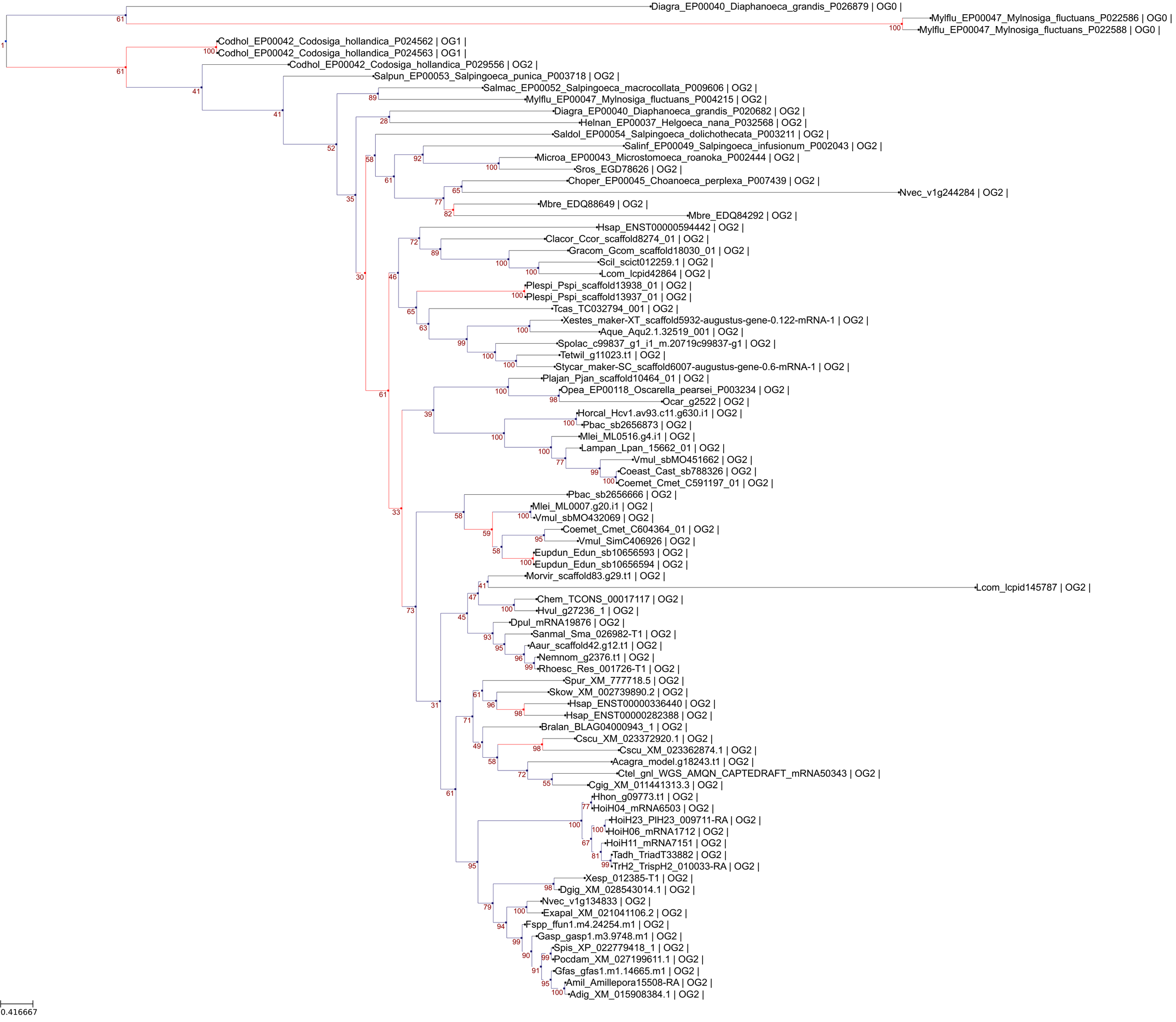
Diagra_EP00040_Diaphanoeca_grandis_P026879 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P022586 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P022588 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P024562 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P024563 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P029556 | OG2 |
Salpun_EP00053_Salpingoeca_punica_P003718 | OG2 |
Salmac_EP00052_Salpingoeca_macrocollata_P009606 | OG2 |
Mylflu_EP00047_Mylnosiga_fluctuans_P004215 | OG2 |
Diagra_EP00040_Diaphanoeca_grandis_P020682 | OG2 |
Helnan_EP00037_Helgoeca_nana_P032568 | OG2 |
Saldol_EP00054_Salpingoeca_dolichothecata_P003211 | OG2 |
Salinf_EP00049_Salpingoeca_infusionum_P002043 | OG2 |
Microa_EP00043_Microstomoeca_roanoka_P002444 | OG2 |
Sros_EGD78626 | OG2 |
Choper_EP00045_Choanoeca_perplexa_P007439 | OG2 |
Nvec_v1g244284 | OG2 |
Mbre_EDQ88649 | OG2 |
Mbre_EDQ84292 | OG2 |
Hsap_ENST00000594442 | OG2 |
Clacor_Ccor_scaffold8274_01 | OG2 |
Gracom_Gcom_scaffold18030_01 | OG2 |
Scil_scict012259.1 | OG2 |
Lcom_lcpid42864 | OG2 |
Plespi_Pspi_scaffold13938_01 | OG2 |
Plespi_Pspi_scaffold13937_01 | OG2 |
Tcas_TC032794_001 | OG2 |
Xestes_maker-XT_scaffold5932-augustus-gene-0.122-mRNA-1 | OG2 |
Aque_Aqu2.1.32519_001 | OG2 |
Spolac_c99837_g1_i1_m.20719c99837-g1 | OG2 |
Tetwil_g11023.t1 | OG2 |
Stycar_maker-SC_scaffold6007-augustus-gene-0.6-mRNA-1 | OG2 |
Plajan_Pjan_scaffold10464_01 | OG2 |
Opea_EP00118_Oscarella_pearsei_P003234 | OG2 |
Ocar_g2522 | OG2 |
Horcal_Hcv1.av93.c11.g630.i1 | OG2 |
Pbac_sb2656873 | OG2 |
Mlei_ML0516.g4.i1 | OG2 |
Lampan_Lpan_15662_01 | OG2 |
Vmul_sbMO451662 | OG2 |
Coeast_Cast_sb788326 | OG2 |
Coemet_Cmet_C591197_01 | OG2 |
Pbac_sb2656666 | OG2 |
Mlei_ML0007.g20.i1 | OG2 |
Vmul_sbMO432069 | OG2 |
Coemet_Cmet_C604364_01 | OG2 |
Vmul_SimC406926 | OG2 |
Eupdun_Edun_sb10656593 | OG2 |
Eupdun_Edun_sb10656594 | OG2 |
Morvir_scaffold83.g29.t1 | OG2 |
Lcom_lcpid145787 | OG2 |
Chem_TCONS_00017117 | OG2 |
Hvul_g27236_1 | OG2 |
Dpul_mRNA19876 | OG2 |
Sanmal_Sma_026982-T1 | OG2 |
Aaur_scaffold42.g12.t1 | OG2 |
Nemnom_g2376.t1 | OG2 |
Rhoesc_Res_001726-T1 | OG2 |
Spur_XM_777718.5 | OG2 |
Skow_XM_002739890.2 | OG2 |
Hsap_ENST00000336440 | OG2 |
Hsap_ENST00000282388 | OG2 |
Bralan_BLAG04000943_1 | OG2 |
Cscu_XM_023372920.1 | OG2 |
Cscu_XM_023362874.1 | OG2 |
Acagra_model.g18243.t1 | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA50343 | OG2 |
Cgig_XM_011441313.3 | OG2 |
Hhon_g09773.t1 | OG2 |
HoiH04_mRNA6503 | OG2 |
HoiH23_PlH23_009711-RA | OG2 |
HoiH06_mRNA1712 | OG2 |
HoiH11_mRNA7151 | OG2 |
Tadh_TriadT33882 | OG2 |
TrH2_TrispH2_010033-RA | OG2 |
Xesp_012385-T1 | OG2 |
Dgig_XM_028543014.1 | OG2 |
Nvec_v1g134833 | OG2 |
Exapal_XM_021041106.2 | OG2 |
Fspp_ffun1.m4.24254.m1 | OG2 |
Gasp_gasp1.m3.9748.m1 | OG2 |
Spis_XP_022779418_1 | OG2 |
Pocdam_XM_027199611.1 | OG2 |
Gfas_gfas1.m1.14665.m1 | OG2 |
Amil_Amillepora15508-RA | OG2 |
Adig_XM_015908384.1 | OG2 |
0.416667