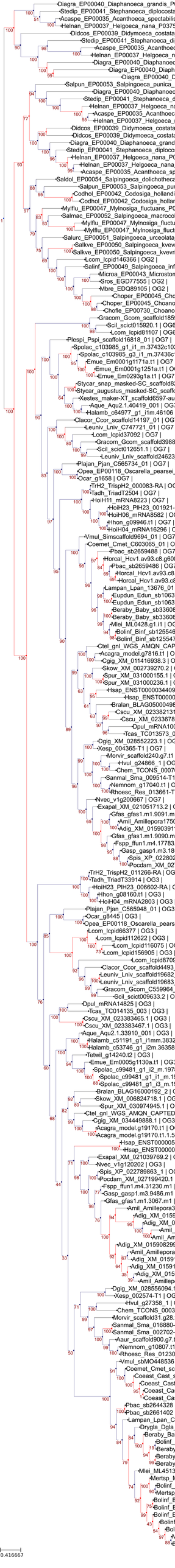
Diagra_EP00040_Diaphanoeca_grandis_P031049 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P007128 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P020964 | OG0 |
Helnan_EP00037_Helgoeca_nana_P037521 | OG0 |
Didcos_EP00039_Didymoeca_costata_P018208 | OG1 |
Stedip_EP00041_Stephanoeca_diplocostata_P007019 | OG1 |
Acaspe_EP00035_Acanthoeca_spectabilis_P038891 | OG1 |
Helnan_EP00037_Helgoeca_nana_P017515 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P030995 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P014427 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P022621 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P023587 | OG4 |
Diagra_EP00040_Diaphanoeca_grandis_P036767 | OG4 |
Stedip_EP00041_Stephanoeca_diplocostata_P026433 | OG4 |
Helnan_EP00037_Helgoeca_nana_P022663 | OG4 |
Acaspe_EP00035_Acanthoeca_spectabilis_P034957 | OG4 |
Helnan_EP00037_Helgoeca_nana_P022662 | OG4 |
Didcos_EP00039_Didymoeca_costata_P018895 | OG5 |
Didcos_EP00039_Didymoeca_costata_P013585 | OG5 |
Diagra_EP00040_Diaphanoeca_grandis_P007054 | OG5 |
Stedip_EP00041_Stephanoeca_diplocostata_P017126 | OG5 |
Helnan_EP00037_Helgoeca_nana_P010821 | OG5 |
Helnan_EP00037_Helgoeca_nana_P010820 | OG5 |
Acaspe_EP00035_Acanthoeca_spectabilis_P035889 | OG5 |
Saldol_EP00054_Salpingoeca_dolichothecata_P003588 | OG2 |
Salpun_EP00053_Salpingoeca_punica_P000121 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P037085 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P037086 | OG2 |
Mylflu_EP00047_Mylnosiga_fluctuans_P024692 | OG2 |
Salmac_EP00052_Salpingoeca_macrocollata_P018257 | OG2 |
Mylflu_EP00047_Mylnosiga_fluctuans_P024694 | OG2 |
Mylflu_EP00047_Mylnosiga_fluctuans_P007978 | OG2 |
Salurc_EP00051_Salpingoeca_urceolata_P013311 | OG2 |
Salkve_EP00050_Salpingoeca_kvevrii_P016593 | OG2 |
Salkve_EP00050_Salpingoeca_kvevrii_P016594 | OG2 |
Lcom_lcpid146366 | OG2 |
Salinf_EP00049_Salpingoeca_infusionum_P003737 | OG2 |
Microa_EP00043_Microstomoeca_roanoka_P008439 | OG2 |
Sros_EGD77555 | OG2 |
Mbre_EDQ89105 | OG2 |
Choper_EP00045_Choanoeca_perplexa_P002238 | OG2 |
Choper_EP00045_Choanoeca_perplexa_P002239 | OG2 |
Chofle_EP00730_Choanoeca_flexa_P013355 | OG2 |
Gracom_Gcom_scaffold18598_01 | OG6 |
Scil_scict015920.1 | OG6 |
Lcom_lcpid81107 | OG6 |
Plespi_Pspi_scaffold16818_01 | OG7 |
Spolac_c103985_g1_i1_m.37432c103985-g1 | OG7 |
Spolac_c103985_g3_i1_m.37436c103985-g1 | OG7 |
Emue_Em0001g1171a.t1 | OG7 |
Emue_Em0001g1251a.t1 | OG7 |
Emue_Em0293g1a.t1 | OG7 |
Stycar_snap_masked-SC_scaffold82370-processed-gene-0.0-mRNA-1 | OG7 |
Stycar_augustus_masked-SC_scaffold47344-processed-gene-0.0-mRNA-1 | OG7 |
Xestes_maker-XT_scaffold6597-augustus-gene-0.44-mRNA-1 | OG7 |
Aque_Aqu2.1.40419_001 | OG7 |
Halamb_c64977_g1_i1m.46106 | OG7 |
Clacor_Ccor_scaffold14197_01 | OG7 |
Leuniv_Lniv_C747721_01 | OG7 |
Lcom_lcpid37092 | OG7 |
Gracom_Gcom_scaffold3988_01 | OG7 |
Scil_scict012651.1 | OG7 |
Leuniv_Lniv_scaffold24623_01 | OG7 |
Plajan_Pjan_C565734_01 | OG7 |
Opea_EP00118_Oscarella_pearsei_P026539 | OG7 |
Ocar_g1658 | OG7 |
TrH2_TrispH2_000083-RA | OG7 |
Tadh_TriadT12504 | OG7 |
HoiH11_mRNA8223 | OG7 |
HoiH23_PIH23_001921-RA | OG7 |
HoiH06_mRNA8582 | OG7 |
Hhon_g09946.t1 | OG7 |
HoiH04_mRNA16296 | OG7 |
Vmul_Simscaffold9694_01 | OG7 |
Coemet_Cmet_C603065_01 | OG7 |
Pbac_sb2659488 | OG7 |
Horcal_Hcv1.av93.c8.g608.i1 | OG7 |
Pbac_sb2659486 | OG7 |
Horcal_Hcv1.av93.c8.g608.i3 | OG7 |
Horcal_Hcv1.av93.c8.g608.i2 | OG7 |
Lampan_Lpan_13676_01 | OG7 |
Eupdun_Edun_sb10637605 | OG7 |
Eupdun_Edun_sb10637606 | OG7 |
Beraby_Baby_sb336083 | OG7 |
Beraby_Baby_sb336082 | OG7 |
Mlei_ML0428.g1.i1 | OG7 |
Bolinf_Binf_sb1255469 | OG7 |
Bolinf_Binf_sb1255470 | OG7 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA178665 | OG7 |
Acagra_model.g7816.t1 | OG7 |
Cgig_XM_011416938.3 | OG7 |
Skow_XM_002739270.2 | OG7 |
Spur_XM_031000155.1 | OG7 |
Spur_XM_031000236.1 | OG7 |
Hsap_ENST00000344096 | OG7 |
Hsap_ENST00000036709 | OG7 |
Bralan_BLAG05000498_1 | OG7 |
Cscu_XM_023382131.1 | OG7 |
Cscu_XM_023367826.1 | OG7 |
Dpul_mRNA10038 | OG7 |
Tcas_TC013573_001 | OG7 |
Dgig_XM_028552223.1 | OG7 |
Xesp_004365-T1 | OG7 |
Morvir_scaffold240.g7.t1 | OG7 |
Hvul_g24866_1 | OG7 |
Chem_TCONS_00070178 | OG7 |
Sanmal_Sma_009514-T1 | OG7 |
Nemnom_g17040.t1 | OG7 |
Rhoesc_Res_013661-T1 | OG7 |
Nvec_v1g200667 | OG7 |
Exapal_XM_021051713.2 | OG7 |
Gfas_gfas1.m1.9091.m1 | OG7 |
Amil_Amillepora17505-RA | OG7 |
Adig_XM_015903911.1 | OG7 |
Gfas_gfas1.m1.9090.m1 | OG7 |
Fspp_ffun1.m4.17783.m1 | OG7 |
Gasp_gasp1.m3.16594.m1 | OG7 |
Spis_XP_022802770_1 | OG7 |
Pocdam_XM_027199409.1 | OG7 |
TrH2_TrispH2_011266-RA | OG3 |
Tadh_TriadT33914 | OG3 |
HoiH23_PIH23_006602-RA | OG3 |
Hhon_g08160.t1 | OG3 |
HoiH04_mRNA2803 | OG3 |
Plajan_Pjan_C565948_01 | OG3 |
Ocar_g8445 | OG3 |
Opea_EP00118_Oscarella_pearsei_P005515 | OG3 |
Lcom_lcpid66377 | OG3 |
Lcom_lcpid112622 | OG3 |
Lcom_lcpid116075 | OG3 |
Lcom_lcpid156905 | OG3 |
Lcom_lcpid87091 | OG3 |
Clacor_Ccor_scaffold4493_01 | OG3 |
Leuniv_Lniv_scaffold19682_01 | OG3 |
Leuniv_Lniv_scaffold19683_01 | OG3 |
Gracom_Gcom_C559964_01 | OG3 |
Scil_scict009633.2 | OG3 |
Dpul_mRNA14825 | OG3 |
Tcas_TC014135_003 | OG3 |
Cscu_XM_023383465.1 | OG3 |
Cscu_XM_023383467.1 | OG3 |
Aque_Aqu2.1.33910_001 | OG3 |
Halamb_c51191_g1_i1mm.3832 | OG3 |
Halamb_c53746_g1_i2m.36358 | OG3 |
Tetwil_g14240.t2 | OG3 |
Emue_Em0005g1130a.t1 | OG3 |
Spolac_c99481_g1_i2_m.19781c99481-g1 | OG3 |
Spolac_c99481_g1_i1_m.19780c99481-g1 | OG3 |
Spolac_c99481_g1_i3_m.19782c99481-g1 | OG3 |
Bralan_BLAG16000192_2 | OG3 |
Skow_XM_006824718.1 | OG3 |
Spur_XM_030974945.1 | OG3 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA149906 | OG3 |
Cgig_XM_034449888.1 | OG3 |
Acagra_model.g19170.t1 | OG3 |
Acagra_model.g19170.t1.1.5d3b8992 | OG3 |
Hsap_ENST00000543431 | OG3 |
Hsap_ENST00000536588 | OG3 |
Exapal_XM_021039769.2 | OG3 |
Nvec_v1g120202 | OG3 |
Spis_XP_022789863_1 | OG3 |
Pocdam_XM_027199420.1 | OG3 |
Fspp_ffun1.m4.31230.m1 | OG3 |
Gasp_gasp1.m3.9486.m1 | OG3 |
Gfas_gfas1.m1.3067.m1 | OG3 |
Amil_Amillepora33882-RA | OG3 |
Adig_XM_015916572.1 | OG3 |
Adig_XM_015892994.1 | OG3 |
Amil_Amillepora30126-RA | OG3 |
Amil_Amillepora30145-RA | OG3 |
Adig_XM_015908299.1 | OG3 |
Amil_Amillepora30115-RA | OG3 |
Adig_XM_015918223.1 | OG3 |
Adig_XM_015915899.1 | OG3 |
Adig_XM_015919923.1 | OG3 |
Amil_Amillepora35415-RA | OG3 |
Dgig_XM_028556094.1 | OG3 |
Xesp_002574-T1 | OG3 |
Hvul_g27358_1 | OG3 |
Chem_TCONS_00030495 | OG3 |
Morvir_scaffold31.g28.t1 | OG3 |
Sanmal_Sma_016880-T1 | OG3 |
Sanmal_Sma_002702-T1 | OG3 |
Aaur_scaffold900.g7.t2 | OG3 |
Nemnom_g10807.t1 | OG3 |
Rhoesc_Res_012304-T1 | OG3 |
Vmul_sbMO448536 | OG3 |
Coemet_Cmet_scaffold2295_01 | OG3 |
Coeast_Cast_sb801098 | OG3 |
Coeast_Cast_sb801096 | OG3 |
Coeast_Cast_sb801095 | OG3 |
Coeast_Cast_sb801097 | OG3 |
Pbac_sb2644328 | OG3 |
Pbac_sb2661402 | OG3 |
Lampan_Lpan_C416451_01 | OG3 |
Drygla_Dgla_sb292319 | OG3 |
Beraby_Baby_sb341357 | OG3 |
Bolinf_Binf_sb1191863 | OG3 |
Beraby_Baby_sb341358 | OG3 |
Beraby_Baby_sb341355 | OG3 |
Beraby_Baby_sb341356 | OG3 |
Mlei_ML4513.g12.i1 | OG3 |
Mertsp_Mesp_sb561976 | OG3 |
Bolinf_Binf_sb1219763 | OG3 |
Mertsp_Mesp_sb561975 | OG3 |
Bolinf_Binf_sb1236136 | OG3 |
Bolinf_Binf_sb1236137 | OG3 |
Bolinf_Binf_sb1236133 | OG3 |
Bolinf_Binf_sb1236134 | OG3 |
Bolinf_Binf_sb1236135 | OG3 |
Mertsp_Mesp_sb539675 | OG3 |
Bolinf_Binf_sb1236138 | OG3 |
0.416667