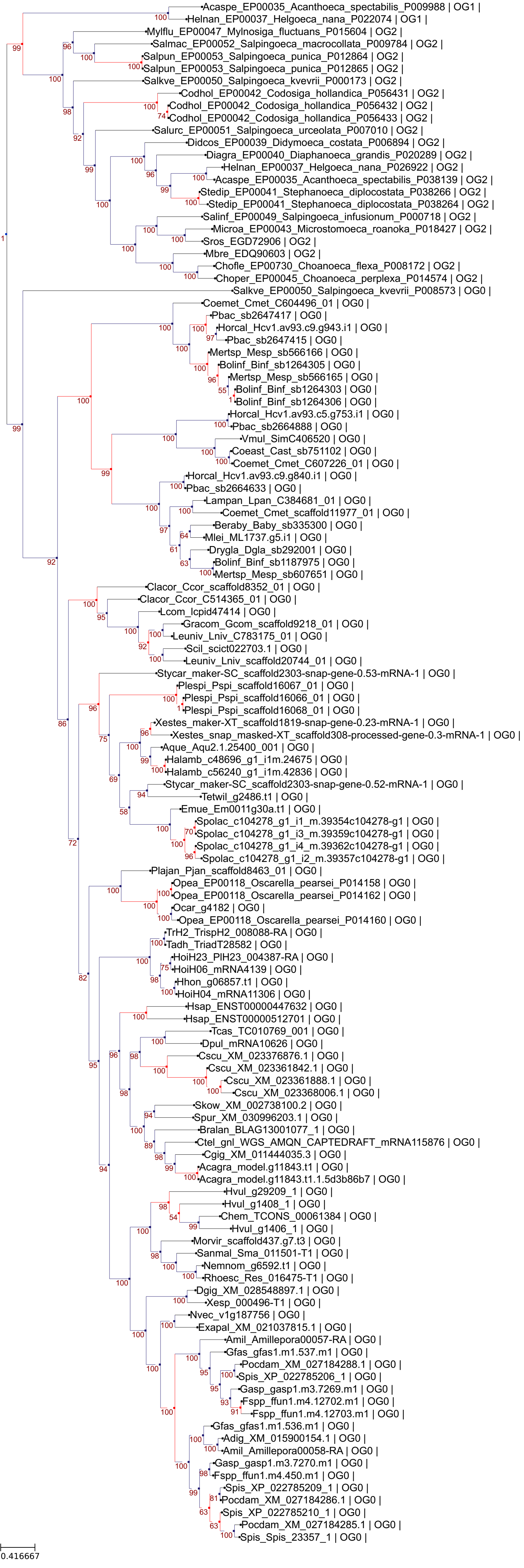
Acaspe_EP00035_Acanthoeca_spectabilis_P009988 | OG1 |
Helnan_EP00037_Helgoeca_nana_P022074 | OG1 |
Mylflu_EP00047_Mylnosiga_fluctuans_P015604 | OG2 |
Salmac_EP00052_Salpingoeca_macrocollata_P009784 | OG2 |
Salpun_EP00053_Salpingoeca_punica_P012864 | OG2 |
Salpun_EP00053_Salpingoeca_punica_P012865 | OG2 |
Salkve_EP00050_Salpingoeca_kvevrii_P000173 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P056431 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P056432 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P056433 | OG2 |
Salurc_EP00051_Salpingoeca_urceolata_P007010 | OG2 |
Didcos_EP00039_Didymoeca_costata_P006894 | OG2 |
Diagra_EP00040_Diaphanoeca_grandis_P020289 | OG2 |
Helnan_EP00037_Helgoeca_nana_P026922 | OG2 |
Acaspe_EP00035_Acanthoeca_spectabilis_P038139 | OG2 |
Stedip_EP00041_Stephanoeca_diplocostata_P038266 | OG2 |
Stedip_EP00041_Stephanoeca_diplocostata_P038264 | OG2 |
Salinf_EP00049_Salpingoeca_infusionum_P000718 | OG2 |
Microa_EP00043_Microstomoeca_roanoka_P018427 | OG2 |
Sros_EGD72906 | OG2 |
Mbre_EDQ90603 | OG2 |
Chofle_EP00730_Choanoeca_flexa_P008172 | OG2 |
Choper_EP00045_Choanoeca_perplexa_P014574 | OG2 |
Salkve_EP00050_Salpingoeca_kvevrii_P008573 | OG0 |
Coemet_Cmet_C604496_01 | OG0 |
Pbac_sb2647417 | OG0 |
Horcal_Hcv1.av93.c9.g943.i1 | OG0 |
Pbac_sb2647415 | OG0 |
Mertsp_Mesp_sb566166 | OG0 |
Bolinf_Binf_sb1264305 | OG0 |
Mertsp_Mesp_sb566165 | OG0 |
Bolinf_Binf_sb1264303 | OG0 |
Bolinf_Binf_sb1264306 | OG0 |
Horcal_Hcv1.av93.c5.g753.i1 | OG0 |
Pbac_sb2664888 | OG0 |
Vmul_SimC406520 | OG0 |
Coeast_Cast_sb751102 | OG0 |
Coemet_Cmet_C607226_01 | OG0 |
Horcal_Hcv1.av93.c9.g840.i1 | OG0 |
Pbac_sb2664633 | OG0 |
Lampan_Lpan_C384681_01 | OG0 |
Coemet_Cmet_scaffold11977_01 | OG0 |
Beraby_Baby_sb335300 | OG0 |
Mlei_ML1737.g5.i1 | OG0 |
Drygla_Dgla_sb292001 | OG0 |
Bolinf_Binf_sb1187975 | OG0 |
Mertsp_Mesp_sb607651 | OG0 |
Clacor_Ccor_scaffold8352_01 | OG0 |
Clacor_Ccor_C514365_01 | OG0 |
Lcom_lcpid47414 | OG0 |
Gracom_Gcom_scaffold9218_01 | OG0 |
Leuniv_Lniv_C783175_01 | OG0 |
Scil_scict022703.1 | OG0 |
Leuniv_Lniv_scaffold20744_01 | OG0 |
Stycar_maker-SC_scaffold2303-snap-gene-0.53-mRNA-1 | OG0 |
Plespi_Pspi_scaffold16067_01 | OG0 |
Plespi_Pspi_scaffold16066_01 | OG0 |
Plespi_Pspi_scaffold16068_01 | OG0 |
Xestes_maker-XT_scaffold1819-snap-gene-0.23-mRNA-1 | OG0 |
Xestes_snap_masked-XT_scaffold308-processed-gene-0.3-mRNA-1 | OG0 |
Aque_Aqu2.1.25400_001 | OG0 |
Halamb_c48696_g1_i1m.24675 | OG0 |
Halamb_c56240_g1_i1m.42836 | OG0 |
Stycar_maker-SC_scaffold2303-snap-gene-0.52-mRNA-1 | OG0 |
Tetwil_g2486.t1 | OG0 |
Emue_Em0011g30a.t1 | OG0 |
Spolac_c104278_g1_i1_m.39354c104278-g1 | OG0 |
Spolac_c104278_g1_i3_m.39359c104278-g1 | OG0 |
Spolac_c104278_g1_i4_m.39362c104278-g1 | OG0 |
Spolac_c104278_g1_i2_m.39357c104278-g1 | OG0 |
Plajan_Pjan_scaffold8463_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P014158 | OG0 |
Opea_EP00118_Oscarella_pearsei_P014162 | OG0 |
Ocar_g4182 | OG0 |
Opea_EP00118_Oscarella_pearsei_P014160 | OG0 |
TrH2_TrispH2_008088-RA | OG0 |
Tadh_TriadT28582 | OG0 |
HoiH23_PlH23_004387-RA | OG0 |
HoiH06_mRNA4139 | OG0 |
Hhon_g06857.t1 | OG0 |
HoiH04_mRNA11306 | OG0 |
Hsap_ENST00000447632 | OG0 |
Hsap_ENST00000512701 | OG0 |
Tcas_TC010769_001 | OG0 |
Dpul_mRNA10626 | OG0 |
Cscu_XM_023376876.1 | OG0 |
Cscu_XM_023361842.1 | OG0 |
Cscu_XM_023361888.1 | OG0 |
Cscu_XM_023368006.1 | OG0 |
Skow_XM_002738100.2 | OG0 |
Spur_XM_030996203.1 | OG0 |
Bralan_BLAG13001077_1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA115876 | OG0 |
Cgig_XM_011444035.3 | OG0 |
Acagra_model.g11843.t1 | OG0 |
Acagra_model.g11843.t1.1.5d3b86b7 | OG0 |
Hvul_g29209_1 | OG0 |
Hvul_g1408_1 | OG0 |
Chem_TCONS_00061384 | OG0 |
Hvul_g1406_1 | OG0 |
Morvir_scaffold437.g7.t3 | OG0 |
Sanmal_Sma_011501-T1 | OG0 |
Nemnom_g6592.t1 | OG0 |
Rhoesc_Res_016475-T1 | OG0 |
Dgig_XM_028548897.1 | OG0 |
Xesp_000496-T1 | OG0 |
Nvec_v1g187756 | OG0 |
Exapal_XM_021037815.1 | OG0 |
Amil_Amillepora00057-RA | OG0 |
Gfas_gfas1.m1.537.m1 | OG0 |
Pocdam_XM_027184288.1 | OG0 |
Spis_XP_022785206_1 | OG0 |
Gasp_gasp1.m3.7269.m1 | OG0 |
Fspp_ffun1.m4.12702.m1 | OG0 |
Fspp_ffun1.m4.12703.m1 | OG0 |
Gfas_gfas1.m1.536.m1 | OG0 |
Adig_XM_015900154.1 | OG0 |
Amil_Amillepora00058-RA | OG0 |
Gasp_gasp1.m3.7270.m1 | OG0 |
Fspp_ffun1.m4.450.m1 | OG0 |
Spis_XP_022785209_1 | OG0 |
Pocdam_XM_027184286.1 | OG0 |
Spis_XP_022785210_1 | OG0 |
Pocdam_XM_027184285.1 | OG0 |
Spis_Spis_23357_1 | OG0 |
0.416667