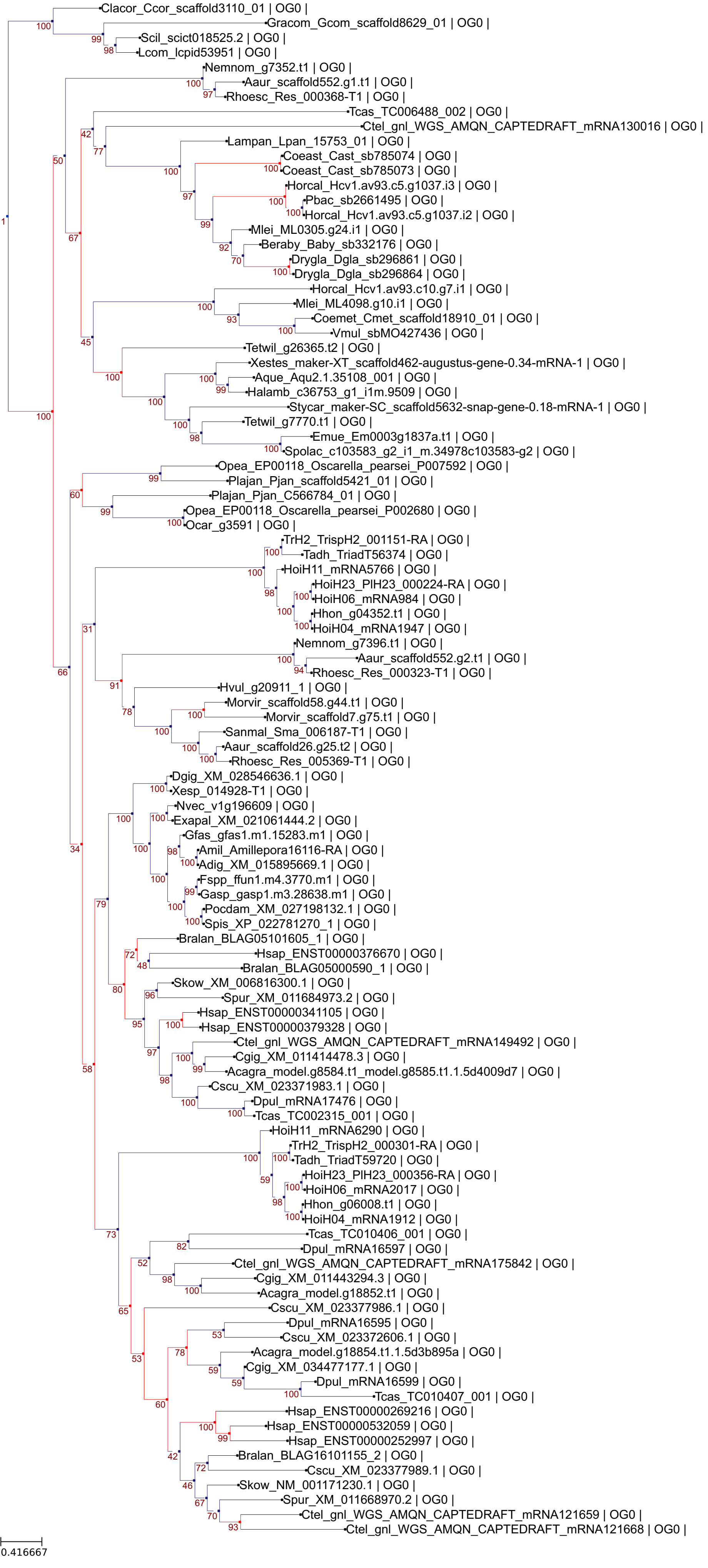
Clacor_Ccor_scaffold3110_01 | OG0 |
Gracom_Gcom_scaffold8629_01 | OG0 |
Scil_scict018525.2 | OG0 |
Lcom_lcpid53951 | OG0 |
Nemnom_g7352.t1 | OG0 |
Aaur_scaffold552.g1.t1 | OG0 |
Rhoesc_Res_000368-T1 | OG0 |
Tcas_TC006488_002 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA130016 | OG0 |
Lampan_Lpan_15753_01 | OG0 |
Coeast_Cast_sb785074 | OG0 |
Coeast_Cast_sb785073 | OG0 |
Horcal_Hcv1.av93.c5.g1037.i3 | OG0 |
Pbac_sb2661495 | OG0 |
Horcal_Hcv1.av93.c5.g1037.i2 | OG0 |
Mlei_ML0305.g24.i1 | OG0 |
Beraby_Baby_sb332176 | OG0 |
Drygla_Dgla_sb296861 | OG0 |
Drygla_Dgla_sb296864 | OG0 |
Horcal_Hcv1.av93.c10.g7.i1 | OG0 |
Mlei_ML4098.g10.i1 | OG0 |
Coemet_Cmet_scaffold18910_01 | OG0 |
Vmul_sbMO427436 | OG0 |
Tetwil_g26365.t2 | OG0 |
Xestes_maker-XT_scaffold462-augustus-gene-0.34-mRNA-1 | OG0 |
Aque_Aqu2.1.35108_001 | OG0 |
Halamb_c36753_g1_i1m.9509 | OG0 |
Stycar_maker-SC_scaffold5632-snap-gene-0.18-mRNA-1 | OG0 |
Tetwil_g7770.t1 | OG0 |
Emue_Em0003g1837a.t1 | OG0 |
Spolac_c103583_g2_i1_m.34978c103583-g2 | OG0 |
Opea_EP00118_Oscarella_pearsei_P007592 | OG0 |
Plajan_Pjan_scaffold5421_01 | OG0 |
Plajan_Pjan_C566784_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P002680 | OG0 |
Ocar_g3591 | OG0 |
TrH2_TrispH2_001151-RA | OG0 |
Tadh_TriadT56374 | OG0 |
HoiH11_mRNA5766 | OG0 |
HoiH23_PIH23_000224-RA | OG0 |
HoiH06_mRNA984 | OG0 |
Hhon_g04352.t1 | OG0 |
HoiH04_mRNA1947 | OG0 |
Nemnom_g7396.t1 | OG0 |
Aaur_scaffold552.g2.t1 | OG0 |
Rhoesc_Res_000323-T1 | OG0 |
Hvul_g20911_1 | OG0 |
Morvir_scaffold58.g44.t1 | OG0 |
Morvir_scaffold7.g75.t1 | OG0 |
Sanmal_Sma_006187-T1 | OG0 |
Aaur_scaffold26.g25.t2 | OG0 |
Rhoesc_Res_005369-T1 | OG0 |
Dgig_XM_028546636.1 | OG0 |
Xesp_014928-T1 | OG0 |
Nvec_v1g196609 | OG0 |
Exapal_XM_021061444.2 | OG0 |
Gfas_gfas1.m1.15283.m1 | OG0 |
Amil_Amillepora16116-RA | OG0 |
Adig_XM_015895669.1 | OG0 |
Fspp_ffun1.m4.3770.m1 | OG0 |
Gasp_gasp1.m3.28638.m1 | OG0 |
Pocdam_XM_027198132.1 | OG0 |
Spis_XP_022781270_1 | OG0 |
Bralan_BLAG05101605_1 | OG0 |
Hsap_ENST00000376670 | OG0 |
Bralan_BLAG05000590_1 | OG0 |
Skow_XM_006816300.1 | OG0 |
Spur_XM_011684973.2 | OG0 |
Hsap_ENST00000341105 | OG0 |
Hsap_ENST00000379328 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA149492 | OG0 |
Cgig_XM_011414478.3 | OG0 |
Acagra_model.g8584.t1_model.g8585.t1.1.5d4009d7 | OG0 |
Cscu_XM_023371983.1 | OG0 |
Dpul_mRNA17476 | OG0 |
Tcas_TC002315_001 | OG0 |
HoiH11_mRNA6290 | OG0 |
TrH2_TrispH2_000301-RA | OG0 |
Tadh_TriadT59720 | OG0 |
HoiH23_PIH23_000356-RA | OG0 |
HoiH06_mRNA2017 | OG0 |
Hhon_g06008.t1 | OG0 |
HoiH04_mRNA1912 | OG0 |
Tcas_TC010406_001 | OG0 |
Dpul_mRNA16597 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA175842 | OG0 |
Cgig_XM_011443294.3 | OG0 |
Acagra_model.g18852.t1 | OG0 |
Cscu_XM_023377986.1 | OG0 |
Dpul_mRNA16595 | OG0 |
Cscu_XM_023372606.1 | OG0 |
Acagra_model.g18854.t1.1.5d3b895a | OG0 |
Cgig_XM_034477177.1 | OG0 |
Dpul_mRNA16599 | OG0 |
Tcas_TC010407_001 | OG0 |
Hsap_ENST00000269216 | OG0 |
Hsap_ENST00000532059 | OG0 |
Hsap_ENST00000252997 | OG0 |
Bralan_BLAG16101155_2 | OG0 |
Cscu_XM_023377989.1 | OG0 |
Skow_NM_001171230.1 | OG0 |
Spur_XM_011668970.2 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA121659 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA121668 | OG0 |
0.416667