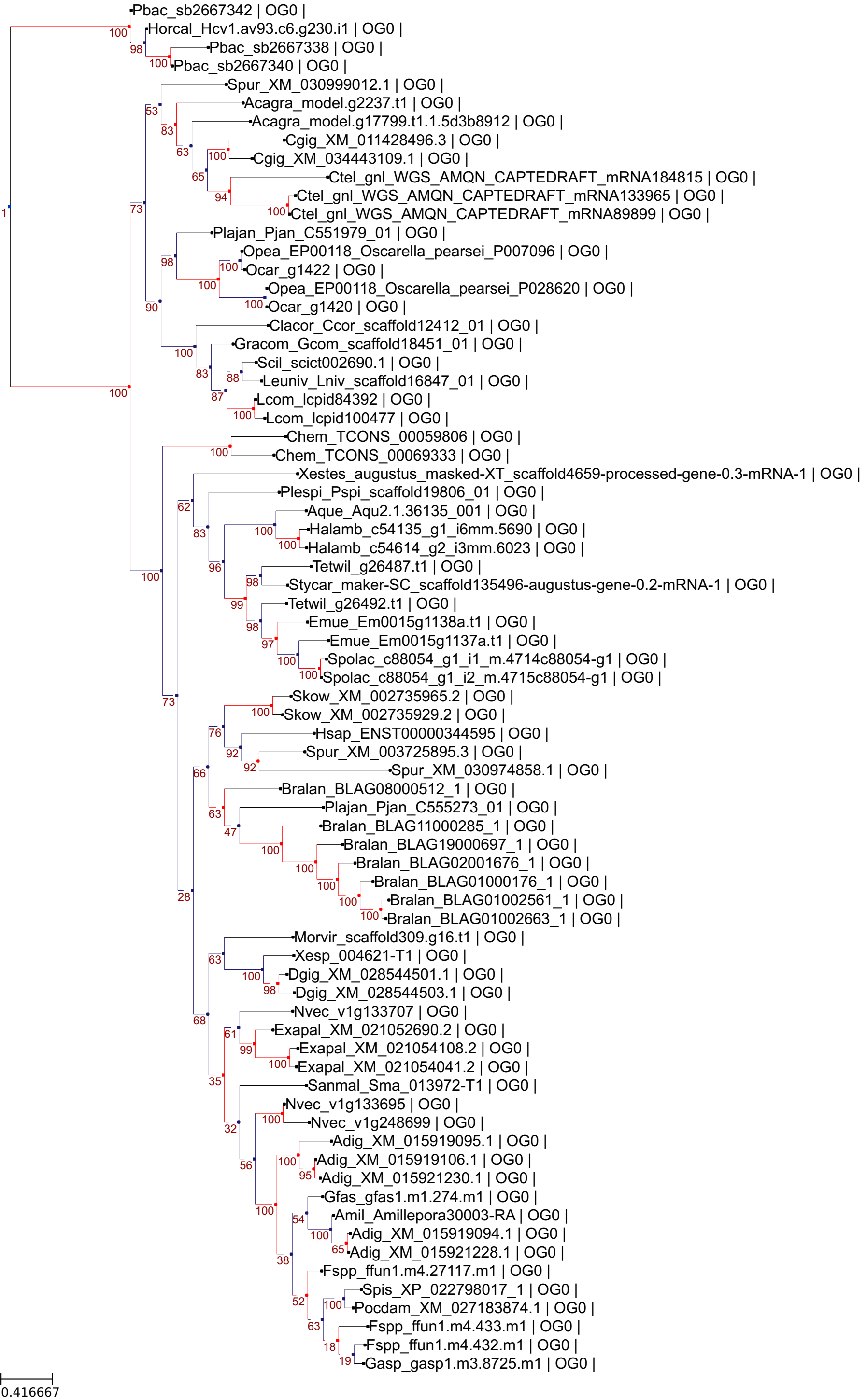
Pbac_sb2667342 | OG0 |
Horcal_Hcv1.av93.c6.g230.i1 | OG0 |
Pbac_sb2667338 | OG0 |
Pbac_sb2667340 | OG0 |
Spur_XM_030999012.1 | OG0 |
Acagra_model.g2237.t1 | OG0 |
Acagra_model.g17799.t1.1.5d3b8912 | OG0 |
Cgig_XM_011428496.3 | OG0 |
Cgig_XM_034443109.1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA184815 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA133965 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA89899 | OG0 |
Plajan_Pjan_C551979_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P007096 | OG0 |
Ocar_g1422 | OG0 |
Opea_EP00118_Oscarella_pearsei_P028620 | OG0 |
Ocar_g1420 | OG0 |
Clacor_Ccor_scaffold12412_01 | OG0 |
Gracom_Gcom_scaffold18451_01 | OG0 |
Scil_scict002690.1 | OG0 |
Leuniv_Lniv_scaffold16847_01 | OG0 |
Lcom_lcpid84392 | OG0 |
Lcom_lcpid100477 | OG0 |
Chem_TCONS_00059806 | OG0 |
Chem_TCONS_00069333 | OG0 |
Xestes_augustus_masked-XT_scaffold4659-processed-gene-0.3-mRNA-1 | OG0 |
Plespi_Pspi_scaffold19806_01 | OG0 |
Aque_Aqu2.1.36135_001 | OG0 |
Halamb_c54135_g1_i6mm.5690 | OG0 |
Halamb_c54614_g2_i3mm.6023 | OG0 |
Tetwil_g26487.t1 | OG0 |
Stycar_maker-SC_scaffold135496-augustus-gene-0.2-mRNA-1 | OG0 |
Tetwil_g26492.t1 | OG0 |
Emue_Em0015g1138a.t1 | OG0 |
Emue_Em0015g1137a.t1 | OG0 |
Spolac_c88054_g1_i1_m.4714c88054-g1 | OG0 |
Spolac_c88054_g1_i2_m.4715c88054-g1 | OG0 |
Skow_XM_002735965.2 | OG0 |
Skow_XM_002735929.2 | OG0 |
Hsap_ENST00000344595 | OG0 |
Spur_XM_003725895.3 | OG0 |
Spur_XM_030974858.1 | OG0 |
Bralan_BLAG08000512_1 | OG0 |
Plajan_Pjan_C555273_01 | OG0 |
Bralan_BLAG11000285_1 | OG0 |
Bralan_BLAG19000697_1 | OG0 |
Bralan_BLAG02001676_1 | OG0 |
Bralan_BLAG01000176_1 | OG0 |
Bralan_BLAG01002561_1 | OG0 |
Bralan_BLAG01002663_1 | OG0 |
Morvir_scaffold309.g16.t1 | OG0 |
Xesp_004621-T1 | OG0 |
Dgig_XM_028544501.1 | OG0 |
Dgig_XM_028544503.1 | OG0 |
Nvec_v1g133707 | OG0 |
Exapal_XM_021052690.2 | OG0 |
Exapal_XM_021054108.2 | OG0 |
Exapal_XM_021054041.2 | OG0 |
Sanmal_Sma_013972-T1 | OG0 |
Nvec_v1g133695 | OG0 |
Nvec_v1g248699 | OG0 |
Adig_XM_015919095.1 | OG0 |
Adig_XM_015919106.1 | OG0 |
Adig_XM_015921230.1 | OG0 |
Gfas_gfas1.m1.274.m1 | OG0 |
Amil_Amillepora30003-RA | OG0 |
Adig_XM_015919094.1 | OG0 |
Adig_XM_015921228.1 | OG0 |
Fspp_ffun1.m4.27117.m1 | OG0 |
Spis_XP_022798017_1 | OG0 |
Pocdam_XM_027183874.1 | OG0 |
Fspp_ffun1.m4.433.m1 | OG0 |
Fspp_ffun1.m4.432.m1 | OG0 |
Gasp_gasp1.m3.8725.m1 | OG0 |
0.416667