Bralan_BLAG02001201_4 | OG0 |
Xestes_maker-XT_scaffold2462-augustus-gene-0.3-mRNA-1 | OG0 |
Skow_XM_006825738.1 | OG0 |
Spur_XM_003728513.3 | OG0 |
Spur_XM_030983327.1 | OG0 |
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Cgig_XM_011452712.3 | OG1 |
Acagra_model.g12288.t1 | OG1 |
Hsap_ENST00000360131 | OG1 |
Coemet_Cmet_scaffold18772_01 | OG1 |
Eupdun_Edun_sb10706302 | OG1 |
Lampan_Lpan_10852_01 | OG1 |
Mlei_ML0069.g28.i1 | OG1 |
Lampan_Lpan_C404097_01 | OG1 |
Pbac_sb2662604 | OG1 |
Horcal_Hcv1.av93.c7.g103.i1 | OG1 |
Mertsp_Mesp_sb554247 | OG1 |
Mertsp_Mesp_sb554248 | OG1 |
Bolinf_Binf_sb1193133 | OG1 |
Bolinf_Binf_sb1193134 | OG1 |
Drygla_Dgla_sb293900 | OG1 |
Drygla_Dgla_sb293898 | OG1 |
Drygla_Dgla_sb293897 | OG1 |
Drygla_Dgla_sb293899 | OG1 |
Mertsp_Mesp_sb522245 | OG1 |
Beraby_Baby_sb349466 | OG1 |
Beraby_Baby_sb349468 | OG1 |
Beraby_Baby_sb349467 | OG1 |
Beraby_Baby_sb349464 | OG1 |
Beraby_Baby_sb349461 | OG1 |
Beraby_Baby_sb349463 | OG1 |
Beraby_Baby_sb349462 | OG1 |
Beraby_Baby_sb349465 | OG1 |
Morvir_scaffold45.g55.t3 | OG1 |
Sanmal_Sma_007942-T1 | OG1 |
Sanmal_Sma_007946-T1 | OG1 |
Nemnom_g1898.t1 | OG1 |
Rhoesc_Res_012083-T1 | OG1 |
Nemnom_g1897.t1 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P005382 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P024180 | OG1 |
Tadh_TriadT19682 | OG1 |
TrH2_TrispH2_000082-RA | OG1 |
HoiH06_mRNA8583 | OG1 |
HoiH23_PIH23_001920-RA | OG1 |
HoiH11_mRNA1705 | OG1 |
HoiH04_mRNA16294 | OG1 |
Hhon_g09945.t1 | OG1 |
Dgig_XM_028546921.1 | OG1 |
Dgig_XM_028546923.1 | OG1 |
Exapal_XM_028663440.1 | OG1 |
Nvec_NVE3440 | OG1 |
Adig_XM_015919036.1 | OG1 |
Amil_Amillepora18195-RA | OG1 |
Gfas_gfas1.m1.1978.m1 | OG1 |
Spis_XP_022807740_1 | OG1 |
Pocdam_XM_027197172.1 | OG1 |
Gasp_gasp1.m3.698.m1 | OG1 |
Fspp_ffun1.m4.19420.m1 | OG1 |
Cscu_XM_023388204.1 | OG1 |
Dpul_mRNA11247 | OG1 |
Tcas_TC016122_001 | OG1 |
Clacor_Ccor_C539263_01 | OG1 |
Chem_TCONS_00054611 | OG1 |
Leuniv_Lniv_scaffold11915_01 | OG1 |
Gracom_Gcom_scaffold17470_01 | OG1 |
Scil_scict024642.1 | OG1 |
Leuniv_Lniv_C764319_01 | OG1 |
Lcom_lcpid37994 | OG1 |
Aque_Aqu2.1.22478_001 | OG1 |
Halamb_c16979_g1_i1m.2881 | OG1 |
Halamb_c38876_g1_i1m.11186 | OG1 |
Xestes_maker-XT_scaffold34933-augustus-gene-0.3-mRNA-1 | OG1 |
Emue_Em0003g1423a.t1 | OG1 |
Plespi_Pspi_C614672_01 | OG1 |
Stycar_maker-SC_scaffold105008-snap-gene-0.2-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold86663-snap-gene-0.4-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold128537-snap-gene-0.5-mRNA-1 | OG1 |
Tetwil_g11093.t1 | OG1 |
Emue_Em0774g4a.t1 | OG1 |
Emue_Em0003g1424a.t1 | OG1 |
Spolac_c97084_g1_i3_m.14254c97084-g1 | OG1 |
Spolac_c97084_g1_i4_m.14257c97084-g1 | OG1 |
Plajan_Pjan_scaffold20137_01 | OG1 |
Opea_EP00118_Oscarella_pearsei_P013571 | OG1 |
Ocar_g1987 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P023518 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P023519 | OG1 |
Mylflu_EP00047_Mylnosiga_fluctuans_P008572 | OG1 |
Tetwil_g25629.t1 | OG1 |
Tetwil_g24578.t1 | OG1 |
Tetwil_g25369.t1 | OG1 |
Tetwil_g4723.t1 | OG1 |
Tetwil_g26655.t1 | OG1 |
Tetwil_g9377.t1 | OG1 |
Tetwil_g10460.t1 | OG1 |
Tetwil_g7528.t1 | OG1 |
Salkve_EP00050_Salpingoeca_kvevrii_P008851 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P024181 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P024178 | OG1 |
Didcos_EP00039_Didymoeca_costata_P015400 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P027347 | OG1 |
Stedip_EP00041_Stephanoeca_diplocostata_P011954 | OG1 |
Helnan_EP00037_Helgoeca_nana_P006711 | OG1 |
Acaspe_EP00035_Acanthoeca_spectabilis_P025748 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P016202 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P016203 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P009941 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P009942 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P006542 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P010276 | OG1 |
Sros_EGD80021 | OG1 |
Mbre_EDQ88114 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P006637 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P011162 | OG1 |
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