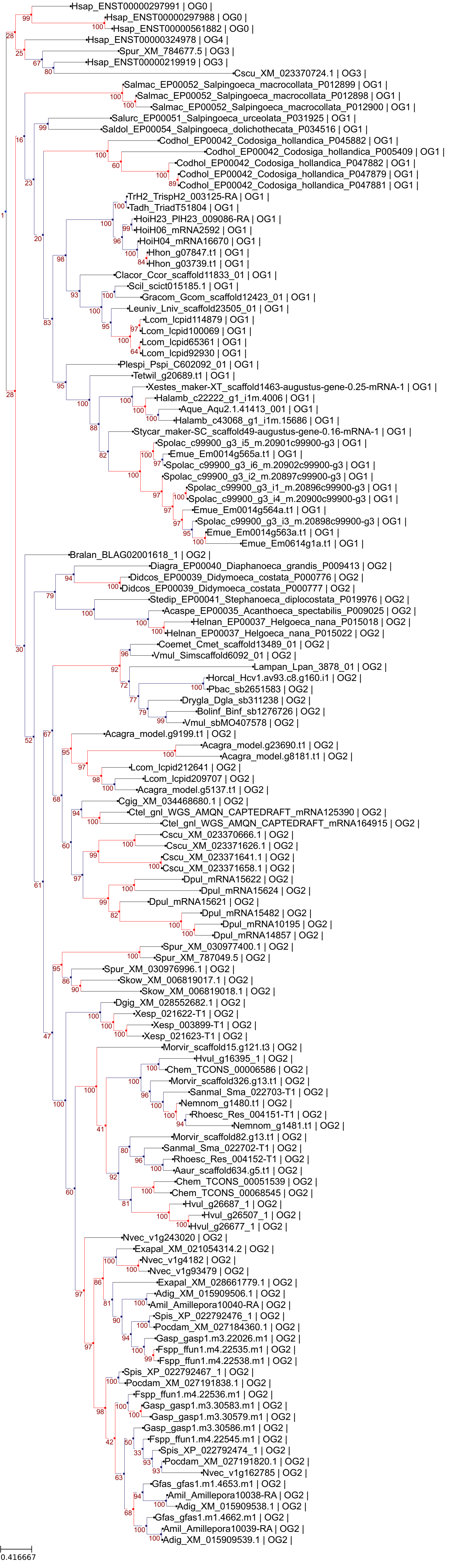
Hsap_ENST00000297991 | OG0 |
Hsap_ENST00000297988 | OG0 |
Hsap_ENST00000561882 | OG0 |
Hsap_ENST00000324978 | OG4 |
Spur_XM_784677.5 | OG3 |
Hsap_ENST00000219919 | OG3 |
Cscu_XM_023370724.1 | OG3 |
Salmac_EP00052_Salpingoeca_macrocollata_P012899 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P012898 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P012900 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P031925 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P034516 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P045882 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P005409 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P047882 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P047879 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P047881 | OG1 |
TrH2_TrispH2_003125-RA | OG1 |
Tadh_TriadT51804 | OG1 |
HoiH23_PlH23_009086-RA | OG1 |
HoiH06_mRNA2592 | OG1 |
HoiH04_mRNA16670 | OG1 |
Hhon_g07847.t1 | OG1 |
Hhon_g03739.t1 | OG1 |
Clacor_Ccor_scaffold11833_01 | OG1 |
Scil_scict015185.1 | OG1 |
Gracom_Gcom_scaffold12423_01 | OG1 |
Leuniv_Lniv_scaffold23505_01 | OG1 |
Lcom_lcpid114879 | OG1 |
Lcom_lcpid100069 | OG1 |
Lcom_lcpid65361 | OG1 |
Lcom_lcpid92930 | OG1 |
Plespi_Pspi_C602092_01 | OG1 |
Tetwil_g20689.t1 | OG1 |
Xestes_maker-XT_scaffold1463-augustus-gene-0.25-mRNA-1 | OG1 |
Halamb_c22222_g1_i1m.4006 | OG1 |
Aque_Aqu2.1.41413_001 | OG1 |
Halamb_c43068_g1_i1m.15686 | OG1 |
Stycar_maker-SC_scaffold49-augustus-gene-0.16-mRNA-1 | OG1 |
Spolac_c99900_g3_i5_m.20901c99900-g3 | OG1 |
Emue_Em0014g565a.t1 | OG1 |
Spolac_c99900_g3_i6_m.20902c99900-g3 | OG1 |
Spolac_c99900_g3_i2_m.20897c99900-g3 | OG1 |
Spolac_c99900_g3_i1_m.20896c99900-g3 | OG1 |
Spolac_c99900_g3_i4_m.20900c99900-g3 | OG1 |
Emue_Em0014g564a.t1 | OG1 |
Spolac_c99900_g3_i3_m.20898c99900-g3 | OG1 |
Emue_Em0014g563a.t1 | OG1 |
Emue_Em0614g1a.t1 | OG1 |
Bralan_BLAG02001618_1 | OG2 |
Diagra_EP00040_Diaphanoeca_grandis_P009413 | OG2 |
Didcos_EP00039_Didymoeca_costata_P000776 | OG2 |
Didcos_EP00039_Didymoeca_costata_P000777 | OG2 |
Stedip_EP00041_Stephanoeca_diplocostata_P019976 | OG2 |
Acaspe_EP00035_Acanthoeca_spectabilis_P009025 | OG2 |
Helnan_EP00037_Helgoeca_nana_P015018 | OG2 |
Helnan_EP00037_Helgoeca_nana_P015022 | OG2 |
Coemet_Cmet_scaffold13489_01 | OG2 |
Vmul_Simscaffold6092_01 | OG2 |
Lampan_Lpan_3878_01 | OG2 |
Horcal_Hcv1.av93.c8.g160.i1 | OG2 |
Pbac_sb2651583 | OG2 |
Drygla_Dgla_sb311238 | OG2 |
Bolinf_Binf_sb1276726 | OG2 |
Vmul_sbMO407578 | OG2 |
Acagra_model.g9199.t1 | OG2 |
Acagra_model.g23690.t1 | OG2 |
Acagra_model.g8181.t1 | OG2 |
Lcom_lcpid212641 | OG2 |
Lcom_lcpid209707 | OG2 |
Acagra_model.g5137.t1 | OG2 |
Cgig_XM_034468680.1 | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA125390 | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA164915 | OG2 |
Cscu_XM_023370666.1 | OG2 |
Cscu_XM_023371626.1 | OG2 |
Cscu_XM_023371641.1 | OG2 |
Cscu_XM_023371658.1 | OG2 |
Dpul_mRNA15622 | OG2 |
Dpul_mRNA15624 | OG2 |
Dpul_mRNA15621 | OG2 |
Dpul_mRNA15482 | OG2 |
Dpul_mRNA10195 | OG2 |
Dpul_mRNA14857 | OG2 |
Spur_XM_030977400.1 | OG2 |
Spur_XM_787049.5 | OG2 |
Spur_XM_030976996.1 | OG2 |
Skow_XM_006819017.1 | OG2 |
Skow_XM_006819018.1 | OG2 |
Dgig_XM_028552682.1 | OG2 |
Xesp_021622-T1 | OG2 |
Xesp_003899-T1 | OG2 |
Xesp_021623-T1 | OG2 |
Morvir_scaffold15.g121.t3 | OG2 |
Hvul_g16395_1 | OG2 |
Chem_TCONS_00006586 | OG2 |
Morvir_scaffold326.g13.t1 | OG2 |
Sanmal_Sma_022703-T1 | OG2 |
Nemnom_g1480.t1 | OG2 |
Rhoesc_Res_004151-T1 | OG2 |
Nemnom_g1481.t1 | OG2 |
Morvir_scaffold82.g13.t1 | OG2 |
Sanmal_Sma_022702-T1 | OG2 |
Rhoesc_Res_004152-T1 | OG2 |
Aaur_scaffold634.g5.t1 | OG2 |
Chem_TCONS_00051539 | OG2 |
Chem_TCONS_00068545 | OG2 |
Hvul_g26687_1 | OG2 |
Hvul_g26507_1 | OG2 |
Hvul_g26677_1 | OG2 |
Nvec_v1g243020 | OG2 |
Exapal_XM_021054314.2 | OG2 |
Nvec_v1g4182 | OG2 |
Nvec_v1g93479 | OG2 |
Exapal_XM_028661779.1 | OG2 |
Adig_XM_015909506.1 | OG2 |
Amil_Amillepora10040-RA | OG2 |
Spis_XP_022792476_1 | OG2 |
Pocdam_XM_027184360.1 | OG2 |
Gasp_gasp1.m3.22026.m1 | OG2 |
Fspp_ffun1.m4.22535.m1 | OG2 |
Fspp_ffun1.m4.22538.m1 | OG2 |
Spis_XP_022792467_1 | OG2 |
Pocdam_XM_027191838.1 | OG2 |
Fspp_ffun1.m4.22536.m1 | OG2 |
Gasp_gasp1.m3.30583.m1 | OG2 |
Gasp_gasp1.m3.30579.m1 | OG2 |
Gasp_gasp1.m3.30586.m1 | OG2 |
Fspp_ffun1.m4.22545.m1 | OG2 |
Spis_XP_022792474_1 | OG2 |
Pocdam_XM_027191820.1 | OG2 |
Nvec_v1g162785 | OG2 |
Gfas_gfas1.m1.4653.m1 | OG2 |
Amil_Amillepora10038-RA | OG2 |
Adig_XM_015909538.1 | OG2 |
Gfas_gfas1.m1.4662.m1 | OG2 |
Amil_Amillepora10039-RA | OG2 |
Adig_XM_015909539.1 | OG2 |
0.416667