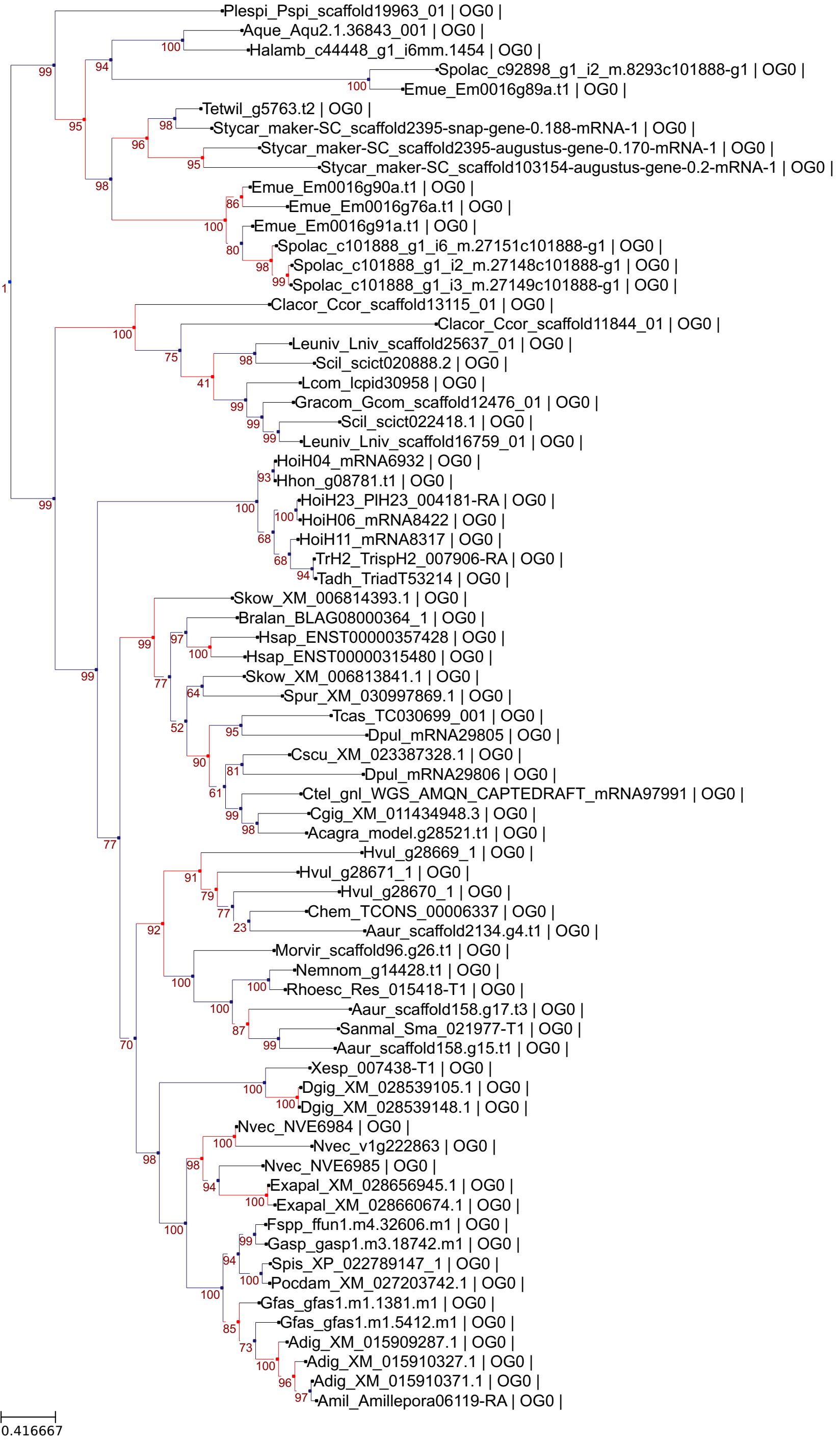
Plespi_Pspi_scaffold19963_01 | OG0 |
Aque_Aqu2.1.36843_001 | OG0 |
Halamb_c44448_g1_i6mm.1454 | OG0 |
Spolac_c92898_g1_i2_m.8293c101888-g1 | OG0 |
Emue_Em0016g89a.t1 | OG0 |
Tetwil_g5763.t2 | OG0 |
Stycar_maker-SC_scaffold2395-snap-gene-0.188-mRNA-1 | OG0 |
Stycar_maker-SC_scaffold2395-augustus-gene-0.170-mRNA-1 | OG0 |
Stycar_maker-SC_scaffold103154-augustus-gene-0.2-mRNA-1 | OG0 |
Emue_Em0016g90a.t1 | OG0 |
Emue_Em0016g76a.t1 | OG0 |
Emue_Em0016g91a.t1 | OG0 |
Spolac_c101888_g1_i6_m.27151c101888-g1 | OG0 |
Spolac_c101888_g1_i2_m.27148c101888-g1 | OG0 |
Spolac_c101888_g1_i3_m.27149c101888-g1 | OG0 |
Clacor_Ccor_scaffold13115_01 | OG0 |
Clacor_Ccor_scaffold11844_01 | OG0 |
Leuniv_Lniv_scaffold25637_01 | OG0 |
Scil_scict020888.2 | OG0 |
Lcom_lcpid30958 | OG0 |
Gracom_Gcom_scaffold12476_01 | OG0 |
Scil_scict022418.1 | OG0 |
Leuniv_Lniv_scaffold16759_01 | OG0 |
HoiH04_mRNA6932 | OG0 |
Hhon_g08781.t1 | OG0 |
HoiH23_PlH23_004181-RA | OG0 |
HoiH06_mRNA8422 | OG0 |
HoiH11_mRNA8317 | OG0 |
TrH2_TrispH2_007906-RA | OG0 |
Tadh_TriadT53214 | OG0 |
Skow_XM_006814393.1 | OG0 |
Bralan_BLAG08000364_1 | OG0 |
Hsap_ENST00000357428 | OG0 |
Hsap_ENST00000315480 | OG0 |
Skow_XM_006813841.1 | OG0 |
Spur_XM_030997869.1 | OG0 |
Tcas_TC030699_001 | OG0 |
Dpul_mRNA29805 | OG0 |
Cscu_XM_023387328.1 | OG0 |
Dpul_mRNA29806 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA97991 | OG0 |
Cgig_XM_011434948.3 | OG0 |
Acagra_model.g28521.t1 | OG0 |
Hvul_g28669_1 | OG0 |
Hvul_g28671_1 | OG0 |
Hvul_g28670_1 | OG0 |
Chem_TCONS_00006337 | OG0 |
Aaur_scaffold2134.g4.t1 | OG0 |
Morvir_scaffold96.g26.t1 | OG0 |
Nemnom_g14428.t1 | OG0 |
Rhoesc_Res_015418-T1 | OG0 |
Aaur_scaffold158.g17.t3 | OG0 |
Sanmal_Sma_021977-T1 | OG0 |
Aaur_scaffold158.g15.t1 | OG0 |
Xesp_007438-T1 | OG0 |
Dgig_XM_028539105.1 | OG0 |
Dgig_XM_028539148.1 | OG0 |
Nvec_NVE6984 | OG0 |
Nvec_v1g222863 | OG0 |
Nvec_NVE6985 | OG0 |
Exapal_XM_028656945.1 | OG0 |
Exapal_XM_028660674.1 | OG0 |
Fspp_ffun1.m4.32606.m1 | OG0 |
Gasp_gasp1.m3.18742.m1 | OG0 |
Spis_XP_022789147_1 | OG0 |
Pocdam_XM_027203742.1 | OG0 |
Gfas_gfas1.m1.1381.m1 | OG0 |
Gfas_gfas1.m1.5412.m1 | OG0 |
Adig_XM_015909287.1 | OG0 |
Adig_XM_015910327.1 | OG0 |
Adig_XM_015910371.1 | OG0 |
Amil_Amillepora06119-RA | OG0 |
0.416667