | OG0 |
Gfas_gfas1.m1.9910.m1 | OG0 |
Amil_Amillepora03882-RA | OG0 |
0.416667