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•Plajan_Pjan_scaffold23205_01 | OG0 |
                            Opea_EP00118_Oscarella_pearsei_P017223 | OG0 |
         100
                        Opea_EP00118_Oscarella_pearsei_P017221 | OG0 |
                   100
                           Opea_EP00118_Oscarella_pearsei_P017222 | OG0 |
   89
                       •Clacor_Ccor_scaffold15455_01 | OG0 |
                                   •Leuniv_Lniv_C764283_01 | OG0 |
              100
                            Gracom_Gcom_C561592_01 | OG0 |
                    82
                             •Leuniv_Lniv_scaffold23855_01 | OG0 |
                               Lcom_lcpid10015 | OG0 |
                                 •Plespi_Pspi_scaffold20653_01 | OG0 |
                                     •Lcom_lcpid186686 | OG0 |
                               •Xestes_maker-XT_scaffold2432-snap-gene-0.4-mRNA-1 | OG0 |
                                •Xestes_maker-XT_scaffold10825-augustus-gene-0.23-mRNA-1 | OG0 |
                      100
            97
                                 •Halamb_c56470_g2_i1m.43419 | OG0 |
                                 <sub>s</sub>Aque_Aqu2.1.12605_001 | OG0 |
                               <sup>1001</sup>Aque _Aqu2.1.26818_001 | OG0 |
                 100
                                 Emue_Em0023g835a.t1 | OG0 |
                                •Spolac_c103728_g1_i3_m.35810c103728-g1 | OG0 |
                             •Tetwil_g16178.t2 | OG0 |
                    98
                          <sup>97</sup>-Tetwil_g3125.t1 | OG0 |
                                 Stycar_maker-SC_scaffold649-augustus-gene-0.8-mRNA-1 | OG0
                                           -Stycar_maker-SC_scaffold2988-snap-gene-0.7-mRNA-1 | OG0 |
                         TrH2_TrispH2_004364-RA | OG0 |
                      <sup>1001</sup>-Tadh_TriadT55997 | OG0 |
                          •HoiH11 mRNA7973 | OG0 |
                   100
                               <sub>"</sub>HoiH23 PIH23_006140-RA | OG0 |
                      100
                            Hhon g07995.t1 | OG0 |
                            <sup>100</sup> HoiH04_mRNA1164 | OG0 |
   70
                                    •Eupdun_Edun_sb10618055 | OG0 |
                                     -Lampan_Lpan_9817_01 | OG0 |
                                       •Vmul_sbMO447976 | OG0 |
                               65
                                       -Coeast_Cast_sb789657 | OG0 |
                        100
                                 100
                                         Coemet_Cmet_scaffold866_01 | OG0 |
                                    99
                                       <sup>53</sup> Coeast_Cast_sb789655 | OG0 |
                                         Pbac_sb2667248 | OG0 |
                                            Horcal_Hcv1.av93.c3.g349.i1 | OG0 |
                                         100 Horcal Hcv1.av93.c3.g349.i2 | OG0 |
                                        <sub>!</sub>Drygla_Dgla_sb299432 | OG0 |
                               100
                                     <sup>100</sup> Drygla_Dgla_sb299434 | OG0 |
                                          •Beraby_Baby_sb333146 | OG0 |
                                              <sub>"</sub>Bolinf_Binf_sb1232826 | OG0 |
                                           <sup>100</sup> Mertsp_Mesp_sb577973 | OG0 |
                                             -Vmul_sbMO389661 | OG0 |
                                         <sup>100</sup> Mlei ML1959.g14.i1 | OG0 |
                                            Mertsp_Mesp_sb563506 | OG0 |
                                         <sup>100</sup> Bolinf_Binf_sb1261701 | OG0 |
                    Bralan_BLAG90000841_2 | OG0 |
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                     •Hsap_ENST00000255882 | OG0 |
                     •Skow_XM_006818625.1 | OG0 |
                       Spur XM 030991581.1 | OG0 |
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           100
                    <sup>100</sup> Spur_XM_030991598.1 | OG0 |
                        •Cscu_XM_023366259.1 | OG0 |
         85
                             •Dpul_mRNA24323 | OG0 |
              100
                           •Tcas_TC014935_001 | OG0 |
           100
                          ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA184437 | OG0 |
                            •Acagra_model.g14160.t1 | OG0 |
              100
                             Cgig_XM_034480392.1 | OG0 |
   56
                          <sup>100</sup> Cgig_XM_034480406.1 | OG0 |
                             •Chem_TCONS_00061661 | OG0 |
                           •Morvir_scaffold119.g29.t3 | OG0 |
            100
                                 •Sanmal_Sma_011504-T1 | OG0 |
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                              •Aaur_scaffold1588.g4.t1 | OG0 |
                               _-Rhoesc_Res_016192-T1 | OG0 |
                              <sup>99</sup><sup>_</sup>•Nemnom_g3865.t1 | OG0 |
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                       Dgig_XM_028547829.1 | OG0 |
                         •Xesp_000984-T1 | OG0 |
                      •Exapal XM 028658872.1 | OG0 |
       99
              100
                     •Nvec_v1g165173 | OG0 |
                    Gfas gfas1.m1.402.m1 | OG0 |
                       -Adig XM 015901701.1 | OG0 |
                            -Fspp_ffun1.m4.11128.m1 | OG0 |
                     Fspp_ffun1.m4.11127.m1 | OG0 |
                      -Spis_XP_022802324_1 | OG0 |
                   <sup>98</sup> Pocdam_XM_027184895.1 | OG0 |
0.416667
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