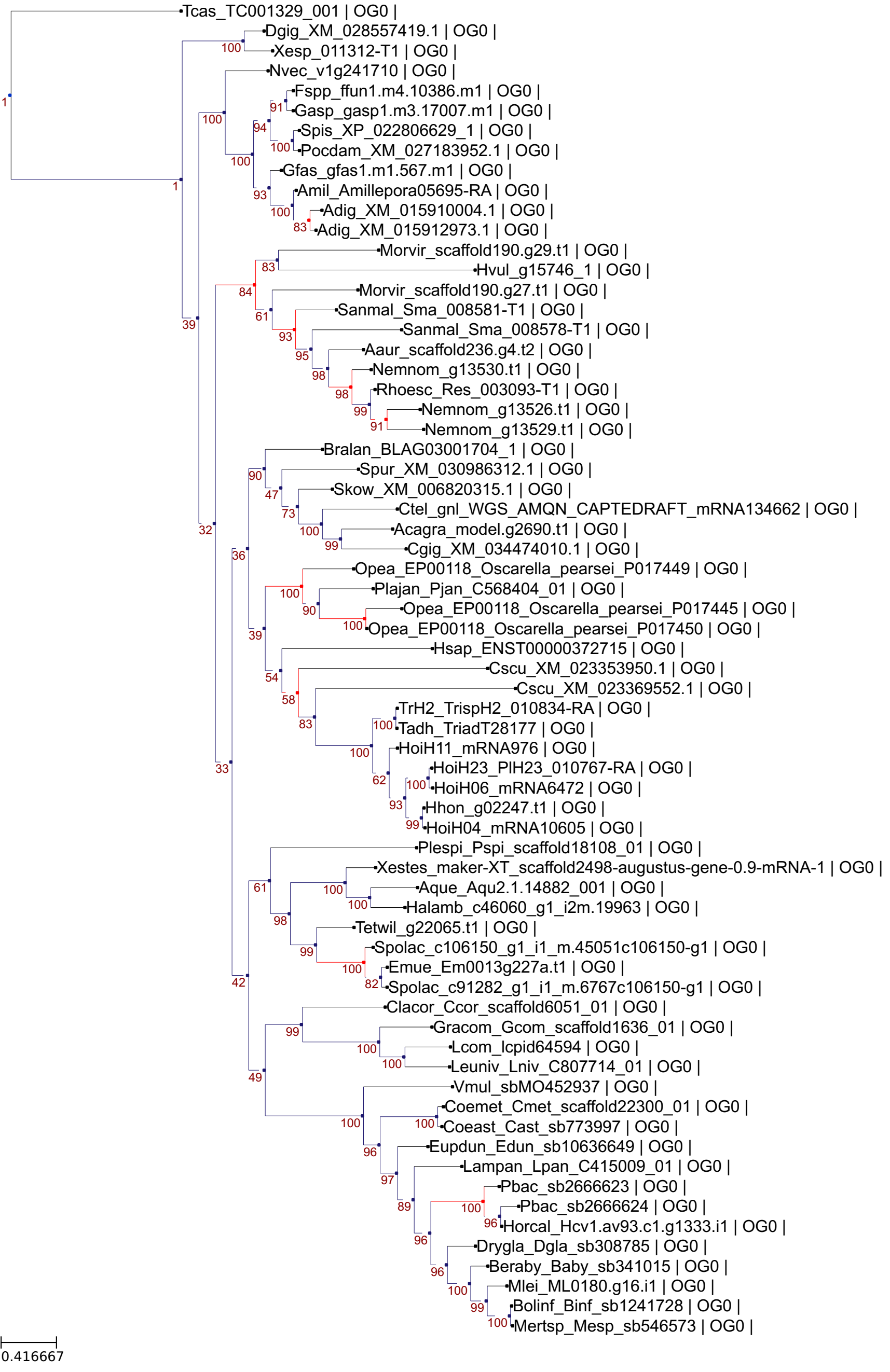
Tcas_TC001329_001 | OG0 |
Dgig_XM_028557419.1 | OG0 |
Xesp_011312-T1 | OG0 |
Nvec_v1g241710 | OG0 |
Fspp_ffun1.m4.10386.m1 | OG0 |
Gasp_gasp1.m3.17007.m1 | OG0 |
Spis_XP_022806629_1 | OG0 |
Pocdam_XM_027183952.1 | OG0 |
Gfas_gfas1.m1.567.m1 | OG0 |
Amil_Amillepora05695-RA | OG0 |
Adig_XM_015910004.1 | OG0 |
Adig_XM_015912973.1 | OG0 |
Morvir_scaffold190.g29.t1 | OG0 |
Hvul_g15746_1 | OG0 |
Morvir_scaffold190.g27.t1 | OG0 |
Sanmal_Sma_008581-T1 | OG0 |
Sanmal_Sma_008578-T1 | OG0 |
Aaur_scaffold236.g4.t2 | OG0 |
Nemnom_g13530.t1 | OG0 |
Rhoesc_Res_003093-T1 | OG0 |
Nemnom_g13526.t1 | OG0 |
Nemnom_g13529.t1 | OG0 |
Bralan_BLAG03001704_1 | OG0 |
Spur_XM_030986312.1 | OG0 |
Skow_XM_006820315.1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA134662 | OG0 |
Acagra_model.g2690.t1 | OG0 |
Cgig_XM_034474010.1 | OG0 |
Opea_EP00118_Oscarella_pearsei_P017449 | OG0 |
Plajan_Pjan_C568404_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P017445 | OG0 |
Opea_EP00118_Oscarella_pearsei_P017450 | OG0 |
Hsap_ENST00000372715 | OG0 |
Cscu_XM_023353950.1 | OG0 |
Cscu_XM_023369552.1 | OG0 |
TrH2_TrispH2_010834-RA | OG0 |
Tadh_TriadT28177 | OG0 |
HoiH11_mRNA976 | OG0 |
HoiH23_PlH23_010767-RA | OG0 |
HoiH06_mRNA6472 | OG0 |
Hhon_g02247.t1 | OG0 |
HoiH04_mRNA10605 | OG0 |
Plespi_Pspi_scaffold18108_01 | OG0 |
Xestes_maker-XT_scaffold2498-augustus-gene-0.9-mRNA-1 | OG0 |
Aque_Aqu2.1.14882_001 | OG0 |
Halamb_c46060_g1_i2m.19963 | OG0 |
Tetwil_g22065.t1 | OG0 |
Spolac_c106150_g1_i1_m.45051c106150-g1 | OG0 |
Emue_Em0013g227a.t1 | OG0 |
Spolac_c91282_g1_i1_m.6767c106150-g1 | OG0 |
Clacor_Ccor_scaffold6051_01 | OG0 |
Gracom_Gcom_scaffold1636_01 | OG0 |
Lcom_lcpid64594 | OG0 |
Leuniv_Lniv_C807714_01 | OG0 |
Vmul_sbMO452937 | OG0 |
Coemet_Cmet_scaffold22300_01 | OG0 |
Coeast_Cast_sb773997 | OG0 |
Eupdun_Edun_sb10636649 | OG0 |
Lampan_Lpan_C415009_01 | OG0 |
Pbac_sb2666623 | OG0 |
Pbac_sb2666624 | OG0 |
Horcal_Hcv1.av93.c1.g1333.i1 | OG0 |
Drygla_Dgla_sb308785 | OG0 |
Beraby_Baby_sb341015 | OG0 |
Mlei_ML0180.g16.i1 | OG0 |
Bolinf_Binf_sb1241728 | OG0 |
Mertsp_Mesp_sb546573 | OG0 |
0.416667