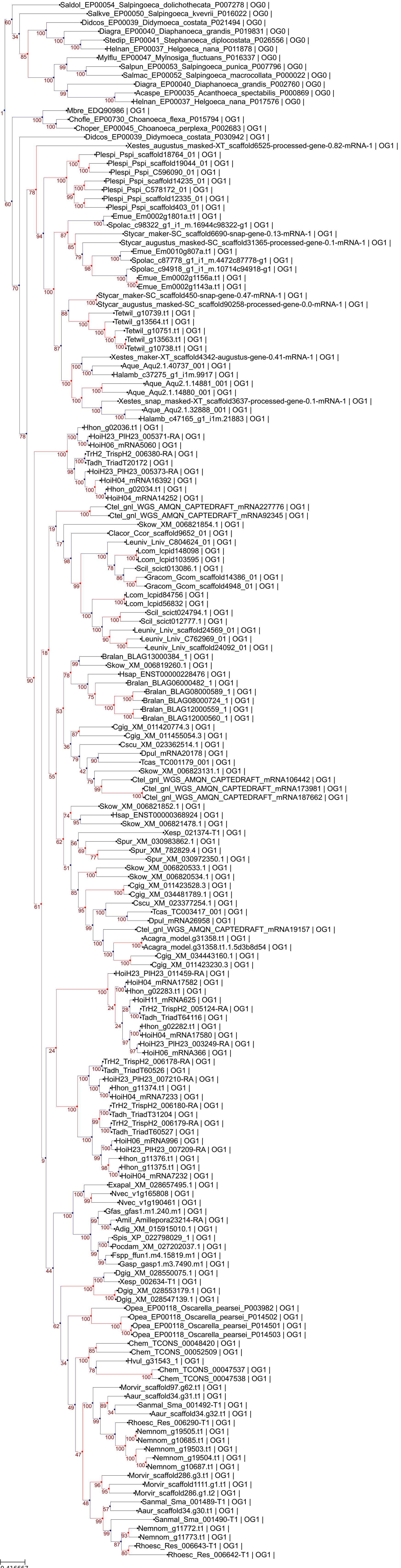
Saldol_EP00054_Salpingoeca_dolichothecata_P007278 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P016022 | OG0 |
Didcos_EP00039_Didymoeca_costata_P021494 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P019831 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P026556 | OG0 |
Helnan_EP00037_Helgoeca_nana_P011878 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P016337 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P007796 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P000022 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P002760 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P000869 | OG0 |
Helnan_EP00037_Helgoeca_nana_P017576 | OG0 |
Mbre_EDQ90986 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P015794 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P002683 | OG1 |
Didcos_EP00039_Didymoeca_costata_P030942 | OG1 |
Xestes_augustus_masked-XT_scaffold6525-processed-gene-0.82-mRNA-1 | OG1 |
Plespi_Pspi_scaffold18764_01 | OG1 |
Plespi_Pspi_scaffold19044_01 | OG1 |
Plespi_Pspi_C596090_01 | OG1 |
Plespi_Pspi_scaffold14235_01 | OG1 |
Plespi_Pspi_C578172_01 | OG1 |
Plespi_Pspi_scaffold12335_01 | OG1 |
Plespi_Pspi_scaffold403_01 | OG1 |
Emue_Em0002g1801a.t1 | OG1 |
Spolac_c98322_g1_i1_m.16944c98322-g1 | OG1 |
Stycar_maker-SC_scaffold6690-snap-gene-0.13-mRNA-1 | OG1 |
Stycar_augustus_masked-SC_scaffold31365-processed-gene-0.1-mRNA-1 | OG1 |
Emue_Em0010g807a.t1 | OG1 |
Spolac_c87778_g1_i1_m.4472c87778-g1 | OG1 |
Spolac_c94918_g1_i1_m.10714c94918-g1 | OG1 |
Emue_Em0002g1156a.t1 | OG1 |
Emue_Em0002g1143a.t1 | OG1 |
Stycar_maker-SC_scaffold450-snap-gene-0.47-mRNA-1 | OG1 |
Stycar_augustus_masked-SC_scaffold90258-processed-gene-0.0-mRNA-1 | OG1 |
Tetwil_g10739.t1 | OG1 |
Tetwil_g13564.t1 | OG1 |
Tetwil_g10751.t1 | OG1 |
Tetwil_g13563.t1 | OG1 |
Tetwil_g10738.t1 | OG1 |
Xestes_maker-XT_scaffold4342-augustus-gene-0.41-mRNA-1 | OG1 |
Aque_Aqu2.1.40737_001 | OG1 |
Halamb_c37275_g1_i1m.9917 | OG1 |
Aque_Aqu2.1.14881_001 | OG1 |
Aque_Aqu2.1.14880_001 | OG1 |
Xestes_snap_masked-XT_scaffold3637-processed-gene-0.1-mRNA-1 | OG1 |
Aque_Aqu2.1.32888_001 | OG1 |
Halamb_c47165_g1_i1m.21883 | OG1 |
Hhon_g02036.t1 | OG1 |
HoiH23_PIH23_005371-RA | OG1 |
HoiH06_mRNA5060 | OG1 |
TrH2_TrispH2_006380-RA | OG1 |
Tadh_TriadT20172 | OG1 |
HoiH23_PIH23_005373-RA | OG1 |
HoiH04_mRNA16392 | OG1 |
Hhon_g02034.t1 | OG1 |
HoiH04_mRNA14252 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA227776 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA92345 | OG1 |
Skow_XM_006821854.1 | OG1 |
Clacor_Ccor_scaffold9652_01 | OG1 |
Leuniv_Lniv_C804624_01 | OG1 |
Lcom_lcpid148098 | OG1 |
Lcom_lcpid103595 | OG1 |
Scil_scict013086.1 | OG1 |
Gracom_Gcom_scaffold14386_01 | OG1 |
Gracom_Gcom_scaffold4948_01 | OG1 |
Lcom_lcpid84756 | OG1 |
Lcom_lcpid56832 | OG1 |
Scil_scict024794.1 | OG1 |
Scil_scict012777.1 | OG1 |
Leuniv_Lniv_scaffold24569_01 | OG1 |
Leuniv_Lniv_C762969_01 | OG1 |
Leuniv_Lniv_scaffold24092_01 | OG1 |
Bralan_BLAG13000384_1 | OG1 |
Skow_XM_006819260.1 | OG1 |
Hsap_ENST00000228476 | OG1 |
Bralan_BLAG06000482_1 | OG1 |
Bralan_BLAG08000589_1 | OG1 |
Bralan_BLAG08000724_1 | OG1 |
Bralan_BLAG12000559_1 | OG1 |
Bralan_BLAG12000560_1 | OG1 |
Cgig_XM_011420774.3 | OG1 |
Cgig_XM_011455054.3 | OG1 |
Cscu_XM_023362514.1 | OG1 |
Dpul_mRNA20178 | OG1 |
Tcas_TC001179_001 | OG1 |
Skow_XM_006823131.1 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA106442 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA173981 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA187662 | OG1 |
Skow_XM_006821852.1 | OG1 |
Hsap_ENST00000368924 | OG1 |
Skow_XM_006821478.1 | OG1 |
Xesp_021374-T1 | OG1 |
Spur_XM_030983862.1 | OG1 |
Spur_XM_782829.4 | OG1 |
Spur_XM_030972350.1 | OG1 |
Skow_XM_006820533.1 | OG1 |
Skow_XM_006820534.1 | OG1 |
Cgig_XM_011423528.3 | OG1 |
Cgig_XM_034481789.1 | OG1 |
Cscu_XM_023377254.1 | OG1 |
Tcas_TC003417_001 | OG1 |
Dpul_mRNA26958 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA19157 | OG1 |
Acagra_model.g31358.t1 | OG1 |
Acagra_model.g31358.t1.1.5d3b8d54 | OG1 |
Cgig_XM_034443160.1 | OG1 |
Cgig_XM_011423230.3 | OG1 |
HoiH23_PIH23_011459-RA | OG1 |
HoiH04_mRNA17582 | OG1 |
Hhon_g02283.t1 | OG1 |
HoiH11_mRNA625 | OG1 |
TrH2_TrispH2_005124-RA | OG1 |
Tadh_TriadT64116 | OG1 |
Hhon_g02282.t1 | OG1 |
HoiH04_mRNA17580 | OG1 |
HoiH23_PIH23_003249-RA | OG1 |
HoiH06_mRNA366 | OG1 |
TrH2_TrispH2_006178-RA | OG1 |
Tadh_TriadT60526 | OG1 |
HoiH23_PIH23_007210-RA | OG1 |
Hhon_g11374.t1 | OG1 |
HoiH04_mRNA7233 | OG1 |
TrH2_TrispH2_006180-RA | OG1 |
Tadh_TriadT31204 | OG1 |
TrH2_TrispH2_006179-RA | OG1 |
Tadh_TriadT60527 | OG1 |
HoiH06_mRNA996 | OG1 |
HoiH23_PIH23_007209-RA | OG1 |
Hhon_g11376.t1 | OG1 |
Hhon_g11375.t1 | OG1 |
HoiH04_mRNA7232 | OG1 |
Exapal_XM_028657495.1 | OG1 |
Nvec_v1g165808 | OG1 |
Nvec_v1g190461 | OG1 |
Gfas_gfas1.m1.240.m1 | OG1 |
Amil_Amillepora23214-RA | OG1 |
Adig_XM_015915010.1 | OG1 |
Spis_XP_022798029_1 | OG1 |
Pocdam_XM_027202037.1 | OG1 |
Fspp_ffun1.m4.15819.m1 | OG1 |
Gasp_gasp1.m3.7490.m1 | OG1 |
Dgig_XM_028550075.1 | OG1 |
Xesp_002634-T1 | OG1 |
Dgig_XM_028553179.1 | OG1 |
Dgig_XM_028547139.1 | OG1 |
Opea_EP00118_Oscarella_pearsei_P003982 | OG1 |
Opea_EP00118_Oscarella_pearsei_P014502 | OG1 |
Opea_EP00118_Oscarella_pearsei_P014501 | OG1 |
Opea_EP00118_Oscarella_pearsei_P014503 | OG1 |
Chem_TCONS_00048420 | OG1 |
Chem_TCONS_00052509 | OG1 |
Hvul_g31543_1 | OG1 |
Chem_TCONS_00047537 | OG1 |
Chem_TCONS_00047538 | OG1 |
Morvir_scaffold97.g62.t1 | OG1 |
Aaur_scaffold34.g31.t1 | OG1 |
Sanmal_Sma_001492-T1 | OG1 |
Aaur_scaffold34.g32.t1 | OG1 |
Rhoesc_Res_006290-T1 | OG1 |
Nemnom_g19505.t1 | OG1 |
Nemnom_g10685.t1 | OG1 |
Nemnom_g19503.t1 | OG1 |
Nemnom_g19504.t1 | OG1 |
Nemnom_g10687.t1 | OG1 |
Morvir_scaffold286.g3.t1 | OG1 |
Morvir_scaffold1111.g1.t1 | OG1 |
Morvir_scaffold286.g1.t2 | OG1 |
Sanmal_Sma_001489-T1 | OG1 |
Aaur_scaffold34.g30.t1 | OG1 |
Sanmal_Sma_001490-T1 | OG1 |
Nemnom_g11772.t1 | OG1 |
Nemnom_g11773.t1 | OG1 |
Rhoesc_Res_006643-T1 | OG1 |
Rhoesc_Res_006642-T1 | OG1 |
0.416667