Stycar_augustus_masked-SC_scaffold146955-processed-gene-0.3-mRNA-1 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P005248 | OG0 |
Xestes_augustus_masked-XT_scaffold144966-processed-gene-0.0-mRNA-1 | OG0 |
Xestes_augustus_masked-XT_scaffold56060-processed-gene-0.1-mRNA-1 | OG0 |
Xestes_augustus_masked-XT_scaffold129349-processed-gene-0.2-mRNA-1 | OG0 |
Xestes_augustus_masked-XT_scaffold174193-processed-gene-0.0-mRNA-1 | OG0 |
Xestes_augustus_masked-XT_scaffold25523-processed-gene-0.0-mRNA-1 | OG2 |
Spis_XP_022778814_1 | OG3 |
Opea_EP00118_Oscarella_pearsei_P002274 | OG1 |
Xestes_maker-XT_scaffold2912-augustus-gene-0.29-mRNA-1 | OG1 |
Spolac_c102628_g3_i1_m.30259c102628-g3 | OG1 |
Emue_Em0015g1009a.t1 | OG1 |
Tetwil_g12819.t1 | OG1 |
Halamb_c41932_g1_i2mm.1026 | OG1 |
Aque_Aqu2.1.20545_001 | OG1 |
Exapal_XM_021059021.2 | OG1 |
Nvec_v1g111058 | OG1 |
Spis_XP_022781549_1 | OG1 |
Pocdam_XM_027204006.1 | OG1 |
Fspp_ffun1.m4.6237.m1 | OG1 |
Gasp_gasp1.m3.28951.m1 | OG1 |
Gfas_gfas1.m1.8885.m1 | OG1 |
Adig_XM_015906923.1 | OG1 |
Amil_Amillepora37607-RA | OG1 |
Amil_Amillepora22313-RA | OG1 |
Amil_Amillepora37605-RA | OG1 |
Chem_TCONS_00048013 | OG1 |
Hvul_g19444_1 | OG1 |
Morvir_scaffold41.g36.t1 | OG1 |
Sanmal_Sma_019218-T1 | OG1 |
Aaur_scaffold363.g12.t1 | OG1 |
Nemnom_g14772.t1 | OG1 |
Rhoesc_Res_006485-T1 | OG1 |
Dgig_XM_028545153.1 | OG1 |
Xesp_004790-T1 | OG1 |
Bralan_BLAG01002606_1 | OG1 |
Spur_XM_030991438.1 | OG1 |
Skow_XM_002742338.2 | OG1 |
Hsap_ENST00000313468 | OG1 |
Dpul_mRNA29060 | OG1 |
Cscu_XM_023360643.1 | OG1 |
Tcas_TC015022_001 | OG1 |
Lcom_lcpid192784 | OG1 |
Acagra_model.g14078.t1 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA149229 | OG1 |
Cgig_XM_011451750.3 | OG1 |
Lcom_lcpid113767 | OG1 |
Salurc_EP00051_Salpingoeca_urceolata_P027351 | OG1 |
Saldol_EP00054_Salpingoeca_dolichothecata_P037246 | OG1 |
Codhol_EP00042_Codosiga_hollandica_P002745 | OG1 |
Mylflu_EP00047_Mylnosiga_fluctuans_P003057 | OG1 |
Salmac_EP00052_Salpingoeca_macrocollata_P034180 | OG1 |
Salpun_EP00053_Salpingoeca_punica_P004562 | OG1 |
Diagra_EP00040_Diaphanoeca_grandis_P005288 | OG1 |
Acaspe_EP00035_Acanthoeca_spectabilis_P020727 | OG1 |
Helnan_EP00037_Helgoeca_nana_P038064 | OG1 |
Stedip_EP00041_Stephanoeca_diplocostata_P010370 | OG1 |
Didcos_EP00039_Didymoeca_costata_P018450 | OG1 |
Mbre_EDQ86165 | OG1 |
Chofle_EP00730_Choanoeca_flexa_P019815 | OG1 |
Choper_EP00045_Choanoeca_perplexa_P004097 | OG1 |
Sros_EGD74326 | OG1 |
Microa_EP00043_Microstomoeca_roanoka_P014572 | OG1 |
Salinf_EP00049_Salpingoeca_infusionum_P017596 | OG1 |
Clacor_Ccor_scaffold13659_01 | OG1 |
Lcom_lcpid125914 | OG1 |
Leuniv_Lniv_scaffold2370_01 | OG1 |
Scil_scict022240.1 | OG1 |
Gracom_Gcom_scaffold16419_01 | OG1 |
HoiH11_mRNA4869 | OG1 |
HoiH06_mRNA621 | OG1 |
TrH2_TrispH2_009631-RA | OG1 |
HoiH23_PlH23_011743-RA | OG1 |
Hhon_g04528.t1 | OG1 |
HoiH04_mRNA2175 | OG1 |
Eupdun_Edun_sb10636702 | OG1 |
Lampan_Lpan_5633_01 | OG1 |
Vmul_sbMO441675 | OG1 |
Coemet_Cmet_scaffold7887_01 | OG1 |
Coemet_Cmet_scaffold7888_01 | OG1 |
Coeast_Cast_sb789995 | OG1 |
Coeast_Cast_sb789996 | OG1 |
Horcal_Hcv1.av93.c5.g1067.i1 | OG1 |
Bolinf_Binf_sb1226703 | OG1 |
Bolinf_Binf_sb1226352 | OG1 |
Mertsp_Mesp_sb570841 | OG1 |
Drygla_Dgla_sb299213 | OG1 |
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Beraby_Baby_sb338645 | OG1 |
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