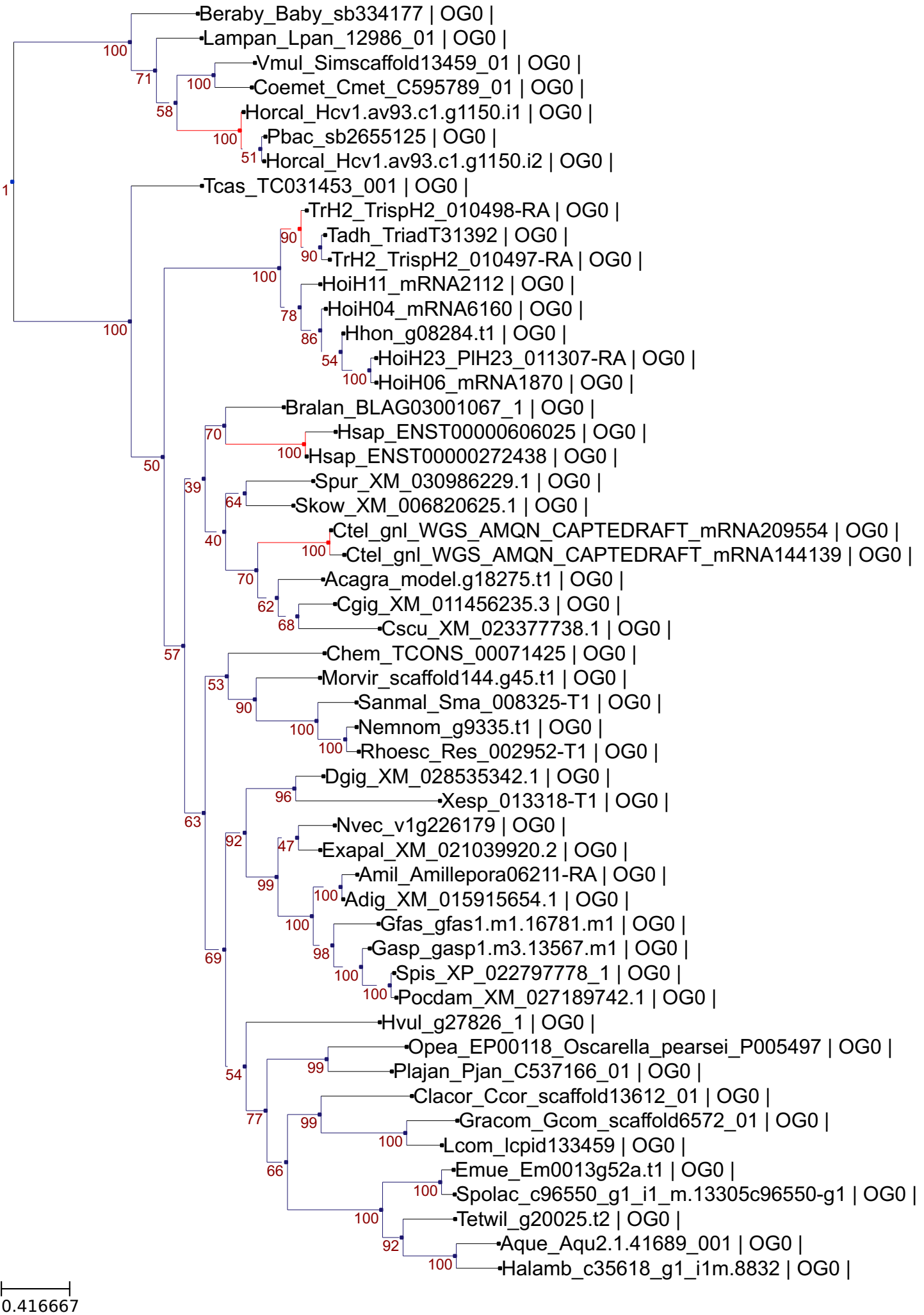
Beraby_Baby_sb334177 | OG0 |
Lampan_Lpan_12986_01 | OG0 |
Vmul_Simscaffold13459_01 | OG0 |
Coemet_Cmet_C595789_01 | OG0 |
Horcal_Hcv1.av93.c1.g1150.i1 | OG0 |
Pbac_sb2655125 | OG0 |
Horcal_Hcv1.av93.c1.g1150.i2 | OG0 |
Tcas_TC031453_001 | OG0 |
TrH2_TrispH2_010498-RA | OG0 |
Tadh_TriadT31392 | OG0 |
TrH2_TrispH2_010497-RA | OG0 |
HoiH11_mRNA2112 | OG0 |
HoiH04_mRNA6160 | OG0 |
Hhon_g08284.t1 | OG0 |
HoiH23_PlH23_011307-RA | OG0 |
HoiH06_mRNA1870 | OG0 |
Bralan_BLAG03001067_1 | OG0 |
Hsap_ENST00000606025 | OG0 |
Hsap_ENST00000272438 | OG0 |
Spur_XM_030986229.1 | OG0 |
Skow_XM_006820625.1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA209554 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA144139 | OG0 |
Acagra_model.g18275.t1 | OG0 |
Cgig_XM_011456235.3 | OG0 |
Cscu_XM_023377738.1 | OG0 |
Chem_TCONS_00071425 | OG0 |
Morvir_scaffold144.g45.t1 | OG0 |
Sanmal_Sma_008325-T1 | OG0 |
Nemnom_g9335.t1 | OG0 |
Rhoesc_Res_002952-T1 | OG0 |
Dgig_XM_028535342.1 | OG0 |
Xesp_013318-T1 | OG0 |
Nvec_v1g226179 | OG0 |
Exapal_XM_021039920.2 | OG0 |
Amil_Amillepora06211-RA | OG0 |
Adig_XM_015915654.1 | OG0 |
Gfas_gfas1.m1.16781.m1 | OG0 |
Gasp_gasp1.m3.13567.m1 | OG0 |
Spis_XP_022797778_1 | OG0 |
Pocdam_XM_027189742.1 | OG0 |
Hvul_g27826_1 | OG0 |
Opea_EP00118_Oscarella_pearsei_P005497 | OG0 |
Plajan_Pjan_C537166_01 | OG0 |
Clacor_Ccor_scaffold13612_01 | OG0 |
Gracom_Gcom_scaffold6572_01 | OG0 |
Lcom_lcpid133459 | OG0 |
Emue_Em0013g52a.t1 | OG0 |
Spolac_c96550_g1_i1_m.13305c96550-g1 | OG0 |
Tetwil_g20025.t2 | OG0 |
Aque_Aqu2.1.41689_001 | OG0 |
Halamb_c35618_g1_i1m.8832 | OG0 |
0.416667