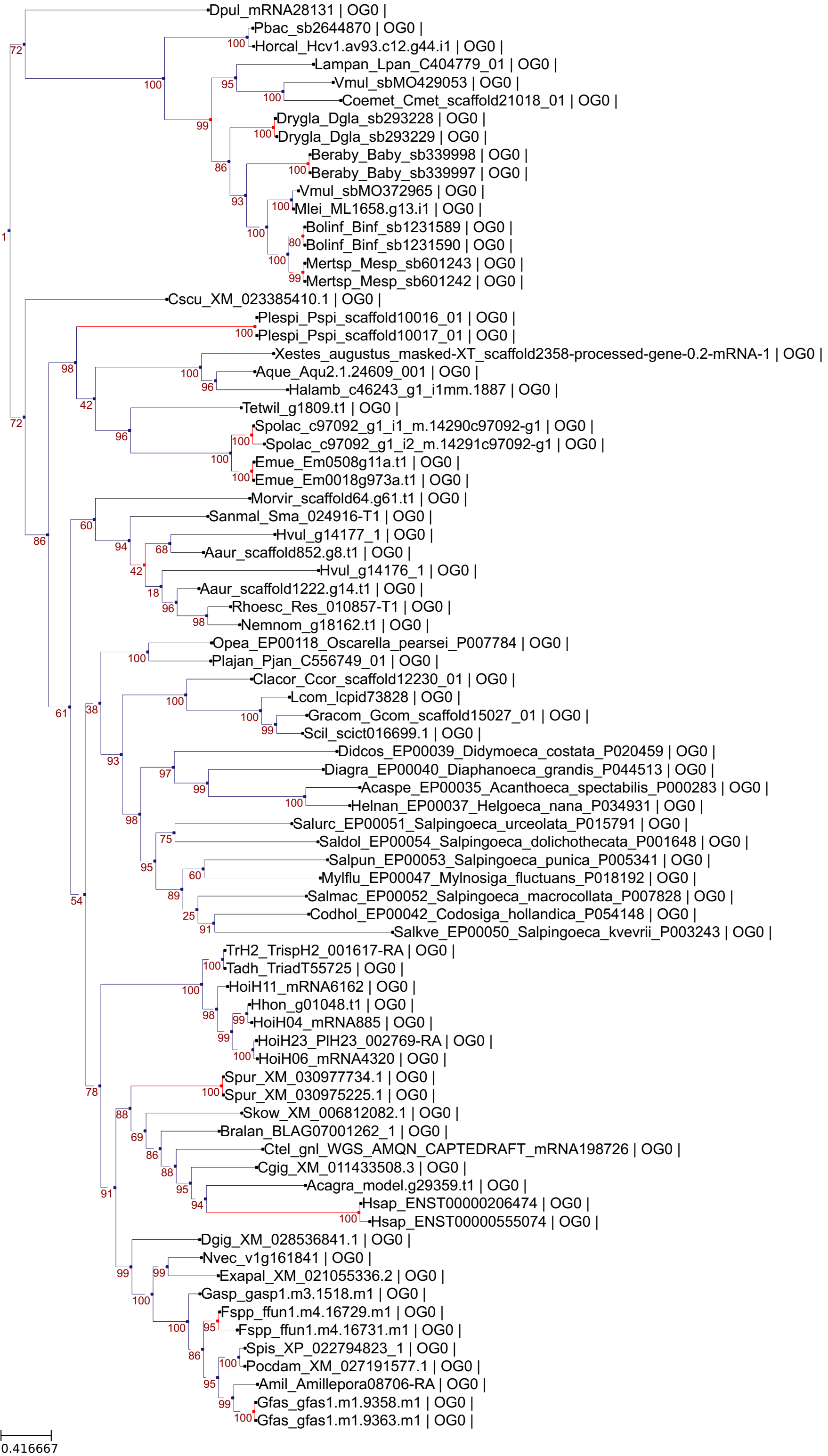
Dpul_mRNA28131 | OG0 |
Pbac_sb2644870 | OG0 |
Horcal_Hcv1.av93.c12.g44.i1 | OG0 |
Lampan_Lpan_C404779_01 | OG0 |
Vmul_sbMO429053 | OG0 |
Coemet_Cmet_scaffold21018_01 | OG0 |
Drygla_Dgla_sb293228 | OG0 |
Drygla_Dgla_sb293229 | OG0 |
Beraby_Baby_sb339998 | OG0 |
Beraby_Baby_sb339997 | OG0 |
Vmul_sbMO372965 | OG0 |
Mlei_ML1658.g13.i1 | OG0 |
Bolinf_Binf_sb1231589 | OG0 |
Bolinf_Binf_sb1231590 | OG0 |
Mertsp_Mesp_sb601243 | OG0 |
Mertsp_Mesp_sb601242 | OG0 |
Cscu_XM_023385410.1 | OG0 |
Plespi_Pspi_scaffold10016_01 | OG0 |
Plespi_Pspi_scaffold10017_01 | OG0 |
Xestes_augustus_masked-XT_scaffold2358-processed-gene-0.2-mRNA-1 | OG0 |
Aque_Aqu2.1.24609_001 | OG0 |
Halamb_c46243_g1_i1mm.1887 | OG0 |
Tetwil_g1809.t1 | OG0 |
Spolac_c97092_g1_i1_m.14290c97092-g1 | OG0 |
Spolac_c97092_g1_i2_m.14291c97092-g1 | OG0 |
Emue_Em0508g11a.t1 | OG0 |
Emue_Em0018g973a.t1 | OG0 |
Morvir_scaffold64.g61.t1 | OG0 |
Sanmal_Sma_024916-T1 | OG0 |
Hvul_g14177_1 | OG0 |
Aaur_scaffold852.g8.t1 | OG0 |
Hvul_g14176_1 | OG0 |
Aaur_scaffold1222.g14.t1 | OG0 |
Rhoesc_Res_010857-T1 | OG0 |
Nemnom_g18162.t1 | OG0 |
Opea_EP00118_Oscarella_pearsei_P007784 | OG0 |
Plajan_Pjan_C556749_01 | OG0 |
Clacor_Ccor_scaffold12230_01 | OG0 |
Lcom_lcpid73828 | OG0 |
Gracom_Gcom_scaffold15027_01 | OG0 |
Scil_scict016699.1 | OG0 |
Didcos_EP00039_Didymoeca_costata_P020459 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P044513 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P000283 | OG0 |
Helnan_EP00037_Helgoeca_nana_P034931 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P015791 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P001648 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P005341 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P018192 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P007828 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P054148 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P003243 | OG0 |
TrH2_TrispH2_001617-RA | OG0 |
Tadh_TriadT55725 | OG0 |
HoiH11_mRNA6162 | OG0 |
Hhon_g01048.t1 | OG0 |
HoiH04_mRNA885 | OG0 |
HoiH23_PlH23_002769-RA | OG0 |
HoiH06_mRNA4320 | OG0 |
Spur_XM_030977734.1 | OG0 |
Spur_XM_030975225.1 | OG0 |
Skow_XM_006812082.1 | OG0 |
Bralan_BLAG07001262_1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA198726 | OG0 |
Cgig_XM_011433508.3 | OG0 |
Acagra_model.g29359.t1 | OG0 |
Hsap_ENST00000206474 | OG0 |
Hsap_ENST00000555074 | OG0 |
Dgig_XM_028536841.1 | OG0 |
Nvec_v1g161841 | OG0 |
Exapal_XM_021055336.2 | OG0 |
Gasp_gasp1.m3.1518.m1 | OG0 |
Fspp_ffun1.m4.16729.m1 | OG0 |
Fspp_ffun1.m4.16731.m1 | OG0 |
Spis_XP_022794823_1 | OG0 |
Pocdam_XM_027191577.1 | OG0 |
Amil_Amillepora08706-RA | OG0 |
Gfas_gfas1.m1.9358.m1 | OG0 |
Gfas_gfas1.m1.9363.m1 | OG0 |
0.416667