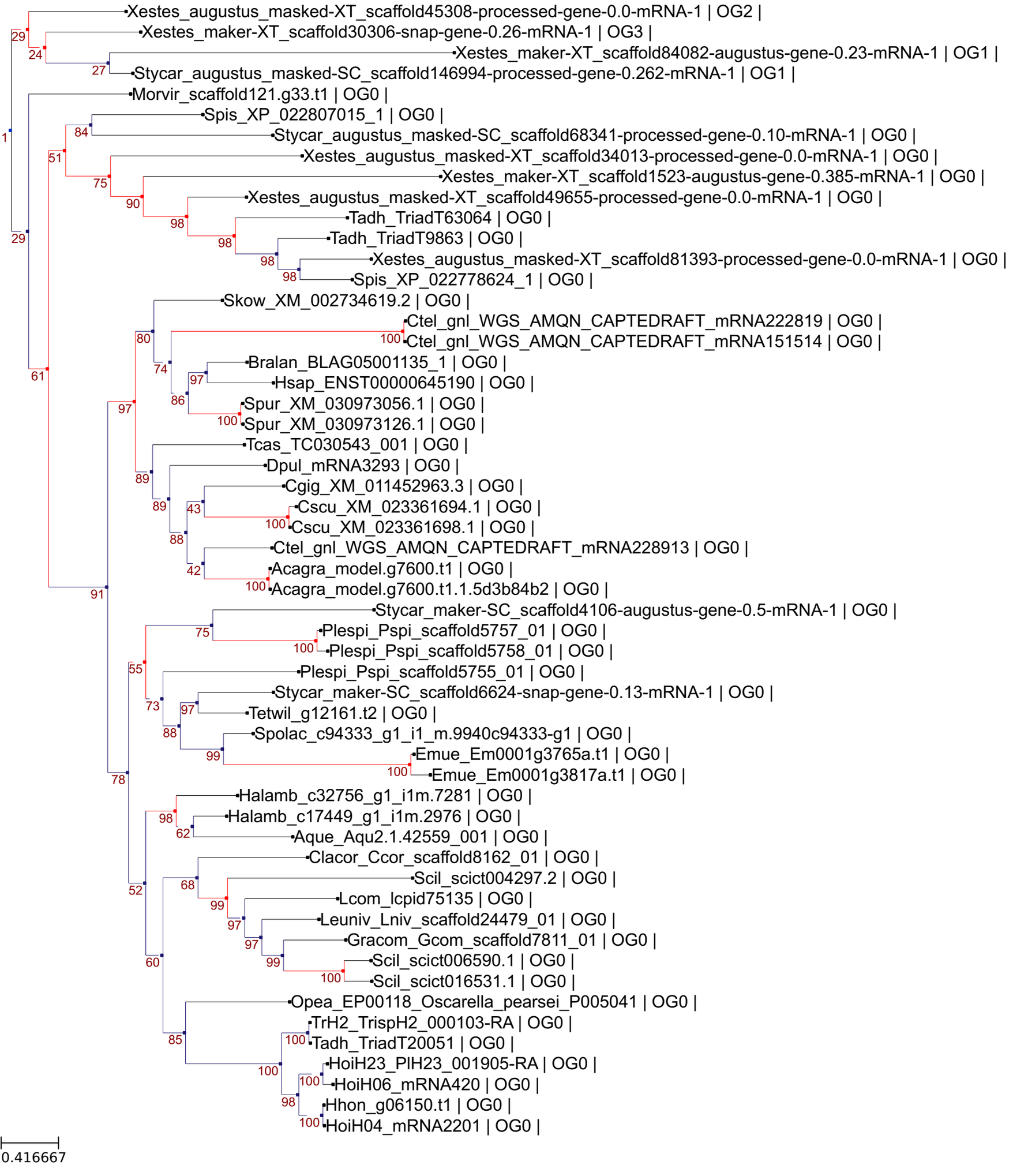
Xestes_augustus_masked-XT_scaffold45308-processed-gene-0.0-mRNA-1 | OG2 |
Xestes_maker-XT_scaffold30306-snap-gene-0.26-mRNA-1 | OG3 |
Xestes_maker-XT_scaffold84082-augustus-gene-0.23-mRNA-1 | OG1 |
Stycar_augustus_masked-SC_scaffold146994-processed-gene-0.262-mRNA-1 | OG1 |
Morvir_scaffold121.g33.t1 | OG0 |
Spis_XP_022807015_1 | OG0 |
Stycar_augustus_masked-SC_scaffold68341-processed-gene-0.10-mRNA-1 | OG0 |
Xestes_augustus_masked-XT_scaffold34013-processed-gene-0.0-mRNA-1 | OG0 |
Xestes_maker-XT_scaffold1523-augustus-gene-0.385-mRNA-1 | OG0 |
Xestes_augustus_masked-XT_scaffold49655-processed-gene-0.0-mRNA-1 | OG0 |
Tadh_TriadT63064 | OG0 |
Tadh_TriadT9863 | OG0 |
Xestes_augustus_masked-XT_scaffold81393-processed-gene-0.0-mRNA-1 | OG0 |
Spis_XP_022778624_1 | OG0 |
Skow_XM_002734619.2 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA222819 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA151514 | OG0 |
Bralan_BLAG05001135_1 | OG0 |
Hsap_ENST00000645190 | OG0 |
Spur_XM_030973056.1 | OG0 |
Spur_XM_030973126.1 | OG0 |
Tcas_TC030543_001 | OG0 |
Dpul_mRNA3293 | OG0 |
Cgig_XM_011452963.3 | OG0 |
Cscu_XM_023361694.1 | OG0 |
Cscu_XM_023361698.1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA228913 | OG0 |
Acagra_model.g7600.t1 | OG0 |
Acagra_model.g7600.t1.1.5d3b84b2 | OG0 |
Stycar_maker-SC_scaffold4106-augustus-gene-0.5-mRNA-1 | OG0 |
Plespi_Pspi_scaffold5757_01 | OG0 |
Plespi_Pspi_scaffold5758_01 | OG0 |
Plespi_Pspi_scaffold5755_01 | OG0 |
Stycar_maker-SC_scaffold6624-snap-gene-0.13-mRNA-1 | OG0 |
Tetwil_g12161.t2 | OG0 |
Spolac_c94333_g1_i1_m.9940c94333-g1 | OG0 |
Emue_Em0001g3765a.t1 | OG0 |
Emue_Em0001g3817a.t1 | OG0 |
Halamb_c32756_g1_i1m.7281 | OG0 |
Halamb_c17449_g1_i1m.2976 | OG0 |
Aque_Aqu2.1.42559_001 | OG0 |
Clacor_Ccor_scaffold8162_01 | OG0 |
Scil_scict004297.2 | OG0 |
Lcom_lcpid75135 | OG0 |
Leuniv_Lniv_scaffold24479_01 | OG0 |
Gracom_Gcom_scaffold7811_01 | OG0 |
Scil_scict006590.1 | OG0 |
Scil_scict016531.1 | OG0 |
Opea_EP00118_Oscarella_pearsei_P005041 | OG0 |
TrH2_TrispH2_000103-RA | OG0 |
Tadh_TriadT20051 | OG0 |
HoiH23_PlH23_001905-RA | OG0 |
HoiH06_mRNA420 | OG0 |
Hhon_g06150.t1 | OG0 |
HoiH04_mRNA2201 | OG0 |
0.416667