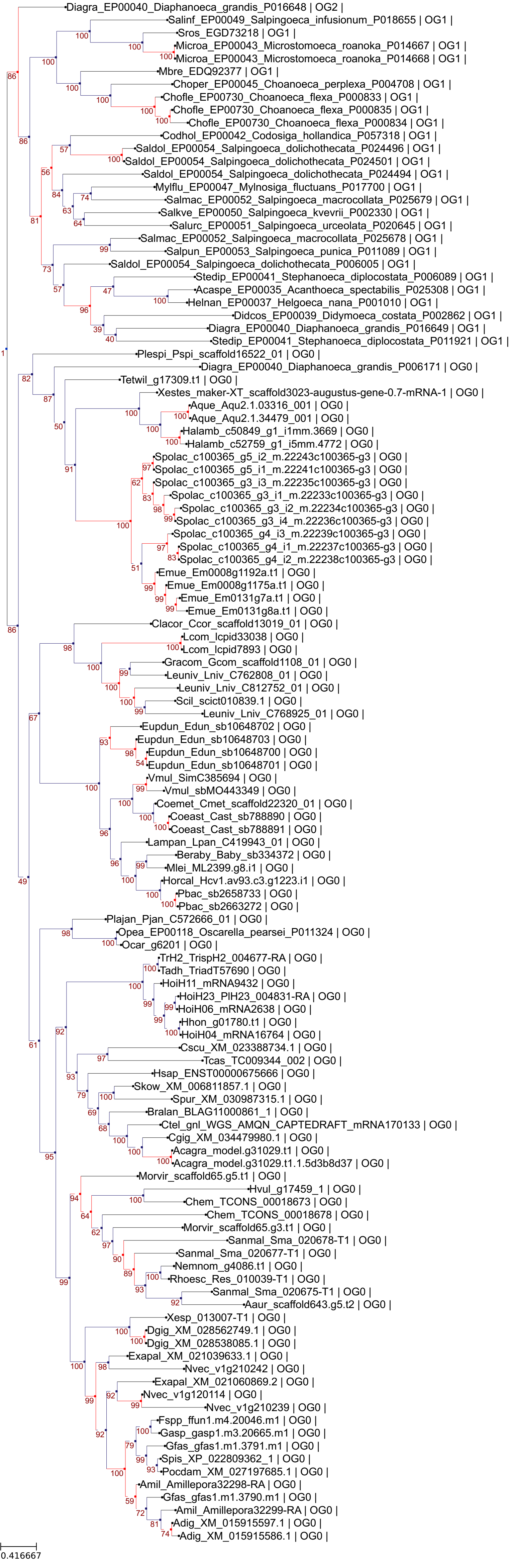
Diagra_EP00040_Diaphanoeca_grandis_P016648 | OG2 |
Salinf_EP00049_Salpingoeca_infusionum_P018655 | OG1 |
Sros_EGD73218 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P014667 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P014668 | OG1 |
Mbre_EDQ92377 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P004708 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P000833 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P000835 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P000834 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P057318 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P024496 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P024501 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P024494 | OG1 |
Mylflu_EP00047_Mylnosiga_fluctuans_P017700 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P025679 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P002330 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P020645 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P025678 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P011089 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P006005 | OG1 |
Stedip_EP00041_Stephanoeca_diplocostata_P006089 | OG1 |
Acaspe_EP00035_Acanthoeca_spectabilis_P025308 | OG1 |
Helnan_EP00037_Helgoeca_nana_P001010 | OG1 |
Didcos_EP00039_Didymoeca_costata_P002862 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P016649 | OG1 |
Stedip_EP00041_Stephanoeca_diplocostata_P011921 | OG1 |
Plespi_Pspi_scaffold16522_01 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P006171 | OG0 |
Tetwil_g17309.t1 | OG0 |
Xestes_maker-XT_scaffold3023-augustus-gene-0.7-mRNA-1 | OG0 |
Aque_Aqu2.1.03316_001 | OG0 |
Aque_Aqu2.1.34479_001 | OG0 |
Halamb_c50849_g1_i1mm.3669 | OG0 |
Halamb_c52759_g1_i5mm.4772 | OG0 |
Spolac_c100365_g5_i2_m.22243c100365-g3 | OG0 |
Spolac_c100365_g5_i1_m.22241c100365-g3 | OG0 |
Spolac_c100365_g3_i3_m.22235c100365-g3 | OG0 |
Spolac_c100365_g3_i1_m.22233c100365-g3 | OG0 |
Spolac_c100365_g3_i2_m.22234c100365-g3 | OG0 |
Spolac_c100365_g3_i4_m.22236c100365-g3 | OG0 |
Spolac_c100365_g4_i3_m.22239c100365-g3 | OG0 |
Spolac_c100365_g4_i1_m.22237c100365-g3 | OG0 |
Spolac_c100365_g4_i2_m.22238c100365-g3 | OG0 |
Emue_Em0008g1192a.t1 | OG0 |
Emue_Em0008g1175a.t1 | OG0 |
Emue_Em0131g7a.t1 | OG0 |
Emue_Em0131g8a.t1 | OG0 |
Clacor_Ccor_scaffold13019_01 | OG0 |
Lcom_lcpid33038 | OG0 |
Lcom_lcpid7893 | OG0 |
Gracom_Gcom_scaffold1108_01 | OG0 |
Leuniv_Lniv_C762808_01 | OG0 |
Leuniv_Lniv_C812752_01 | OG0 |
Scil_scict010839.1 | OG0 |
Leuniv_Lniv_C768925_01 | OG0 |
Eupdun_Edun_sb10648702 | OG0 |
Eupdun_Edun_sb10648703 | OG0 |
Eupdun_Edun_sb10648700 | OG0 |
Eupdun_Edun_sb10648701 | OG0 |
Vmul_SimC385694 | OG0 |
Vmul_sbMO443349 | OG0 |
Coemet_Cmet_scaffold22320_01 | OG0 |
Coeast_Cast_sb788890 | OG0 |
Coeast_Cast_sb788891 | OG0 |
Lampan_Lpan_C419943_01 | OG0 |
Beraby_Baby_sb334372 | OG0 |
Mlei_ML2399.g8.i1 | OG0 |
Horcal_Hcv1.av93.c3.g1223.i1 | OG0 |
Pbac_sb2658733 | OG0 |
Pbac_sb2663272 | OG0 |
Plajan_Pjan_C572666_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P011324 | OG0 |
Ocar_g6201 | OG0 |
TrH2_TrispH2_004677-RA | OG0 |
Tadh_TriadT57690 | OG0 |
HoiH11_mRNA9432 | OG0 |
HoiH23_PlH23_004831-RA | OG0 |
HoiH06_mRNA2638 | OG0 |
Hhon_g01780.t1 | OG0 |
HoiH04_mRNA16764 | OG0 |
Cscu_XM_023388734.1 | OG0 |
Tcas_TC009344_002 | OG0 |
Hsap_ENST00000675666 | OG0 |
Skow_XM_006811857.1 | OG0 |
Spur_XM_030987315.1 | OG0 |
Bralan_BLAG11000861_1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA170133 | OG0 |
Cgig_XM_034479980.1 | OG0 |
Acagra_model.g31029.t1 | OG0 |
Acagra_model.g31029.t1.1.5d3b8d37 | OG0 |
Morvir_scaffold65.g5.t1 | OG0 |
Hvul_g17459_1 | OG0 |
Chem_TCONS_00018673 | OG0 |
Chem_TCONS_00018678 | OG0 |
Morvir_scaffold65.g3.t1 | OG0 |
Sanmal_Sma_020678-T1 | OG0 |
Sanmal_Sma_020677-T1 | OG0 |
Nemnom_g4086.t1 | OG0 |
Rhoesc_Res_010039-T1 | OG0 |
Sanmal_Sma_020675-T1 | OG0 |
Aaur_scaffold643.g5.t2 | OG0 |
Xesp_013007-T1 | OG0 |
Dgig_XM_028562749.1 | OG0 |
Dgig_XM_028538085.1 | OG0 |
Exapal_XM_021039633.1 | OG0 |
Nvec_v1g210242 | OG0 |
Exapal_XM_021060869.2 | OG0 |
Nvec_v1g120114 | OG0 |
Nvec_v1g210239 | OG0 |
Fspp_ffun1.m4.20046.m1 | OG0 |
Gasp_gasp1.m3.20665.m1 | OG0 |
Gfas_gfas1.m1.3791.m1 | OG0 |
Spis_XP_022809362_1 | OG0 |
Pocdam_XM_027197685.1 | OG0 |
Amil_Amillepora32298-RA | OG0 |
Gfas_gfas1.m1.3790.m1 | OG0 |
Amil_Amillepora32299-RA | OG0 |
Adig_XM_015915597.1 | OG0 |
Adig_XM_015915586.1 | OG0 |
0.416667