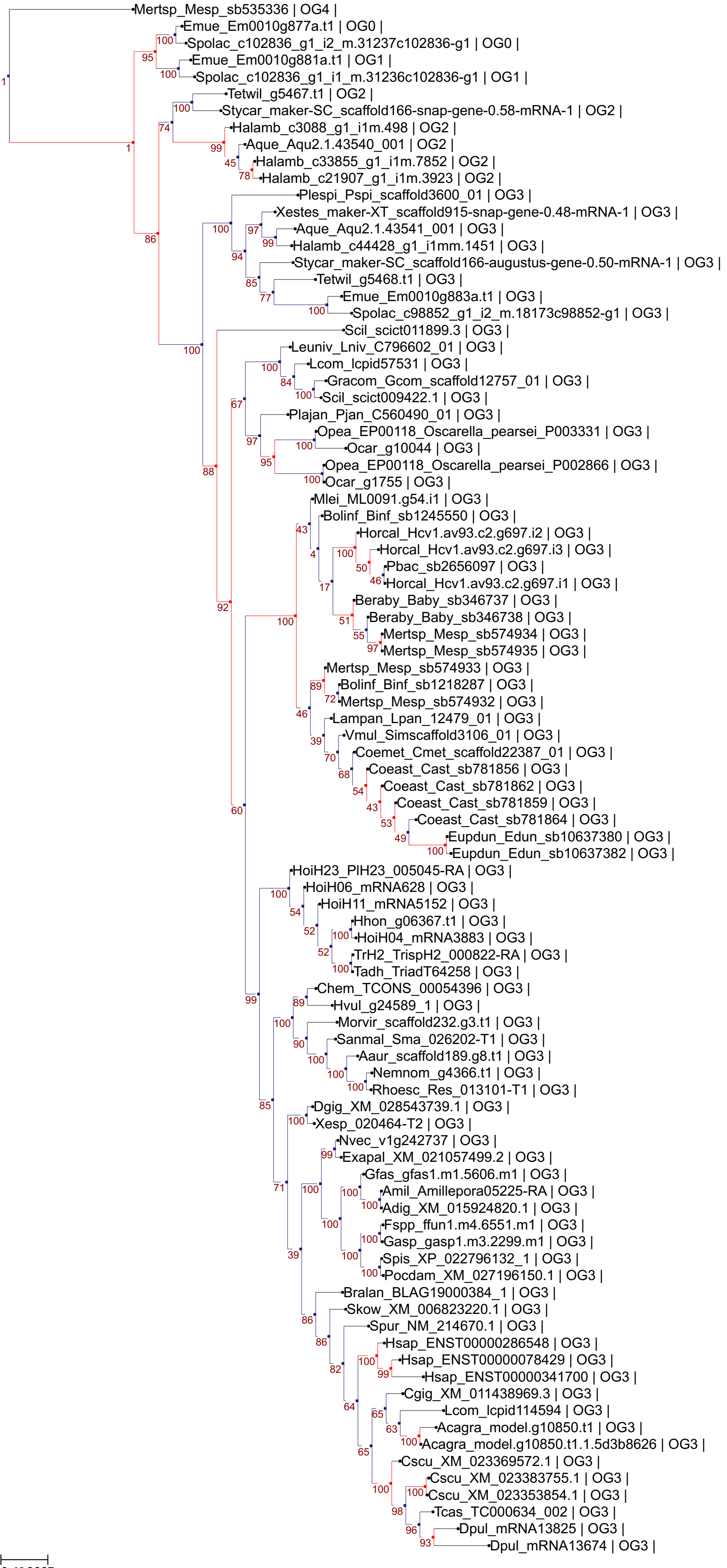
Mertsp_Mesp_sb535336 | OG4 |
Emue_Em0010g877a.t1 | OG0 |
Spolac_c102836_g1_i2_m.31237c102836-g1 | OG0 |
Emue_Em0010g881a.t1 | OG1 |
Spolac_c102836_g1_i1_m.31236c102836-g1 | OG1 |
Tetwil_g5467.t1 | OG2 |
Stycar_maker-SC_scaffold166-snap-gene-0.58-mRNA-1 | OG2 |
Halamb_c3088_g1_i1m.498 | OG2 |
Aque_Aqu2.1.43540_001 | OG2 |
Halamb_c33855_g1_i1m.7852 | OG2 |
Halamb_c21907_g1_i1m.3923 | OG2 |
Plespi_Pspi_scaffold3600_01 | OG3 |
Xestes_maker-XT_scaffold915-snap-gene-0.48-mRNA-1 | OG3 |
Aque_Aqu2.1.43541_001 | OG3 |
Halamb_c44428_g1_i1mm.1451 | OG3 |
Stycar_maker-SC_scaffold166-augustus-gene-0.50-mRNA-1 | OG3 |
Tetwil_g5468.t1 | OG3 |
Emue_Em0010g883a.t1 | OG3 |
Spolac_c98852_g1_i2_m.18173c98852-g1 | OG3 |
Scil_scict011899.3 | OG3 |
Leuniv_Lniv_C796602_01 | OG3 |
Lcom_lcpid57531 | OG3 |
Gracom_Gcom_scaffold12757_01 | OG3 |
Scil_scict009422.1 | OG3 |
Plajan_Pjan_C560490_01 | OG3 |
Opea_EP00118_Oscarella_pearsei_P003331 | OG3 |
Ocar_g10044 | OG3 |
Opea_EP00118_Oscarella_pearsei_P002866 | OG3 |
Ocar_g1755 | OG3 |
Mlei_ML0091.g54.i1 | OG3 |
Bolinf_Binf_sb1245550 | OG3 |
Horcal_Hcv1.av93.c2.g697.i2 | OG3 |
Horcal_Hcv1.av93.c2.g697.i3 | OG3 |
Pbac_sb2656097 | OG3 |
Horcal_Hcv1.av93.c2.g697.i1 | OG3 |
Beraby_Baby_sb346737 | OG3 |
Beraby_Baby_sb346738 | OG3 |
Mertsp_Mesp_sb574934 | OG3 |
Mertsp_Mesp_sb574935 | OG3 |
Mertsp_Mesp_sb574933 | OG3 |
Bolinf_Binf_sb1218287 | OG3 |
Mertsp_Mesp_sb574932 | OG3 |
Lampan_Lpan_12479_01 | OG3 |
Vmul_Simscaffold3106_01 | OG3 |
Coemet_Cmet_scaffold22387_01 | OG3 |
Coeast_Cast_sb781856 | OG3 |
Coeast_Cast_sb781862 | OG3 |
Coeast_Cast_sb781859 | OG3 |
Coeast_Cast_sb781864 | OG3 |
Eupdun_Edun_sb10637380 | OG3 |
Eupdun_Edun_sb10637382 | OG3 |
HoiH23_PIH23_005045-RA | OG3 |
HoiH06_mRNA628 | OG3 |
HoiH11_mRNA5152 | OG3 |
Hhon_g06367.t1 | OG3 |
HoiH04_mRNA3883 | OG3 |
TrH2_TrispH2_000822-RA | OG3 |
Tadh_TriadT64258 | OG3 |
Chem_TCONS_00054396 | OG3 |
Hvul_g24589_1 | OG3 |
Morvir_scaffold232.g3.t1 | OG3 |
Sanmal_Sma_026202-T1 | OG3 |
Aaur_scaffold189.g8.t1 | OG3 |
Nemnom_g4366.t1 | OG3 |
Rhoesc_Res_013101-T1 | OG3 |
Dgig_XM_028543739.1 | OG3 |
Xesp_020464-T2 | OG3 |
Nvec_v1g242737 | OG3 |
Exapal_XM_021057499.2 | OG3 |
Gfas_gfas1.m1.5606.m1 | OG3 |
Amil_Amillepora05225-RA | OG3 |
Adig_XM_015924820.1 | OG3 |
Fspp_ffun1.m4.6551.m1 | OG3 |
Gasp_gasp1.m3.2299.m1 | OG3 |
Spis_XP_022796132_1 | OG3 |
Pocdam_XM_027196150.1 | OG3 |
Bralan_BLAG19000384_1 | OG3 |
Skow_XM_006823220.1 | OG3 |
Spur_NM_214670.1 | OG3 |
Hsap_ENST00000286548 | OG3 |
Hsap_ENST00000078429 | OG3 |
Hsap_ENST00000341700 | OG3 |
Cgig_XM_011438969.3 | OG3 |
Lcom_lcpid114594 | OG3 |
Acagra_model.g10850.t1 | OG3 |
Acagra_model.g10850.t1.1.5d3b8626 | OG3 |
Cscu_XM_023369572.1 | OG3 |
Cscu_XM_023383755.1 | OG3 |
Cscu_XM_023353854.1 | OG3 |
Tcas_TC000634_002 | OG3 |
Dpul_mRNA13825 | OG3 |
Dpul_mRNA13674 | OG3 |
0.416667