Skow_XM_006815240.1 | OG0 |
Cgig_XM_034465963.1 | OG0 |
Acagra_model.g27852.t1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA183430 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA226670 | OG0 |
Cscu_XM_023364661.1 | OG0 |
Dpul_mRNA18999 | OG0 |
Tcas_TC010949_001 | OG0 |
Tcas_TC013058_001 | OG0 |
Spur_NM_214494.1 | OG1 |
Bralan_BLAG19100898_1 | OG1 |
Hsap_ENST00000679126 | OG1 |
Skow_XM_002738858.2 | OG1 |
Vmul_sbMO417535 | OG1 |
Coemet_Cmet_scaffold22697_01 | OG1 |
Coeast_Cast_sb792198 | OG1 |
Coeast_Cast_sb792202 | OG1 |
Coeast_Cast_sb792201 | OG1 |
Coeast_Cast_sb792199 | OG1 |
Coeast_Cast_sb792200 | OG1 |
Lampan_Lpan_5404_01 | OG1 |
Lampan_Lpan_5405_01 | OG1 |
Bolinf_Binf_sb1207271 | OG1 |
Mertsp_Mesp_sb568446 | OG1 |
Vmul_sbMO368117 | OG1 |
Mlei_ML3209.g8.i1 | OG1 |
Beraby_Baby_sb338197 | OG1 |
Beraby_Baby_sb338203 | OG1 |
Pbac_sb2663243 | OG1 |
Pbac_sb2662133 | OG1 |
Pbac_sb2663958 | OG1 |
Horcal_Hcv1.av93.c2.g559.i1 | OG1 |
Horcal_Hcv1.av93.c2.g559.i2 | OG1 |
Horcal_Hcv1.av93.c2.g559.i3 | OG1 |
Morvir_scaffold738.g3.t1 | OG1 |
Chem_TCONS_00021611 | OG1 |
Hvul_g26943_1 | OG1 |
Morvir_scaffold738.g2.t1 | OG1 |
Sanmal_Sma_009858-T1 | OG1 |
Sanmal_Sma_026055-T1 | OG1 |
Aaur_scaffold33.g3.t1 | OG1 |
Aaur_scaffold33.g5.t2 | OG1 |
Nemnom_g19080.t1 | OG1 |
Rhoesc_Res_012992-T1 | OG1 |
Xesp_020386-T1 | OG1 |
Dgig_XM_028557924.1 | OG1 |
Exapal_XM_021043294.2 | OG1 |
Nvec_v1g193454 | OG1 |
Nvec_v1g1771 | OG1 |
Adig_XM_015896557.1 | OG1 |
Adig_XM_015892738.1 | OG1 |
Amil_Amillepora03975-RA | OG1 |
Amil_Amillepora31438-RA | OG1 |
Gfas_gfas1.m1.15878.m1 | OG1 |
Fspp_ffun1.m4.7644.m1 | OG1 |
Gasp_gasp1.m3.12790.m1 | OG1 |
Fspp_ffun1.m4.31279.m1 | OG1 |
Spis_XP_022797908_1 | OG1 |
Pocdam_XM_027187087.1 | OG1 |
TrH2_TrispH2_001050-RA | OG1 |
Tadh_TriadT52286 | OG1 |
HoiH23_PlH23_009656-RA | OG1 |
HoiH06_mRNA9646 | OG1 |
Hhon_g02872.t1 | OG1 |
HoiH04_mRNA8261 | OG1 |
Hhon_g02871.t1 | OG1 |
HoiH04_mRNA2395 | OG1 |
HoiH06_mRNA9691 | OG1 |
HoiH23_PlH23_009655-RA | OG1 |
HoiH11_mRNA6132 | OG1 |
Tadh_TriadT52287 | OG1 |
TrH2_TrispH2_001051-RA | OG1 |
Opea_EP00118_Oscarella_pearsei_P010566 | OG1 |
Opea_EP00118_Oscarella_pearsei_P010565 | OG1 |
Plajan_Pjan_scaffold3355_01 | OG1 |
Plajan_Pjan_scaffold3352_01 | OG1 |
Plajan_Pjan_scaffold3354_01 | OG1 |
Clacor_Ccor_scaffold14058_01 | OG1 |
Gracom_Gcom_C554264_01 | OG1 |
Leuniv_Lniv_scaffold24326_01 | OG1 |
Lcom_lcpid173735 | OG1 |
Lcom_lcpid68329 | OG1 |
Lcom_lcpid119030 | OG1 |
Plespi_Pspi_C615000_01 | OG1 |
Aque_Aqu2.1.38611_001 | OG1 |
Halamb_c53800_g2_i1mm.5454 | OG1 |
Stycar_maker-SC_scaffold5483-augustus-gene-0.24-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold5483-augustus-gene-0.25-mRNA-1 | OG1 |
Tetwil_g4159.t1 | OG1 |
Spolac_c103612_g2_i3_m.35159c103612-g2 | OG1 |
Emue_Em0010g357a.t1 | OG1 |
Emue_Em0010g361a.t1 | OG1 |
0.416667