Spur_XM_030979668.1 | OG0 |
Spur_XM_030986980.1 | OG0 |
Mbre_EDQ86239 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P003150 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P012861 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P000165 | OG1 |
Sros_EGD73956 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P026575 | OG2 |
Saldol_EP00054_Salpingoeca_dolichothecata_P026576 | OG2 |
Salurc_EP00051_Salpingoeca_urceolata_P034957 | OG2 |
Salpun_EP00053_Salpingoeca_punica_P017215 | OG2 |
Mylflu_EP00047_Mylnosiga_fluctuans_P016013 | OG2 |
Codhol_EP00042_Codosiga_hollandica_P016206 | OG2 |
Salpun_EP00053_Salpingoeca_punica_P017216 | OG2 |
Salmac_EP00052_Salpingoeca_macrocollata_P033004 | OG2 |
Cscu_XM_023368864.1 | OG2 |
Tcas_TC014120_001 | OG2 |
Chofle_EP00730_Choanoeca_flexa_P018929 | OG2 |
Choper_EP00045_Choanoeca_perplexa_P004544 | OG2 |
Didcos_EP00039_Didymoeca_costata_P007963 | OG2 |
Stedip_EP00041_Stephanoeca_diplocostata_P018888 | OG2 |
Diagra_EP00040_Diaphanoeca_grandis_P010630 | OG2 |
Acaspe_EP00035_Acanthoeca_spectabilis_P030554 | OG2 |
Helnan_EP00037_Helgoeca_nana_P016644 | OG2 |
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Clacor_Ccor_scaffold5915_01 | OG2 |
Lcom_lcpid82829 | OG2 |
Leuniv_Lniv_scaffold25436_01 | OG2 |
Gracom_Gcom_C552170_01 | OG2 |
Scil_scict000267.1 | OG2 |
TrH2_TrispH2_005177-RA | OG2 |
Tadh_TriadT19869 | OG2 |
HoiH11_mRNA8786 | OG2 |
HoiH23_PIH23_001201-RA | OG2 |
HoiH06_mRNA36 | OG2 |
Hhon_g06211.t1 | OG2 |
HoiH04_mRNA17954 | OG2 |
HoiH04_mRNA8845 | OG2 |
Chem_TCONS_00069212 | OG2 |
Hvul_g9232_1 | OG2 |
Sanmal_Sma_020570-T1 | OG2 |
Nemnom_g6204.t1 | OG2 |
Rhoesc_Res_010269-T1 | OG2 |
Morvir_scaffold279.g4.t1 | OG2 |
Hvul_g3619_1 | OG2 |
Chem_TCONS_00050428 | OG2 |
Dgig_XM_028537964.1 | OG2 |
Xesp_012843-T1 | OG2 |
Exapal_XM_021045304.2 | OG2 |
Nvec_v1g120055 | OG2 |
Gfas_gfas1.m1.9715.m1 | OG2 |
Amil_Amillepora19373-RA | OG2 |
Adig_XM_015900754.1 | OG2 |
Spis_XP_022789849_1 | OG2 |
Pocdam_XM_027188525.1 | OG2 |
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Gasp_gasp1.m3.1326.m1 | OG2 |
Skow_XM_006824403.1 | OG2 |
Spur_XM_030986981.1 | OG2 |
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Hsap_ENST00000310512 | OG2 |
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Acagra_model.g30125.t1 | OG2 |
Cgig_XM_011449935.3 | OG2 |
Dpul_mRNA5035 | OG2 |
Tcas_TC011992_001 | OG2 |
Cscu_XM_023357375.1 | OG2 |
Cscu_XM_023361319.1 | OG2 |
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Coeast_Cast_sb776629 | OG2 |
Coemet_Cmet_scaffold16471_01 | OG2 |
Coeast_Cast_sb776628 | OG2 |
Lampan_Lpan_8649_01 | OG2 |
Horcal_Hcv1.av93.c1.g1510.i1 | OG2 |
Pbac_sb2644947 | OG2 |
Drygla_Dgla_sb280311 | OG2 |
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Opea_EP00118_Oscarella_pearsei_P022189 | OG2 |
Plajan_Pjan_C561206_01 | OG2 |
Aque_Aqu2.1.40730_001 | OG2 |
Halamb_c51640_g1_i1mm.4085 | OG2 |
Plespi_Pspi_C588200_01 | OG2 |
Stycar_maker-SC_scaffold5371-augustus-gene-0.10-mRNA-1 | OG2 |
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Emue_Em0013g92a.t1 | OG2 |
Spolac_c101201_g2_i1_m.24851c101201-g1 | OG2 |
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