Opea_EP00118_Oscarella_pearsei_P000859 | OG0 |
Opea_EP00118_Oscarella_pearsei_P006506 | OG0 |
Ocar_g6721 | OG0 |
Opea_EP00118_Oscarella_pearsei_P011878 | OG0 |
Ocar_g5733 | OG0 |
Opea_EP00118_Oscarella_pearsei_P006505 | OG0 |
Opea_EP00118_Oscarella_pearsei_P018368 | OG0 |
Opea_EP00118_Oscarella_pearsei_P018366 | OG0 |
Opea_EP00118_Oscarella_pearsei_P018365 | OG0 |
Opea_EP00118_Oscarella_pearsei_P013797 | OG0 |
Ocar_g8808 | OG0 |
Morvir_scaffold292.g5.t1 | OG0 |
Sanmal_Sma_020421-T1 | OG0 |
Morvir_scaffold292.g3.t1 | OG0 |
Aaur_scaffold140.g17.t1 | OG0 |
Nemnom_g4039.t1 | OG0 |
Rhoesc_Res_009907-T1 | OG0 |
Morvir_scaffold31.g91.t1 | OG0 |
Sanmal_Sma_027390-T1 | OG0 |
Aaur_scaffold1149.g1.t1 | OG0 |
Nemnom_g2385.t1 | OG0 |
Rhoesc_Res_001734-T1 | OG0 |
Dgig_XM_028549333.1 | OG0 |
Dgig_XM_028560730.1 | OG0 |
Exapal_XM_028661568.1 | OG0 |
Nvec_v1g223838 | OG0 |
Nvec_v1g245338 | OG0 |
Amil_Amillepora16700-RA | OG0 |
Adig_XM_015898911.1 | OG0 |
Gfas_gfas1.m1.7190.m1 | OG0 |
Gasp_gasp1.m1b.13339.m1 | OG0 |
Fspp_ffun1.m4.25061.m1 | OG0 |
Pocdam_XM_027190647.1 | OG0 |
Spis_XP_022802043_1 | OG0 |
Morvir_scaffold78.g18.t1 | OG1 |
Sanmal_Sma_020557-T1 | OG1 |
Nemnom_g3924.t1 | OG1 |
Rhoesc_Res_009861-T1 | OG1 |
Clacor_Ccor_scaffold6618_01 | OG1 |
Spur_XM_778793.5 | OG1 |
Spur_XM_030999764.1 | OG1 |
Lcom_lcpid186553 | OG1 |
Lcom_lcpid81558 | OG1 |
Lcom_lcpid140289 | OG1 |
Clacor_Ccor_C525179_01 | OG1 |
Gracom_Gcom_scaffold9530_01 | OG1 |
Lcom_lcpid71488 | OG1 |
Scil_scict000061.1 | OG1 |
Leuniv_Lniv_scaffold18032_01 | OG1 |
Tetwil_g3829.t2 | OG1 |
Lcom_lcpid204968 | OG1 |
Stycar_snap_masked-SC_scaffold938-processed-gene-0.7-mRNA-1 | OG1 |
Stycar_maker-SC_scaffold938-augustus-gene-0.9-mRNA-1 | OG1 |
Halamb_c5890_g1_i1m.936 | OG1 |
Xestes_snap_masked-XT_scaffold35674-processed-gene-0.3-mRNA-1 | OG1 |
Aque_Aqu2.1.44104_001 | OG1 |
Halamb_c55343_g1_i1mm.6490 | OG1 |
Spolac_c103958_g2_i3_m.37261c103215-g2 | OG1 |
Spolac_c103958_g2_i1_m.37256c103215-g2 | OG1 |
Emue_Em0019g642a.t1 | OG1 |
Emue_Em0695g2a.t1 | OG1 |
Spolac_c103958_g2_i6_m.37270c103215-g2 | OG1 |
Spolac_c103958_g2_i4_m.37265c103215-g2 | OG1 |
Spolac_c103958_g2_i5_m.37267c103215-g2 | OG1 |
Spolac_c103958_g1_i1_m.37254c103958-g1 | OG1 |
Emue_Em0019g641a.t1 | OG1 |
Emue_Em0695g1a.t1 | OG1 |
Spolac_c103958_g2_i2_m.37259c103215-g2 | OG1 |
Spolac_c103215_g2_i2_m.33031c103215-g2 | OG1 |
Spolac_c103215_g2_i3_m.33033c103215-g2 | OG1 |
Lcom_lcpid187797 | OG1 |
Cgig_XM_011426989.3 | OG1 |
Acagra_model.g15850.t1 | OG1 |
Lcom_lcpid139940 | OG1 |
Lcom_lcpid99371 | OG1 |
Lcom_lcpid99385 | OG1 |
Skow_XM_006820420.1 | OG1 |
Skow_XM_006820421.1 | OG1 |
Tcas_TC032215_001 | OG1 |
Bralan_BLAG04002113_1 | OG1 |
Bralan_BLAG03000040_1 | OG1 |
Bralan_BLAG01000019_1 | OG1 |
Spur_XM_030999678.1 | OG1 |
Spur_XM_030998521.1 | OG1 |
Spur_XM_003724338.2 | OG1 |
Hsap_ENST00000327757 | OG1 |
Bralan_BLAG16000594_1 | OG1 |
Lcom_lcpid189662 | OG1 |
Tetwil_g1180.t2 | OG1 |
Skow_XM_006820425.1 | OG1 |
Skow_XM_006820426.1 | OG1 |
Skow_XM_006820424.1 | OG1 |
Skow_XM_006820422.1 | OG1 |
Skow_XM_006820423.1 | OG1 |
Acagra_model.g22341.t1 | OG1 |
Cgig_XM_011452634.3 | OG1 |
Lcom_lcpid100375 | OG1 |
Lcom_lcpid100928 | OG1 |
Lcom_lcpid100126 | OG1 |
Lcom_lcpid100653 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA224221 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA121773 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA188389 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA198659 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA198661 | OG1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA194093 | OG1 |

0.416667