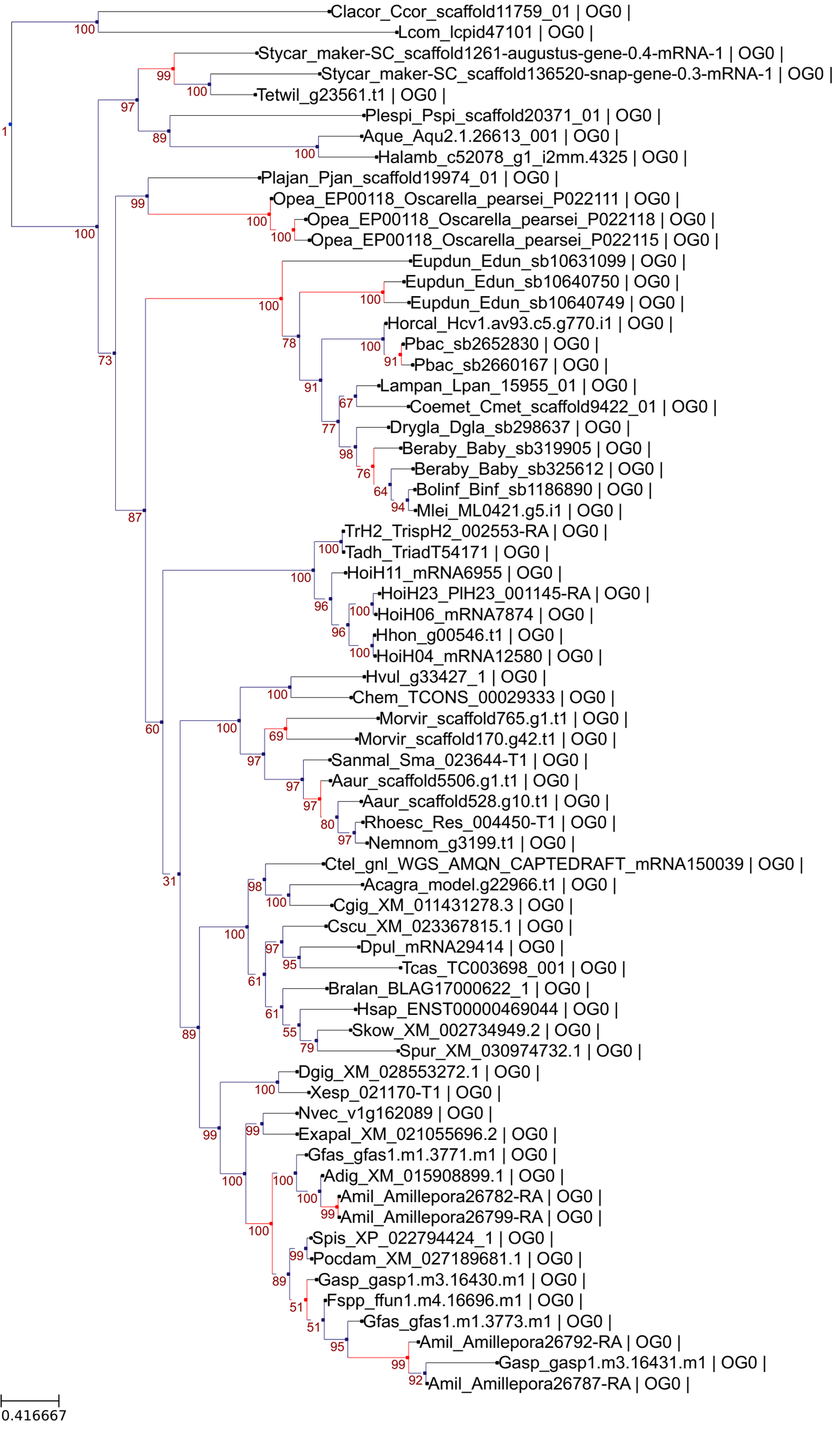
Clacor_Ccor_scaffold11759_01 | OG0 |
Lcom_lcpid47101 | OG0 |
Stycar_maker-SC_scaffold1261-augustus-gene-0.4-mRNA-1 | OG0 |
Stycar_maker-SC_scaffold136520-snap-gene-0.3-mRNA-1 | OG0 |
Tetwil_g23561.t1 | OG0 |
Plespi_Pspi_scaffold20371_01 | OG0 |
Aque_Aqu2.1.26613_001 | OG0 |
Halamb_c52078_g1_i2mm.4325 | OG0 |
Plajan_Pjan_scaffold19974_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P022111 | OG0 |
Opea_EP00118_Oscarella_pearsei_P022118 | OG0 |
Opea_EP00118_Oscarella_pearsei_P022115 | OG0 |
Eupdun_Edun_sb10631099 | OG0 |
Eupdun_Edun_sb10640750 | OG0 |
Eupdun_Edun_sb10640749 | OG0 |
Horcal_Hcv1.av93.c5.g770.i1 | OG0 |
Pbac_sb2652830 | OG0 |
Pbac_sb2660167 | OG0 |
Lampan_Lpan_15955_01 | OG0 |
Coemet_Cmet_scaffold9422_01 | OG0 |
Drygla_Dgla_sb298637 | OG0 |
Beraby_Baby_sb319905 | OG0 |
Beraby_Baby_sb325612 | OG0 |
Bolinf_Binf_sb1186890 | OG0 |
Mlei_ML0421.g5.i1 | OG0 |
TrH2_TrispH2_002553-RA | OG0 |
Tadh_TriadT54171 | OG0 |
HoiH11_mRNA6955 | OG0 |
HoiH23_PlH23_001145-RA | OG0 |
HoiH06_mRNA7874 | OG0 |
Hhon_g00546.t1 | OG0 |
HoiH04_mRNA12580 | OG0 |
Hvul_g33427_1 | OG0 |
Chem_TCONS_00029333 | OG0 |
Morvir_scaffold765.g1.t1 | OG0 |
Morvir_scaffold170.g42.t1 | OG0 |
Sanmal_Sma_023644-T1 | OG0 |
Aaur_scaffold5506.g1.t1 | OG0 |
Aaur_scaffold528.g10.t1 | OG0 |
Rhoesc_Res_004450-T1 | OG0 |
Nemnom_g3199.t1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA150039 | OG0 |
Acagra_model.g22966.t1 | OG0 |
Cgig_XM_011431278.3 | OG0 |
Cscu_XM_023367815.1 | OG0 |
Dpul_mRNA29414 | OG0 |
Tcas_TC003698_001 | OG0 |
Bralan_BLAG17000622_1 | OG0 |
Hsap_ENST00000469044 | OG0 |
Skow_XM_002734949.2 | OG0 |
Spur_XM_030974732.1 | OG0 |
Dgig_XM_028553272.1 | OG0 |
Xesp_021170-T1 | OG0 |
Nvec_v1g162089 | OG0 |
Exapal_XM_021055696.2 | OG0 |
Gfas_gfas1.m1.3771.m1 | OG0 |
Adig_XM_015908899.1 | OG0 |
Amil_Amillepora26782-RA | OG0 |
Amil_Amillepora26799-RA | OG0 |
Spis_XP_022794424_1 | OG0 |
Pocdam_XM_027189681.1 | OG0 |
Gasp_gasp1.m3.16430.m1 | OG0 |
Fspp_ffun1.m4.16696.m1 | OG0 |
Gfas_gfas1.m1.3773.m1 | OG0 |
Amil_Amillepora26792-RA | OG0 |
Gasp_gasp1.m3.16431.m1 | OG0 |
Amil_Amillepora26787-RA | OG0 |
0.416667