Spis_XP_022806584_1 | OG0 |
Salmac_EP00052_Salpingoeca_macrocollata_P016717 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P022973 | OG0 |
Mylflu_EP00047_Mylnosiga_fluctuans_P010935 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P020598 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P052609 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P052610 | OG0 |
Mbre_EDQ87813 | OG0 |
Chofle_EP00730_Choanoeca_flexa_P003621 | OG0 |
Choper_EP00045_Choanoeca_perplexa_P014048 | OG0 |
Microa_EP00043_Microstomoeca_roanoka_P027578 | OG0 |
Sros_EGD77827 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P006574 | OG0 |
Salinf_EP00049_Salpingoeca_infusionum_P006575 | OG0 |
Salkve_EP00050_Salpingoeca_kvevrii_P018205 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P016518 | OG0 |
Salurc_EP00051_Salpingoeca_urceolata_P016519 | OG0 |
Didcos_EP00039_Didymoeca_costata_P015193 | OG0 |
Diagra_EP00040_Diaphanoeca_grandis_P012615 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P040541 | OG0 |
Acaspe_EP00035_Acanthoeca_spectabilis_P037941 | OG0 |
Helnan_EP00037_Helgoeca_nana_P024184 | OG0 |
Opea_EP00118_Oscarella_pearsei_P005947 | OG1 |
Clacor_Ccor_scaffold5831_01 | OG1 |
Leuniv_Lniv_scaffold15594_01 | OG1 |
Leuniv_Lniv_scaffold15595_01 | OG1 |
Lcom_lcpid63022 | OG1 |
Gracom_Gcom_scaffold4450_01 | OG1 |
Scil_scict008495.1 | OG1 |
Cgig_XM_011455995.3 | OG2 |
Salkve_EP00050_Salpingoeca_kvevrii_P016685 | OG2 |
Vmul_SimC400360 | OG2 |
Coeast_Cast_sb773742 | OG2 |
Coemet_Cmet_C604118_01 | OG2 |
Lampan_Lpan_8094_01 | OG2 |
Bolinf_Binf_sb1209295 | OG2 |
Mertsp_Mesp_sb581561 | OG2 |
Bolinf_Binf_sb1249689 | OG2 |
Mertsp_Mesp_sb570990 | OG2 |
Beraby_Baby_sb345394 | OG2 |
Beraby_Baby_sb345396 | OG2 |
Beraby_Baby_sb345393 | OG2 |
Beraby_Baby_sb345395 | OG2 |
Horcal_Hcv1.av93.c10.g524.i1 | OG2 |
Pbac_sb2657462 | OG2 |
Pbac_sb2653756 | OG2 |
Pbac_sb2653755 | OG2 |
Mlei_ML1134.g24.i1 | OG2 |
Mlei_ML4202.g1.i1 | OG2 |
Drygla_Dgla_sb278715 | OG2 |
Drygla_Dgla_sb278716 | OG2 |
Drygla_Dgla_sb278713 | OG2 |
Drygla_Dgla_sb278714 | OG2 |
Plespi_Pspi_scaffold19706_01 | OG2 |
Tetwil_g21337.t2 | OG2 |
Spolac_c103051_g1_i2_m.32247c103051-g1 | OG2 |
Spolac_c103051_g1_i3_m.32249c103051-g1 | OG2 |
Emue_Em0022g152a.t1 | OG2 |
Emue_Em0460g4a.t1 | OG2 |
Emue_Em0655g4a.t1 | OG2 |
Xestes_maker-XT_scaffold102-augustus-gene-0.13-mRNA-1 | OG2 |
Halamb_c49612_g1_i3mm.3161 | OG2 |
Aque_Aqu2.1.06126_001 | OG2 |
Aque_Aqu2.1.04852_001 | OG2 |
Aque_Aqu2.1.25778_001 | OG2 |
Aque_Aqu2.1.06127_001 | OG2 |
Opea_EP00118_Oscarella_pearsei_P005948 | OG2 |
Plajan_Pjan_scaffold16791_01 | OG2 |
Plajan_Pjan_scaffold16790_01 | OG2 |
Opea_EP00118_Oscarella_pearsei_P009454 | OG2 |
Ocar_g6601 | OG2 |
Acagra_model.g13602.t1 | OG2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA183911 | OG2 |
Cscu_XM_023367945.1 | OG2 |
Cscu_XM_023357221.1 | OG2 |
Dpul_mRNA1602 | OG2 |
Tcas_TC007348_001 | OG2 |
Hsap_ENST00000264451 | OG2 |
Skow_XM_006818095.1 | OG2 |
Bralan_BLAG06000725_1 | OG2 |
Skow_XM_006812608.1 | OG2 |
Spur_XM_030980909.1 | OG2 |
Spur_XM_011667429.2 | OG2 |
TrH2_TrispH2_008960-RA | OG2 |
Tadh_TriadT25311 | OG2 |
HoiH23_PlH23_004775-RA | OG2 |
HoiH06_mRNA3159 | OG2 |
Hhon_g10992.t1 | OG2 |
HoiH04_mRNA7622 | OG2 |
Dgig_XM_028541952.1 | OG2 |
Xesp_015983-T1 | OG2 |
Hvul_g25077_1 | OG2 |
Chem_TCONS_00064949 | OG2 |
Chem_TCONS_00064948 | OG2 |
Morvir_scaffold377.g15.t1 | OG2 |
Sanmal_Sma_010914-T1 | OG2 |
Aaur_scaffold67.g24.t2 | OG2 |
Nemnom_g16474.t1 | OG2 |
Rhoesc_Res_000273-T1 | OG2 |
Exapal_XM_021050165.2 | OG2 |
Nvec_v1g207889 | OG2 |
Gfas_gfas1.m1.18587.m1 | OG2 |
Amil_Amillepora36870-RA | OG2 |
Adig_XM_015901230.1 | OG2 |
Spis_XP_022783395_1 | OG2 |
Pocdam_XM_027190035.1 | OG2 |
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Fspp_ffun1.m4.785.m1 | OG2 |
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