Sanmal_Sma_016023-T1 | OG0 |
Sanmal_Sma_008845-T1 | OG0 |
Tetwil_g14196.t1 | OG1 |
Emue_Em0015g563a.t1 | OG1 |
Spolac_c101463_g3_i1_m.25690c101463-g3 | OG1 |
Stycar_maker-SC_scaffold11503-augustus-gene-0.2-mRNA-1 | OG2 |
Plespi_Pspi_scaffold13444_01 | OG2 |
Spolac_c98720_g1_i1_m.17882c101392-g1 | OG2 |
Emue_Em0001g227a.t1 | OG2 |
Emue_Em0001g210a.t1 | OG2 |
Halamb_c28143_g1_i2m.5678 | OG2 |
Aque_Aqu2.1.29899_001 | OG2 |
Xestes_maker-XT_scaffold116987-augustus-gene-0.5-mRNA-1 | OG2 |
Lcom_lcpid174081 | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA164238 | OG2 |
Lcom_lcpid138868 | OG2 |
Acagra_model.g15101.t1 | OG2 |
Lcom_lcpid84182 | OG2 |
Cgig_XM_034482855.1 | OG2 |
Cscu_XM_023369778.1 | OG2 |
Tcas_TC006047_001 | OG2 |
Dpul_mRNA15190 | OG2 |
Dpul_mRNA24920 | OG2 |
Skow_XM_002731289.2 | OG2 |
Bralan_BLAG01001388_1 | OG2 |
Hsap_ENST00000313486 | OG2 |
Hsap_ENST00000634017 | OG2 |
HoiH23_PlH23_008639-RA | OG2 |
HoiH06_mRNA5851 | OG2 |
Hhon_g00762.t1 | OG2 |
HoiH04_mRNA11599 | OG2 |
HoiH11_mRNA6584 | OG2 |
TrH2_TrispH2_002006-RA | OG2 |
Tadh_TriadT59214 | OG2 |
Spur_XM_030992072.1 | OG2 |
Hvul_g21921_1 | OG2 |
Chem_TCONS_00003235 | OG2 |
Chem_TCONS_00044952 | OG2 |
Aaur_scaffold658.g15.t1 | OG2 |
Sanmal_Sma_022148-T1 | OG2 |
Sanmal_Sma_011576-T1 | OG2 |
Rhoesc_Res_002279-T1 | OG2 |
Nemnom_g18752.t1 | OG2 |
Rhoesc_Res_015210-T1 | OG2 |
Aaur_scaffold658.g14.t1 | OG2 |
Morvir_scaffold44.g21.t1 | OG2 |
Dgig_XM_028551874.1 | OG2 |
Xesp_008333-T1 | OG2 |
Spis_XP_022786411_1 | OG2 |
Adig_XM_015910577.1 | OG2 |
Fspp_ffun1.m4.17743.m1 | OG2 |
Pocdam_XM_027198769.1 | OG2 |
Gasp_gasp1.m3.28494.m1 | OG2 |
Gfas_gfas1.m1.14587.m1 | OG2 |
Exapal_XM_021059124.2 | OG2 |
Nvec_v1g183595 | OG2 |
Nvec_v1g204254 | OG2 |
Spur_XM_030981380.1 | OG2 |
Spur_XM_030981884.1 | OG2 |
Spur_XM_030981886.1 | OG2 |
Spur_XM_030982631.1 | OG2 |
Mlei_ML0198.g11.i1 | OG2 |
Vmul_sbMO391066 | OG2 |
Vmul_sbMO484322 | OG2 |
Coeast_Cast_sb769444 | OG2 |
Coemet_Cmet_scaffold10756_01 | OG2 |
Clacor_Ccor_C538331_01 | OG2 |
Lcom_lcpid89415 | OG2 |
Scil_scict001571.1 | OG2 |
Gracom_Gcom_scaffold7474_01 | OG2 |
Leuniv_Lniv_scaffold22375_01 | OG2 |
Leuniv_Lniv_scaffold19498_01 | OG2 |
Lcom_lcpid94929 | OG2 |
Gracom_Gcom_C540602_01 | OG2 |
Gracom_Gcom_C515984_01 | OG2 |
Salmac_EP00052_Salpingoeca_macrocollata_P015082 | OG2 |
Salmac_EP00052_Salpingoeca_macrocollata_P015084 | OG2 |
Salurc_EP00051_Salpingoeca_urceolata_P006015 | OG2 |
Saldol_EP00054_Salpingoeca_dolichothecata_P031540 | OG2 |
Salmac_EP00052_Salpingoeca_macrocollata_P015083 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P022122 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P023929 | OG2 |
Vmul_sbMO424370 | OG2 |
Vmul_Simscaffold6589_01 | OG2 |
Coemet_Cmet_C597491_01 | OG2 |
Coeast_Cast_sb777317 | OG2 |
Coeast_Cast_sb777316 | OG2 |
Coeast_Cast_sb777314 | OG2 |
Coeast_Cast_sb777315 | OG2 |
Coeast_Cast_sb773549 | OG2 |
Horcal_Hcv1.av93.c2.g1052.i1 | OG2 |
Pbac_sb2654486 | OG2 |
Lampan_Lpan_C410555_01 | OG2 |
Mertsp_Mesp_sb567415 | OG2 |
Bolinf_Binf_sb1224228 | OG2 |
Bolinf_Binf_sb1224227 | OG2 |
Bolinf_Binf_sb1224229 | OG2 |
Eupdun_Edun_sb10679340 | OG2 |
Drygla_Dgla_sb287422 | OG2 |
Beraby_Baby_sb338449 | OG2 |
Mertsp_Mesp_sb527676 | OG2 |
Bolinf_Binf_sb1230534 | OG2 |
Bolinf_Binf_sb1230535 | OG2 |
Mlei_ML1316.g5.i1 | OG2 |
Vmul_sbMO427584 | OG2 |
Bolinf_Binf_sb1252962 | OG2 |
Bolinf_Binf_sb1252961 | OG2 |
Bolinf_Binf_sb1252960 | OG2 |
Mertsp_Mesp_sb576550 | OG2 |
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