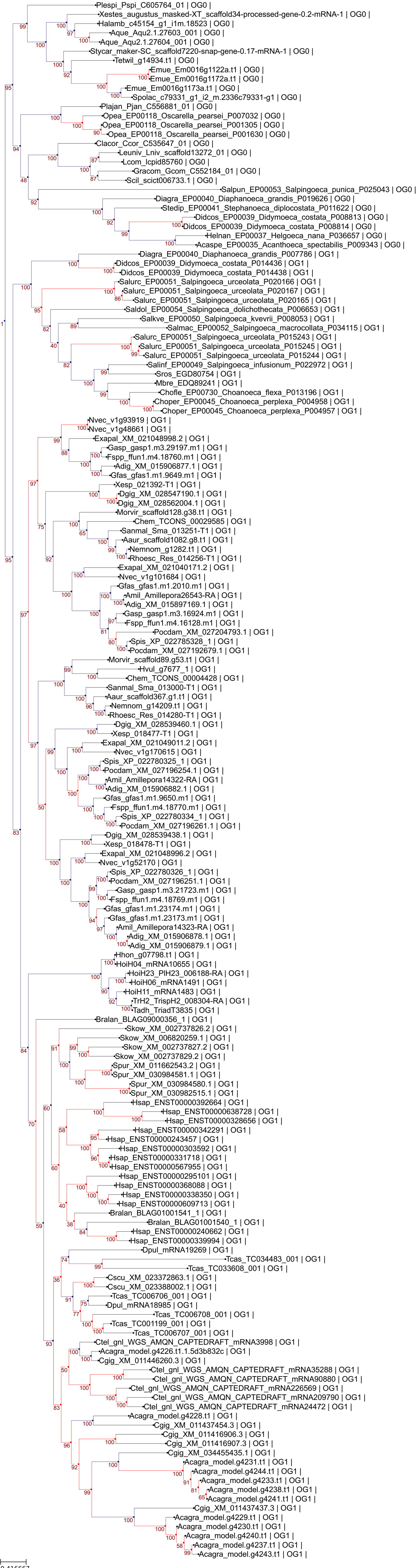
Plespi_Pspi_C605764_01 | OG0 |
Xestes_augustus_masked-XT_scaffold34-processed-gene-0.2-mRNA-1 | OG0 |
Halamb_c45154_g1_i1m.18523 | OG0 |
Aque_Aqu2.1.27603_001 | OG0 |
Aque_Aqu2.1.27604_001 | OG0 |
Stycar_maker-SC_scaffold7220-snap-gene-0.17-mRNA-1 | OG0 |
Tetwil_g14934.t1 | OG0 |
Emue_Em0016g1122a.t1 | OG0 |
Emue_Em0016g1172a.t1 | OG0 |
Emue_Em0016g1173a.t1 | OG0 |
Spolac_c79331_g1_i2_m.2336c79331-g1 | OG0 |
Plajan_Pjan_C556881_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P007032 | OG0 |
Opea_EP00118_Oscarella_pearsei_P001305 | OG0 |
Opea_EP00118_Oscarella_pearsei_P001630 | OG0 |
Clacor_Ccor_C535647_01 | OG0 |
Leuniv_Lniv_scaffold13272_01 | OG0 |
Lcom_lcpid85760 | OG0 |
Gracom_Gcom_C552184_01 | OG0 |
Scil_scict006733.1 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P025043 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P019626 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P011622 | OG0 |
Didcos_EP00039_Didymoeca_costata_P008813 | OG0 |
Didcos_EP00039_Didymoeca_costata_P008814 | OG0 |
Helnan_EP00037_Helgoeca_nana_P036657 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P009343 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P007786 | OG1 |
Didcos_EP00039_Didymoeca_costata_P014436 | OG1 |
Didcos_EP00039_Didymoeca_costata_P014438 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P020166 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P020167 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P020165 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P006653 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P008053 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P034115 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P015243 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P015245 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P015244 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P022972 | OG1 |
Sros_EGD80754 | OG1 |
Mbre_EDQ89241 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P013196 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P004958 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P004957 | OG1 |
Nvec_v1g93919 | OG1 |
Nvec_v1g48661 | OG1 |
Exapal_XM_021048998.2 | OG1 |
Gasp_gasp1.m3.29197.m1 | OG1 |
Fspp_ffun1.m4.18760.m1 | OG1 |
Adig_XM_015906877.1 | OG1 |
Gfas_gfas1.m1.9649.m1 | OG1 |
Xesp_021392-T1 | OG1 |
Dgig_XM_028547190.1 | OG1 |
Dgig_XM_028562004.1 | OG1 |
Morvir_scaffold128.g38.t1 | OG1 |
Chem_TCONS_00029585 | OG1 |
Sanmal_Sma_013251-T1 | OG1 |
Aaur_scaffold1082.g8.t1 | OG1 |
Nemnom_g1282.t1 | OG1 |
Rhoesc_Res_014256-T1 | OG1 |
Exapal_XM_021040171.2 | OG1 |
Nvec_v1g101684 | OG1 |
Gfas_gfas1.m1.2010.m1 | OG1 |
Amil_Amillepora26543-RA | OG1 |
Adig_XM_015897169.1 | OG1 |
Gasp_gasp1.m3.16924.m1 | OG1 |
Fspp_ffun1.m4.16128.m1 | OG1 |
Pocdam_XM_027204793.1 | OG1 |
Spis_XP_022785328_1 | OG1 |
Pocdam_XM_027192679.1 | OG1 |
Morvir_scaffold89.g53.t1 | OG1 |
Hvul_g7677_1 | OG1 |
Chem_TCONS_00004428 | OG1 |
Sanmal_Sma_013000-T1 | OG1 |
Aaur_scaffold367.g1.t1 | OG1 |
Nemnom_g14209.t1 | OG1 |
Rhoesc_Res_014280-T1 | OG1 |
Dgig_XM_028539460.1 | OG1 |
Xesp_018477-T1 | OG1 |
Exapal_XM_021049011.2 | OG1 |
Nvec_v1g170615 | OG1 |
Spis_XP_022780325_1 | OG1 |
Pocdam_XM_027196254.1 | OG1 |
Amil_Amillepora14322-RA | OG1 |
Adig_XM_015906882.1 | OG1 |
Gfas_gfas1.m1.9650.m1 | OG1 |
Fspp_ffun1.m4.18770.m1 | OG1 |
Spis_XP_022780334_1 | OG1 |
Pocdam_XM_027196261.1 | OG1 |
Dgig_XM_028539438.1 | OG1 |
Xesp_018478-T1 | OG1 |
Exapal_XM_021048996.2 | OG1 |
Nvec_v1g52170 | OG1 |
Spis_XP_022780326_1 | OG1 |
Pocdam_XM_027196251.1 | OG1 |
Gasp_gasp1.m3.21723.m1 | OG1 |
Fspp_ffun1.m4.18769.m1 | OG1 |
Gfas_gfas1.m1.23174.m1 | OG1 |
Gfas_gfas1.m1.23173.m1 | OG1 |
Amil_Amillepora14323-RA | OG1 |
Adig_XM_015906878.1 | OG1 |
Adig_XM_015906879.1 | OG1 |
Hhon_g07798.t1 | OG1 |
HoiH04_mRNA10655 | OG1 |
HoiH23_PlH23_006188-RA | OG1 |
HoiH06_mRNA1491 | OG1 |
HoiH11_mRNA1483 | OG1 |
TrH2_TrispH2_008304-RA | OG1 |
Tadh_TriadT3835 | OG1 |
Bralan_BLAG09000356_1 | OG1 |
Skow_XM_002737826.2 | OG1 |
Skow_XM_006820259.1 | OG1 |
Skow_XM_002737827.2 | OG1 |
Skow_XM_002737829.2 | OG1 |
Spur_XM_011662543.2 | OG1 |
Spur_XM_030984581.1 | OG1 |
Spur_XM_030984580.1 | OG1 |
Spur_XM_030982515.1 | OG1 |
Hsap_ENST00000392664 | OG1 |
Hsap_ENST00000638728 | OG1 |
Hsap_ENST00000328656 | OG1 |
Hsap_ENST00000342291 | OG1 |
Hsap_ENST00000243457 | OG1 |
Hsap_ENST00000303592 | OG1 |
Hsap_ENST00000331718 | OG1 |
Hsap_ENST00000567955 | OG1 |
Hsap_ENST00000295101 | OG1 |
Hsap_ENST00000368088 | OG1 |
Hsap_ENST00000338350 | OG1 |
Hsap_ENST00000609713 | OG1 |
Bralan_BLAG01001541_1 | OG1 |
Bralan_BLAG01001540_1 | OG1 |
Hsap_ENST00000240662 | OG1 |
Hsap_ENST00000339994 | OG1 |
Dpul_mRNA19269 | OG1 |
Tcas_TC034483_001 | OG1 |
Tcas_TC033608_001 | OG1 |
Cscu_XM_023372863.1 | OG1 |
Cscu_XM_023388002.1 | OG1 |
Tcas_TC006706_001 | OG1 |
Dpul_mRNA18985 | OG1 |
Tcas_TC006708_001 | OG1 |
Tcas_TC001199_001 | OG1 |
Tcas_TC006707_001 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA3998 | OG1 |
Acagra_model.g4226.t1.1.5d3b832c | OG1 |
Cgig_XM_011446260.3 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA35288 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA90880 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA226569 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA209790 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA24472 | OG1 |
Acagra_model.g4228.t1 | OG1 |
Cgig_XM_011437454.3 | OG1 |
Cgig_XM_011416906.3 | OG1 |
Cgig_XM_011416907.3 | OG1 |
Cgig_XM_034455435.1 | OG1 |
Acagra_model.g4231.t1 | OG1 |
Acagra_model.g4244.t1 | OG1 |
Acagra_model.g4233.t1 | OG1 |
Acagra_model.g4238.t1 | OG1 |
Acagra_model.g4241.t1 | OG1 |
Cgig_XM_011437437.3 | OG1 |
Acagra_model.g4229.t1 | OG1 |
Acagra_model.g4230.t1 | OG1 |
Acagra_model.g4240.t1 | OG1 |
Acagra_model.g4237.t1 | OG1 |
Acagra_model.g4243.t1 | OG1 |
0.416667