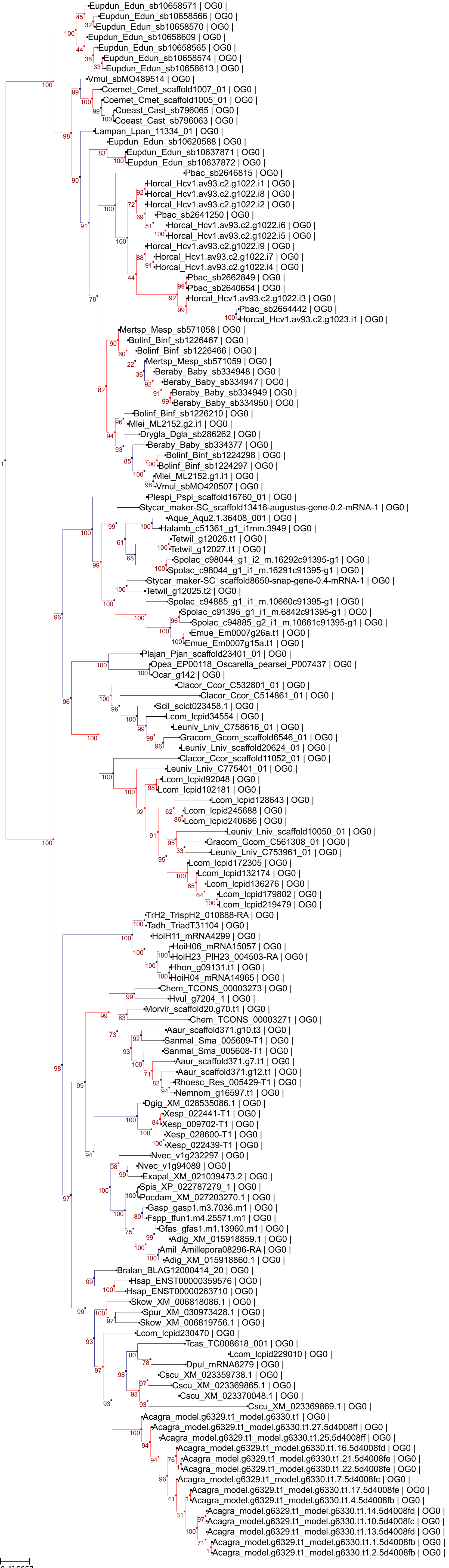
Eupdun_Edun_sb10658571 | OG0 |
Eupdun_Edun_sb10658566 | OG0 |
Eupdun_Edun_sb10658570 | OG0 |
Eupdun_Edun_sb10658609 | OG0 |
Eupdun_Edun_sb10658565 | OG0 |
Eupdun_Edun_sb10658574 | OG0 |
Eupdun_Edun_sb10658613 | OG0 |
Vmul_sbMO489514 | OG0 |
Coemet_Cmet_scaffold1007_01 | OG0 |
Coemet_Cmet_scaffold1005_01 | OG0 |
Coeast_Cast_sb796065 | OG0 |
Coeast_Cast_sb796063 | OG0 |
Lampan_Lpan_11334_01 | OG0 |
Eupdun_Edun_sb10620588 | OG0 |
Eupdun_Edun_sb10637871 | OG0 |
Eupdun_Edun_sb10637872 | OG0 |
Pbac_sb2646815 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i1 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i8 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i2 | OG0 |
Pbac_sb2641250 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i6 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i5 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i9 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i7 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i4 | OG0 |
Pbac_sb2662849 | OG0 |
Pbac_sb2640654 | OG0 |
Horcal_Hcv1.av93.c2.g1022.i3 | OG0 |
Pbac_sb2654442 | OG0 |
Horcal_Hcv1.av93.c2.g1023.i1 | OG0 |
Mertsp_Mesp_sb571058 | OG0 |
Bolinf_Binf_sb1226467 | OG0 |
Bolinf_Binf_sb1226466 | OG0 |
Mertsp_Mesp_sb571059 | OG0 |
Beraby_Baby_sb334948 | OG0 |
Beraby_Baby_sb334947 | OG0 |
Beraby_Baby_sb334949 | OG0 |
Beraby_Baby_sb334950 | OG0 |
Bolinf_Binf_sb1226210 | OG0 |
Mlei_ML2152.g2.i1 | OG0 |
Drygla_Dgla_sb286262 | OG0 |
Beraby_Baby_sb334377 | OG0 |
Bolinf_Binf_sb1224298 | OG0 |
Bolinf_Binf_sb1224297 | OG0 |
Mlei_ML2152.g1.i1 | OG0 |
Vmul_sbMO420507 | OG0 |
Plespi_Pspi_scaffold16760_01 | OG0 |
Stycar_maker-SC_scaffold13416-augustus-gene-0.2-mRNA-1 | OG0 |
Aque_Aqu2.1.36408_001 | OG0 |
Halamb_c51361_g1_i1mm.3949 | OG0 |
Tetwil_g12026.t1 | OG0 |
Tetwil_g12027.t1 | OG0 |
Spolac_c98044_g1_i2_m.16292c91395-g1 | OG0 |
Spolac_c98044_g1_i1_m.16291c91395-g1 | OG0 |
Stycar_maker-SC_scaffold8650-snap-gene-0.4-mRNA-1 | OG0 |
Tetwil_g12025.t2 | OG0 |
Spolac_c94885_g1_i1_m.10660c91395-g1 | OG0 |
Spolac_c91395_g1_i1_m.6842c91395-g1 | OG0 |
Spolac_c94885_g2_i1_m.10661c91395-g1 | OG0 |
Emue_Em0007g26a.t1 | OG0 |
Emue_Em0007g15a.t1 | OG0 |
Plajan_Pjan_scaffold23401_01 | OG0 |
Opea_EP00118_Oscarella_pearsei_P007437 | OG0 |
Ocar_g142 | OG0 |
Clacor_Ccor_C532801_01 | OG0 |
Clacor_Ccor_C514861_01 | OG0 |
Scil_scict023458.1 | OG0 |
Lcom_lcpid34554 | OG0 |
Leuniv_Lniv_C758616_01 | OG0 |
Gracom_Gcom_scaffold6546_01 | OG0 |
Leuniv_Lniv_scaffold20624_01 | OG0 |
Clacor_Ccor_scaffold11052_01 | OG0 |
Leuniv_Lniv_C775401_01 | OG0 |
Lcom_lcpid92048 | OG0 |
Lcom_lcpid102181 | OG0 |
Lcom_lcpid128643 | OG0 |
Lcom_lcpid245688 | OG0 |
Lcom_lcpid240686 | OG0 |
Leuniv_Lniv_scaffold10050_01 | OG0 |
Gracom_Gcom_C561308_01 | OG0 |
Leuniv_Lniv_C753961_01 | OG0 |
Lcom_lcpid172305 | OG0 |
Lcom_lcpid132174 | OG0 |
Lcom_lcpid136276 | OG0 |
Lcom_lcpid179802 | OG0 |
Lcom_lcpid219479 | OG0 |
TrH2_TrispH2_010888-RA | OG0 |
Tadh_TriadT31104 | OG0 |
HoiH11_mRNA4299 | OG0 |
HoiH06_mRNA15057 | OG0 |
HoiH23_PlH23_004503-RA | OG0 |
Hhon_g09131.t1 | OG0 |
HoiH04_mRNA14965 | OG0 |
Chem_TCONS_00003273 | OG0 |
Hvul_g7204_1 | OG0 |
Morvir_scaffold20.g70.t1 | OG0 |
Chem_TCONS_00003271 | OG0 |
Aaur_scaffold371.g10.t3 | OG0 |
Sanmal_Sma_005609-T1 | OG0 |
Sanmal_Sma_005608-T1 | OG0 |
Aaur_scaffold371.g7.t1 | OG0 |
Aaur_scaffold371.g12.t1 | OG0 |
Rhoesc_Res_005429-T1 | OG0 |
Nemnom_g16597.t1 | OG0 |
Dgig_XM_028535086.1 | OG0 |
Xesp_022441-T1 | OG0 |
Xesp_009702-T1 | OG0 |
Xesp_028600-T1 | OG0 |
Xesp_022439-T1 | OG0 |
Nvec_v1g232297 | OG0 |
Nvec_v1g94089 | OG0 |
Exapal_XM_021039473.2 | OG0 |
Spis_XP_022787279_1 | OG0 |
Pocdam_XM_027203270.1 | OG0 |
Gasp_gasp1.m3.7036.m1 | OG0 |
Fspp_ffun1.m4.25571.m1 | OG0 |
Gfas_gfas1.m1.13960.m1 | OG0 |
Adig_XM_015918859.1 | OG0 |
Amil_Amillepora08296-RA | OG0 |
Adig_XM_015918860.1 | OG0 |
Bralan_BLAG12000414_20 | OG0 |
Hsap_ENST00000359576 | OG0 |
Hsap_ENST00000263710 | OG0 |
Skow_XM_006818086.1 | OG0 |
Spur_XM_030973428.1 | OG0 |
Skow_XM_006819756.1 | OG0 |
Lcom_lcpid230470 | OG0 |
Tcas_TC008618_001 | OG0 |
Lcom_lcpid229010 | OG0 |
Dpul_mRNA6279 | OG0 |
Cscu_XM_023359738.1 | OG0 |
Cscu_XM_023369865.1 | OG0 |
Cscu_XM_023370048.1 | OG0 |
Cscu_XM_023369869.1 | OG0 |
Acagra_model.g6329.t1_model.g6330.t1 | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.27.5d4008ff | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.25.5d4008ff | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.16.5d4008fd | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.21.5d4008fe | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.22.5d4008fe | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.7.5d4008fc | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.17.5d4008fe | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.4.5d4008fb | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.14.5d4008fd | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.10.5d4008fc | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.13.5d4008fd | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.1.5d4008fb | OG0 |
Acagra_model.g6329.t1_model.g6330.t1.2.5d4008fb | OG0 |
0.416667