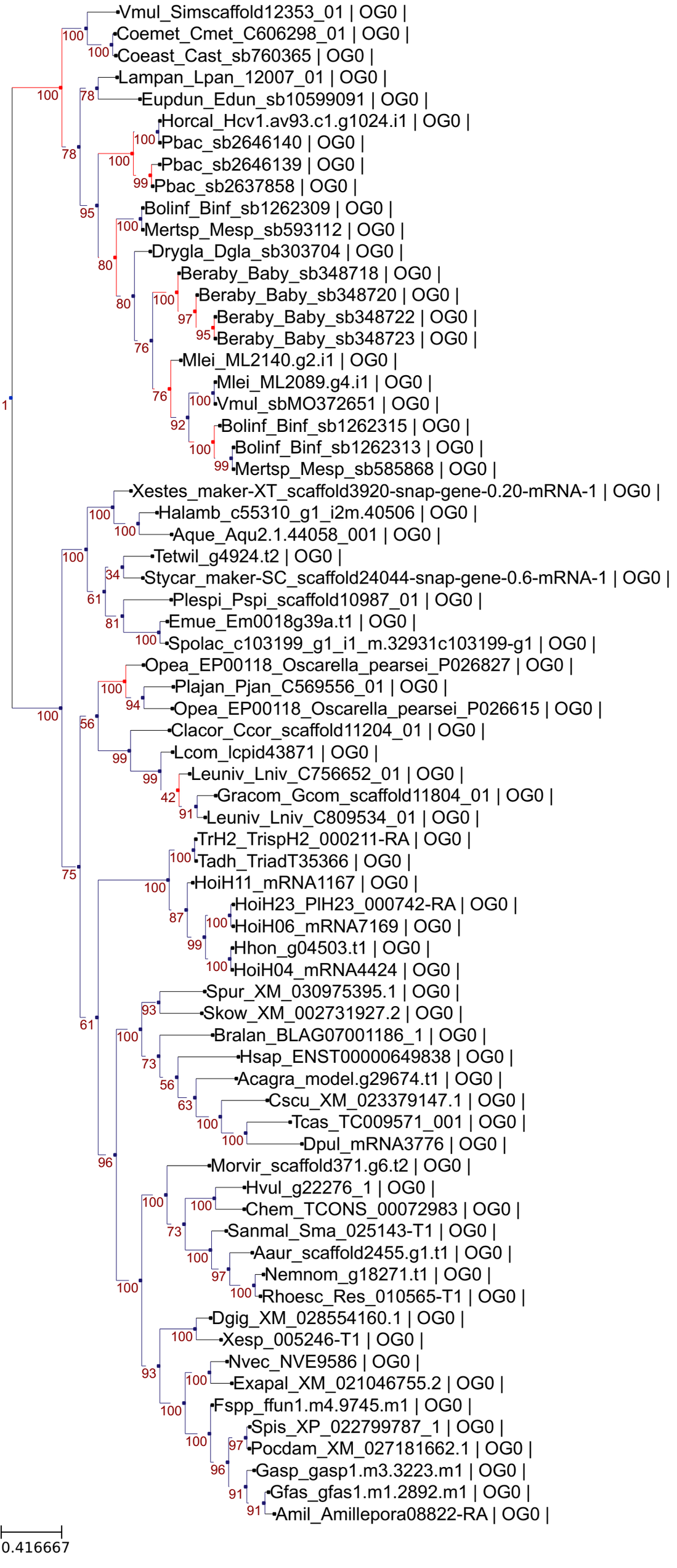
Vmul_Simscaffold12353_01 | OG0 |
Coemet_Cmet_C606298_01 | OG0 |
Coeast_Cast_sb760365 | OG0 |
Lampan_Lpan_12007_01 | OG0 |
Eupdun_Edun_sb10599091 | OG0 |
Horcal_Hcv1.av93.c1.g1024.i1 | OG0 |
Pbac_sb2646140 | OG0 |
Pbac_sb2646139 | OG0 |
Pbac_sb2637858 | OG0 |
Bolinf_Binf_sb1262309 | OG0 |
Mertsp_Mesp_sb593112 | OG0 |
Drygla_Dgla_sb303704 | OG0 |
Beraby_Baby_sb348718 | OG0 |
Beraby_Baby_sb348720 | OG0 |
Beraby_Baby_sb348722 | OG0 |
Beraby_Baby_sb348723 | OG0 |
Mlei_ML2140.g2.i1 | OG0 |
Mlei_ML2089.g4.i1 | OG0 |
Vmul_sbMO372651 | OG0 |
Bolinf_Binf_sb1262315 | OG0 |
Bolinf_Binf_sb1262313 | OG0 |
Mertsp_Mesp_sb585868 | OG0 |
Xestes_maker-XT_scaffold3920-snap-gene-0.20-mRNA-1 | OG0 |
Halamb_c55310_g1_i2m.40506 | OG0 |
Aque_Aqu2.1.44058_001 | OG0 |
Tetwil_g4924.t2 | OG0 |
Stycar_maker-SC_scaffold24044-snap-gene-0.6-mRNA-1 | OG0 |
Plespi_Pspi_scaffold10987_01 | OG0 |
Emue_Em0018g39a.t1 | OG0 |
Spolac_c103199_g1_i1_m.32931c103199-g1 | OG0 |
Opea_EP00118_Oscarella_pearsei_P026827 | OG0 |
Plajan_Pjan_C569556_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P026615 | OG0 |
Clacor_Ccor_scaffold11204_01 | OG0 |
Lcom_lcpid43871 | OG0 |
Leuniv_Lniv_C756652_01 | OG0 |
Gracom_Gcom_scaffold11804_01 | OG0 |
Leuniv_Lniv_C809534_01 | OG0 |
TrH2_TrispH2_000211-RA | OG0 |
Tadh_TriadT35366 | OG0 |
HoiH11_mRNA1167 | OG0 |
HoiH23_PlH23_000742-RA | OG0 |
HoiH06_mRNA7169 | OG0 |
Hhon_g04503.t1 | OG0 |
HoiH04_mRNA4424 | OG0 |
Spur_XM_030975395.1 | OG0 |
Skow_XM_002731927.2 | OG0 |
Bralan_BLAG07001186_1 | OG0 |
Hsap_ENST00000649838 | OG0 |
Acagra_model.g29674.t1 | OG0 |
Cscu_XM_023379147.1 | OG0 |
Tcas_TC009571_001 | OG0 |
Dpul_mRNA3776 | OG0 |
Morvir_scaffold371.g6.t2 | OG0 |
Hvul_g22276_1 | OG0 |
Chem_TCONS_00072983 | OG0 |
Sanmal_Sma_025143-T1 | OG0 |
Aaur_scaffold2455.g1.t1 | OG0 |
Nemnom_g18271.t1 | OG0 |
Rhoesc_Res_010565-T1 | OG0 |
Dgig_XM_028554160.1 | OG0 |
Xesp_005246-T1 | OG0 |
Nvec_NVE9586 | OG0 |
Exapal_XM_021046755.2 | OG0 |
Fspp_ffun1.m4.9745.m1 | OG0 |
Spis_XP_022799787_1 | OG0 |
Pocdam_XM_027181662.1 | OG0 |
Gasp_gasp1.m3.3223.m1 | OG0 |
Gfas_gfas1.m1.2892.m1 | OG0 |
Amil_Amillepora08822-RA | OG0 |
0.416667