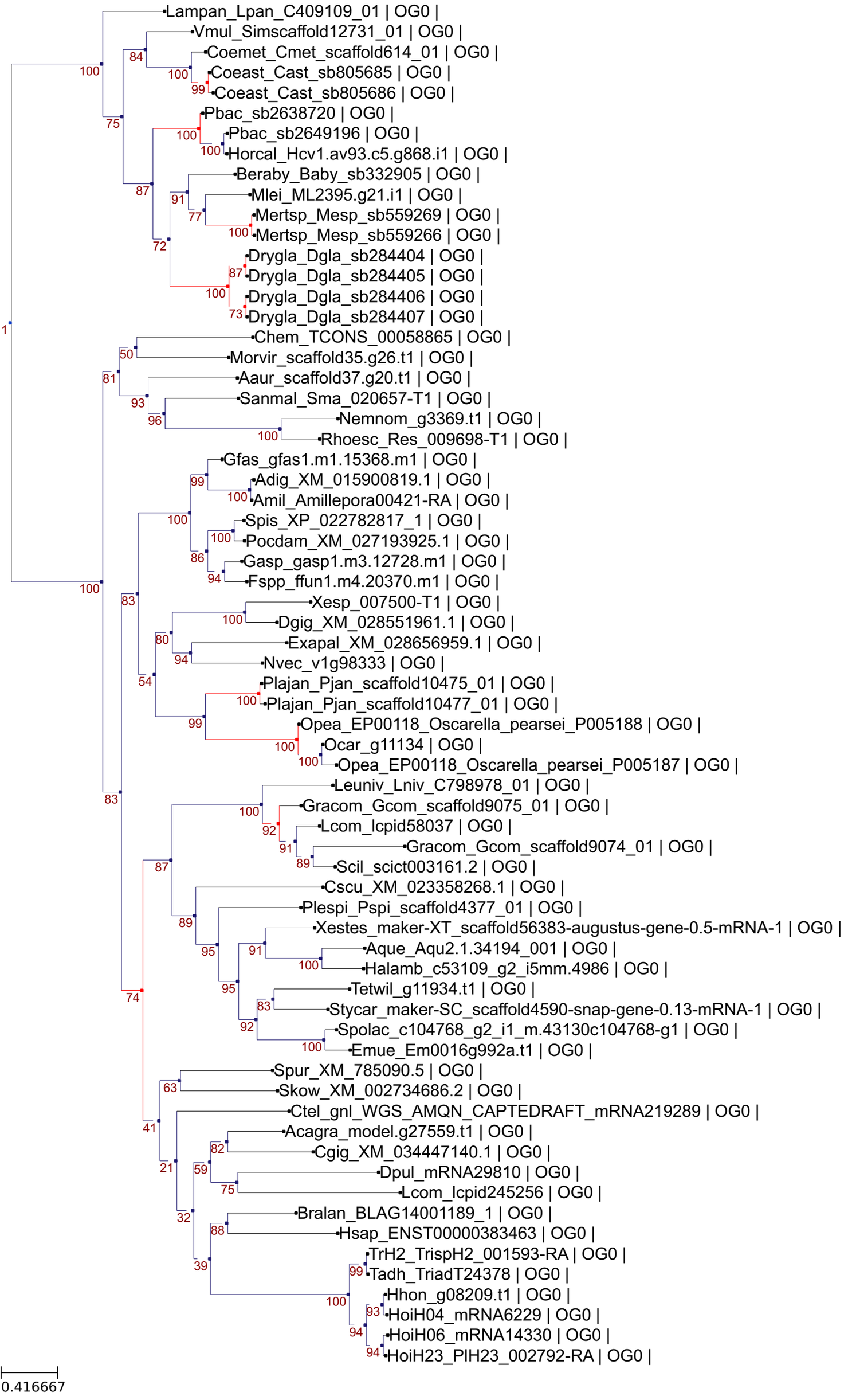
Lampan_Lpan_C409109_01 | OG0 |
Vmul_Simscaffold12731_01 | OG0 |
Coemet_Cmet_scaffold614_01 | OG0 |
Coeast_Cast_sb805685 | OG0 |
Coeast_Cast_sb805686 | OG0 |
Pbac_sb2638720 | OG0 |
Pbac_sb2649196 | OG0 |
Horcal_Hcv1.av93.c5.g868.i1 | OG0 |
Beraby_Baby_sb332905 | OG0 |
Mlei_ML2395.g21.i1 | OG0 |
Mertsp_Mesp_sb559269 | OG0 |
Mertsp_Mesp_sb559266 | OG0 |
Drygla_Dgla_sb284404 | OG0 |
Drygla_Dgla_sb284405 | OG0 |
Drygla_Dgla_sb284406 | OG0 |
Drygla_Dgla_sb284407 | OG0 |
Chem_TCONS_00058865 | OG0 |
Morvir_scaffold35.g26.t1 | OG0 |
Aaur_scaffold37.g20.t1 | OG0 |
Sanmal_Sma_020657-T1 | OG0 |
Nemnom_g3369.t1 | OG0 |
Rhoesc_Res_009698-T1 | OG0 |
Gfas_gfas1.m1.15368.m1 | OG0 |
Adig_XM_015900819.1 | OG0 |
Amil_Amillepora00421-RA | OG0 |
Spis_XP_022782817_1 | OG0 |
Pocdam_XM_027193925.1 | OG0 |
Gasp_gasp1.m3.12728.m1 | OG0 |
Fspp_ffun1.m4.20370.m1 | OG0 |
Xesp_007500-T1 | OG0 |
Dgig_XM_028551961.1 | OG0 |
Exapal_XM_028656959.1 | OG0 |
Nvec_v1g98333 | OG0 |
Plajan_Pjan_scaffold10475_01 | OG0 |
Plajan_Pjan_scaffold10477_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P005188 | OG0 |
Ocar_g11134 | OG0 |
Opea_EP00118_Oscarella_pearsei_P005187 | OG0 |
Leuniv_Lniv_C798978_01 | OG0 |
Gracom_Gcom_scaffold9075_01 | OG0 |
Lcom_lcpid58037 | OG0 |
Gracom_Gcom_scaffold9074_01 | OG0 |
Scil_scict003161.2 | OG0 |
Cscu_XM_023358268.1 | OG0 |
Plespi_Pspi_scaffold4377_01 | OG0 |
Xestes_maker-XT_scaffold56383-augustus-gene-0.5-mRNA-1 | OG0 |
Aque_Aqu2.1.34194_001 | OG0 |
Halamb_c53109_g2_i5mm.4986 | OG0 |
Tetwil_g11934.t1 | OG0 |
Stycar_maker-SC_scaffold4590-snap-gene-0.13-mRNA-1 | OG0 |
Spolac_c104768_g2_i1_m.43130c104768-g1 | OG0 |
Emue_Em0016g992a.t1 | OG0 |
Spur_XM_785090.5 | OG0 |
Skow_XM_002734686.2 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA219289 | OG0 |
Acagra_model.g27559.t1 | OG0 |
Cgig_XM_034447140.1 | OG0 |
Dpul_mRNA29810 | OG0 |
Lcom_lcpid245256 | OG0 |
Bralan_BLAG14001189_1 | OG0 |
Hsap_ENST00000383463 | OG0 |
TrH2_TrispH2_001593-RA | OG0 |
Tadh_TriadT24378 | OG0 |
Hhon_g08209.t1 | OG0 |
HoiH04_mRNA6229 | OG0 |
HoiH06_mRNA14330 | OG0 |
HoiH23_PlH23_002792-RA | OG0 |
0.416667