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•Stycar_maker-SC_scaffold5883-snap-gene-0.134-mRNA-1 | OG0 |
               •Opea_EP00118_Oscarella_pearsei_P018607 | OG0 |
       100
               •Plajan_Pjan_scaffold342_01 | OG0 |
            Clacor Ccor C520491 01 | OG1 |
                 Clacor_Ccor_C532723_01 | OG1 |
   96
               Leuniv Lniv C742181 01 | OG1 |
     100
                   •Gracom_Gcom_scaffold18801_01 | OG1 |
                 •Lcom_lcpid72640 | OG1 |
                 •Plajan_Pjan_scaffold10160_01 | OG1 |
                   Opea EP00118 Oscarella pearsei P005150 | OG1 |
         100
                    -Opea_EP00118_Oscarella_pearsei_P005151 | OG1 |
              100
                   Ocar g9201 | OG1 |
                      Opea EP00118 Oscarella pearsei P005152 | OG1 |
     91
                   •Plespi_Pspi_C609762_01 | OG1 |
                       •Stycar_maker-SC_scaffold1115-augustus-gene-0.2-mRNA-1 | OG1 |
                      Xestes maker-XT scaffold271-snap-gene-0.46-mRNA-1 | OG1 |
       95
                         •Aque_Aqu2.1.28457_001 | OG1 |
              100
                         •Halamb_c47704_g1_i5mm.2357 | OG1 |
          90
                     Tetwil g13018.t3 | OG1 |
                     Stycar_augustus_masked-SC_scaffold5883-processed-gene-0.27-mRNA-1 | OG1 |
                   100 Stycar snap masked-SC scaffold5883-processed-gene-0.53-mRNA-1 | OG1 |
            95
                      -Emue_Em0023g64a.t1 | OG1 |
                        Spolac_c100460_g1_i3_m.22498c100460-g1 | OG1 |
                         _-Spolac_c100460_g1_i1_m.22497c100460-g1 | OG1 |
                        <sup>85</sup> Spolac_c100460_g1_i4_m.22500c100460-g1 | OG1 |
                    Spur XM 030990299.1 | OG1 |
   50
           100
                 Skow_XM_006824606.1 | OG1 |
                       •Dpul mRNA527 | OG1 |
       94
            96
                     •Cscu_XM_023368152.1 | OG1 |
                     Bralan_BLAG10000427_1 | OG1 |
          45
                          +Hsap ENST00000397748 | OG1 |
                    Skow_XM_002733070.2 | OG1 |
            45
                        •Ctel gnl WGS AMQN CAPTEDRAFT mRNA151128 | OG1 |
                          Cgig_XM_011437405.3 | OG1 |
                         Acagra model.g19785.t1 | OG1 |
                    •Morvir scaffold24.g36.t1 | OG1 |
                     •Sanmal Sma 005696-T1 | OG1 |
            100
                         Aaur scaffold298.g13.t1 | OG1 |
               100
                         •Nemnom_g12987.t1 | OG1 |
     56
                     <sup>1001</sup>-Rhoesc_Res_005720-T1 | OG1 |
          88
                     •Exapal XM 021055673.2 | OG1 |
                     Nvec v1g162049 | OG1 |
              100
                  <sup>100</sup> Nvec NVE11935 | OG1 |
                   •Gfas_gfas1.m1.19414.m1 | OG1 |
            98
                      •Gfas gfas1.m1.6375.m1 | OG1 |
                        <sub>-</sub>Amil_Amillepora12790-RA | OG1 |
              100
                           •Adig_XM_015917725.1 | OG1 |
                         •Fspp_ffun1.m4.6913.m1 | OG1 |
                     <sup>99</sup><sup>1</sup>•Gasp_gasp1.m3.26590.m1 | OG1 |
                         <sub></sub>-Spis_XP_022792023_1 | OG1 |
                      <sup>100</sup><sup>1</sup>•Pocdam XM_027193188.1 | OG1 |
                             TrH2_TrispH2_007134-RA | OG1 |
                                <sub>"</sub>HoiH23_PIH23_002985-RA | OG1 |
                                  -HoiH06_mRNA11457 | OG1 |
             91
                                --Hhon_g09749.t1 | OG1 |
                             <sup>96</sup> Hhon_g10396.t1 | OG1 |
               84
                            •TrH2_TrispH2_007132-RA | OG1 |
                         •TrH2_TrispH2_007133-RA | OG1 |
                           •HoiH23_PIH23_002986-RA | OG1 |
                            51
                         <sup>100</sup> HoiH04_mRNA9279 | OG1 |
                                                                                    •Gfas_gfas1.m1.19415.m1 | OG1 |
                        •Dgɪg_XM_028539425.1 | OG1 |
                         -Xesp_009807-T1 | OG1 |
                                            Eupdun_Edun_sb10651763 | OG1 |
                                         <sup>58</sup> Eupdun Edun sb10651764 | OG1 |
            25
                                           Eupdun Edun sb10651765 | OG1 |
                                         941-Eupdun Edun sb10651766 | OG1 |
                                           Bolinf_Binf_sb1230036 | OG1 |
                                        <sup>100</sup> Mertsp_Mesp_sb578918 | OG1 |
                                   100
                                                 -Beraby_Baby_sb344341 | OG1 |
                                                  Drygla Dgla sb288751 | OG1 |
                                               100 Drygla Dgla_sb288753 | OG1 |
                                      80
                                                •Mlei_ML1603.g5.i1 | OG1 |
                                                  Mertsp Mesp sb581360 | OG1 |
                                                     Bolinf_Binf_sb1239056 | OG1 |
                                        48
                                                  <sup>100</sup> Mertsp_Mesp_sb581358 | OG1 |
                                              -Lampan_Lpan_14010_01 | OG1 |
                                                        <sub>-</sub>Pbac_sb2666557 | OG1 |
                                           68
                                                     Horcal Hcv1.av93.c9.g705.i1 | OG1 |
                                                      -Vmul Simscaffold16638 01 | OG1 |
                                             72
                                                       Coemet Cmet C599639 01 | OG1 |
                                                100
                                                            Coeast_Cast_sb786793 | OG1 |
                                                          <sup>57</sup> Coeast_Cast_sb786794 | OG1 |
                                                           Coeast Cast sb786797 | OG1 |
                                                             Coeast_Cast_sb786795 | OG1 |
                                                            481-Coeast Cast sb786796 | OG1 |
0.416667
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