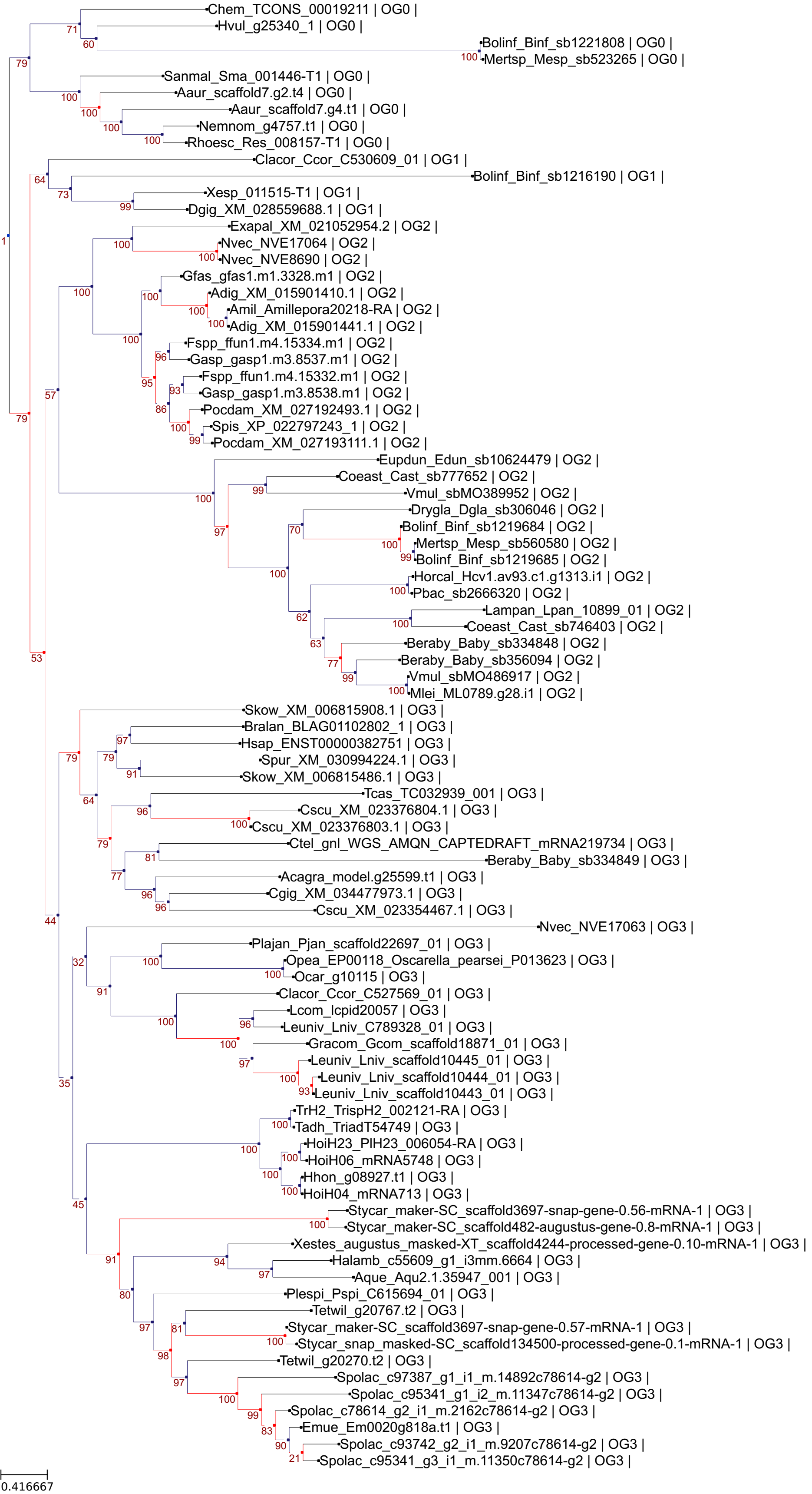
Chem_TCONS_00019211 | OG0 |
Hvul_g25340_1 | OG0 |
Bolinf_Binf_sb1221808 | OG0 |
Mertsp_Mesp_sb523265 | OG0 |
Sanmal_Sma_001446-T1 | OG0 |
Aaur_scaffold7.g2.t4 | OG0 |
Aaur_scaffold7.g4.t1 | OG0 |
Nemnom_g4757.t1 | OG0 |
Rhoesc_Res_008157-T1 | OG0 |
Clacor_Ccor_C530609_01 | OG1 |
Bolinf_Binf_sb1216190 | OG1 |
Xesp_011515-T1 | OG1 |
Dgig_XM_028559688.1 | OG1 |
Exapal_XM_021052954.2 | OG2 |
Nvec_NVE17064 | OG2 |
Nvec_NVE8690 | OG2 |
Gfas_gfas1.m1.3328.m1 | OG2 |
Adig_XM_015901410.1 | OG2 |
Amil_Amillepora20218-RA | OG2 |
Adig_XM_015901441.1 | OG2 |
Fspp_ffun1.m4.15334.m1 | OG2 |
Gasp_gasp1.m3.8537.m1 | OG2 |
Fspp_ffun1.m4.15332.m1 | OG2 |
Gasp_gasp1.m3.8538.m1 | OG2 |
Pocdam_XM_027192493.1 | OG2 |
Spis_XP_022797243_1 | OG2 |
Pocdam_XM_027193111.1 | OG2 |
Eupdun_Edun_sb10624479 | OG2 |
Coeast_Cast_sb777652 | OG2 |
Vmul_sbMO389952 | OG2 |
Drygla_Dgla_sb306046 | OG2 |
Bolinf_Binf_sb1219684 | OG2 |
Mertsp_Mesp_sb560580 | OG2 |
Bolinf_Binf_sb1219685 | OG2 |
Horcal_Hcv1.av93.c1.g1313.i1 | OG2 |
Pbac_sb2666320 | OG2 |
Lampan_Lpan_10899_01 | OG2 |
Coeast_Cast_sb746403 | OG2 |
Beraby_Baby_sb334848 | OG2 |
Beraby_Baby_sb356094 | OG2 |
Vmul_sbMO486917 | OG2 |
Mlei_ML0789.g28.i1 | OG2 |
Skow_XM_006815908.1 | OG3 |
Bralan_BLAG01102802_1 | OG3 |
Hsap_ENST00000382751 | OG3 |
Spur_XM_030994224.1 | OG3 |
Skow_XM_006815486.1 | OG3 |
Tcas_TC032939_001 | OG3 |
Cscu_XM_023376804.1 | OG3 |
Cscu_XM_023376803.1 | OG3 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA219734 | OG3 |
Beraby_Baby_sb334849 | OG3 |
Acagra_model.g25599.t1 | OG3 |
Cgig_XM_034477973.1 | OG3 |
Cscu_XM_023354467.1 | OG3 |
Nvec_NVE17063 | OG3 |
Plajan_Pjan_scaffold22697_01 | OG3 |
Opea_EP00118_Oscarella_pearsei_P013623 | OG3 |
Ocar_g10115 | OG3 |
Clacor_Ccor_C527569_01 | OG3 |
Lcom_lcpid20057 | OG3 |
Leuniv_Lniv_C789328_01 | OG3 |
Gracom_Gcom_scaffold18871_01 | OG3 |
Leuniv_Lniv_scaffold10445_01 | OG3 |
Leuniv_Lniv_scaffold10444_01 | OG3 |
Leuniv_Lniv_scaffold10443_01 | OG3 |
TrH2_TrispH2_002121-RA | OG3 |
Tadh_TriadT54749 | OG3 |
HoiH23_PlH23_006054-RA | OG3 |
HoiH06_mRNA5748 | OG3 |
Hhon_g08927.t1 | OG3 |
HoiH04_mRNA713 | OG3 |
Stycar_maker-SC_scaffold3697-snap-gene-0.56-mRNA-1 | OG3 |
Stycar_maker-SC_scaffold482-augustus-gene-0.8-mRNA-1 | OG3 |
Xestes_augustus_masked-XT_scaffold4244-processed-gene-0.10-mRNA-1 | OG3 |
Halamb_c55609_g1_i3mm.6664 | OG3 |
Aque_Aqu2.1.35947_001 | OG3 |
Plespi_Pspi_C615694_01 | OG3 |
Tetwil_g20767.t2 | OG3 |
Stycar_maker-SC_scaffold3697-snap-gene-0.57-mRNA-1 | OG3 |
Stycar_snap_masked-SC_scaffold134500-processed-gene-0.1-mRNA-1 | OG3 |
Tetwil_g20270.t2 | OG3 |
Spolac_c97387_g1_i1_m.14892c78614-g2 | OG3 |
Spolac_c95341_g1_i2_m.11347c78614-g2 | OG3 |
Spolac_c78614_g2_i1_m.2162c78614-g2 | OG3 |
Emue_Em0020g818a.t1 | OG3 |
Spolac_c93742_g2_i1_m.9207c78614-g2 | OG3 |
Spolac_c95341_g3_i1_m.11350c78614-g2 | OG3 |
0.416667