Lampan_Lpan_C416703_01 | OG0 |
Vmul_SimC400406 | OG0 |
Coeast_Cast_sb816925 | OG0 |
Coemet_Cmet_scaffold13450_01 | OG0 |
Lampan_Lpan_15730_01 | OG1 |
Coeast_Cast_sb760354 | OG1 |
Vmul_sbMO445622 | OG1 |
Horcal_Hcv1.av93.c8.g316.i1 | OG1 |
Pbac_sb2662199 | OG1 |
Drygla_Dgla_sb278729 | OG1 |
Beraby_Baby_sb353218 | OG1 |
Mlei_ML0074.g7.i1 | OG1 |
Vmul_sbMO384689 | OG1 |
Bolinf_Binf_sb1274923 | OG1 |
Bolinf_Binf_sb1274920 | OG1 |
Bolinf_Binf_sb1274922 | OG1 |
Bolinf_Binf_sb1274919 | OG1 |
Bolinf_Binf_sb1274921 | OG1 |
Mertsp_Mesp_sb590625 | OG1 |
Mertsp_Mesp_sb590627 | OG1 |
Mertsp_Mesp_sb590628 | OG1 |
Mertsp_Mesp_sb590624 | OG1 |
Mertsp_Mesp_sb590626 | OG1 |
Pbac_sb2669405 | OG2 |
Horcal_Hcv1.av93.c8.g765.i1 | OG2 |
Pbac_sb2669404 | OG2 |
Beraby_Baby_sb353503 | OG2 |
Drygla_Dgla_sb311048 | OG2 |
Mlei_ML3530.g2.i1 | OG2 |
Beraby_Baby_sb323887 | OG2 |
Beraby_Baby_sb323888 | OG2 |
Eupdun_Edun_sb10603025 | OG3 |
Vmul_SimC404832 | OG3 |
Coemet_Cmet_scaffold20525_01 | OG3 |
Lampan_Lpan_13997_01 | OG3 |
Beraby_Baby_sb323699 | OG3 |
Drygla_Dgla_sb294291 | OG3 |
Beraby_Baby_sb328878 | OG3 |
Beraby_Baby_sb328879 | OG3 |
Pbac_sb2643575 | OG3 |
Horcal_Hcv1.av93.c4.g704.i5 | OG3 |
Horcal_Hcv1.av93.c4.g704.i1 | OG3 |
Horcal_Hcv1.av93.c4.g704.i3 | OG3 |
Horcal_Hcv1.av93.c4.g704.i4 | OG3 |
Horcal_Hcv1.av93.c4.g704.i2 | OG3 |
Vmul_sbMO410644 | OG3 |
Mlei_ML0280.g13.i1 | OG3 |
Bolinf_Binf_sb1260603 | OG3 |
Bolinf_Binf_sb1260600 | OG3 |
Bolinf_Binf_sb1260604 | OG3 |
Bolinf_Binf_sb1260602 | OG3 |
Bolinf_Binf_sb1260601 | OG3 |
Mertsp_Mesp_sb540160 | OG3 |
Clacor_Ccor_scaffold11002_01 | OG3 |
Lcom_lcpid23873 | OG3 |
Leuniv_Lniv_C748369_01 | OG3 |
Leuniv_Lniv_C799810_01 | OG3 |
Scil_scict003282.2 | OG3 |
Leuniv_Lniv_C758544_01 | OG3 |
Gracom_Gcom_C558970_01 | OG3 |
Salurc_EP00051_Salpingoeca_urceolata_P000966 | OG3 |
Salurc_EP00051_Salpingoeca_urceolata_P000967 | OG3 |
Didcos_EP00039_Didymoeca_costata_P012069 | OG3 |
Stedip_EP00041_Stephanoeca_diplocostata_P013746 | OG3 |
Diagra_EP00040_Diaphanoeca_grandis_P030756 | OG3 |
Acaspe_EP00035_Acanthoeca_spectabilis_P033878 | OG3 |
Helnan_EP00037_Helgoeca_nana_P002648 | OG3 |
Salpun_EP00053_Salpingoeca_punica_P015095 | OG3 |
Salmac_EP00052_Salpingoeca_macrocollata_P019474 | OG3 |
Mylflu_EP00047_Mylnosiga_fluctuans_P008870 | OG3 |
Saldol_EP00054_Salpingoeca_dolichothecata_P027674 | OG3 |
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Salkve_EP00050_Salpingoeca_kvevrii_P022204 | OG3 |
Codhol_EP00042_Codosiga_hollandica_P034269 | OG3 |
Codhol_EP00042_Codosiga_hollandica_P034275 | OG3 |
Codhol_EP00042_Codosiga_hollandica_P034265 | OG3 |
Codhol_EP00042_Codosiga_hollandica_P034277 | OG3 |
Salinf_EP00049_Salpingoeca_infusionum_P009143 | OG3 |
Sros_EGD77068 | OG3 |
Sros_EGD72840 | OG3 |
Microa_EP00043_Microstomoeca_roanoka_P009662 | OG3 |
Mbre_EDQ87111 | OG3 |
Chofle_EP00730_Choanoeca_flexa_P001692 | OG3 |
Choper_EP00045_Choanoeca_perplexa_P010733 | OG3 |
Xestes_augustus_masked-XT_scaffold688-processed-gene-0.6-mRNA-1 | OG3 |
Halamb_c19999_g1_i1m.3532 | OG3 |
Aque_Aqu2.1.24265_001 | OG3 |
Halamb_c44242_g2_i1m.17231 | OG3 |
Emue_Em0003g290a.t1 | OG3 |
Spolac_c42340_g1_i1_m.1197c42340-g1 | OG3 |
Tetwil_g25652.t1 | OG3 |
Stycar_augustus_masked-SC_scaffold3398-processed-gene-0.1-mRNA-1 | OG3 |
Stycar_maker-SC_scaffold3398-snap-gene-0.15-mRNA-1 | OG3 |
Plajan_Pjan_C554513_01 | OG3 |
Plajan_Pjan_C546751_01 | OG3 |
Opea_EP00118_Oscarella_pearsei_P004872 | OG3 |
Ocar_g10016 | OG3 |
Opea_EP00118_Oscarella_pearsei_P004871 | OG3 |
TrH2_TrispH2_008139-RA | OG3 |
Tadh_TriadT51263 | OG3 |
HoiH23_PlH23_008780-RA | OG3 |
HoiH06_mRNA11850 | OG3 |
Hhon_g00519.t1 | OG3 |
HoiH04_mRNA16981 | OG3 |
Spur_XM_030981943.1 | OG3 |
Skow_XM_006821225.1 | OG3 |
Spur_XM_785688.5 | OG3 |
Bralan_BLAG14000234_1 | OG3 |
Hsap_ENST00000382172 | OG3 |
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Acagra_model.g21937.t1 | OG3 |
Cgig_XM_034473854.1 | OG3 |
Tcas_TC006380_001 | OG3 |
Cscu_XM_023372354.1 | OG3 |
Dpul_mRNA22481 | OG3 |
Dpul_mRNA22460 | OG3 |
Hvul_g22726_1 | OG3 |
Chem_TCONS_00049978 | OG3 |
Morvir_scaffold42.g27.t1 | OG3 |
Sanmal_Sma_015219-T1 | OG3 |
Aaur_scaffold14.g8.t1 | OG3 |
Nemnom_g17507.t1 | OG3 |
Rhoesc_Res_007224-T1 | OG3 |
Dgig_XM_028540189.1 | OG3 |
Xesp_011971-T1 | OG3 |
Exapal_XM_021046183.2 | OG3 |
Nvec_v1g99065 | OG3 |
Gfas_gfas1.m1.4269.m1 | OG3 |
Amil_Amillepora16215-RA | OG3 |
Adig_XM_015895968.1 | OG3 |
Spis_XP_022777992_1 | OG3 |
Pocdam_XM_027204601.1 | OG3 |
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