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Clacor Ccor scaffold2610 01 | OG0 |
            Clacor_Ccor_C520081_01 | OG0 |
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                .com | cpid38227 | OG0 |
               Scil_scict023965.1 | OG0 |
               Leuniv Lniv scaffold22253 01 | OG0 |
               •Gracom Gcom C559020 01 | OG0 |
79
                •Eupdun_Edun_sb10667993 | OG0 |
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                    Eupdun_Edun_sb10667995 | OG0 |
                  <sup>831</sup>•Eupdun_Edun_sb10667996 | OG0 |
                   •Vmul SimC406180 | OG0 |
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                     Coemet Cmet scaffold22083_01 | OG0 |
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                        -Coeast_Cast_sb760942 | OG0 |
                     <sup>100</sup> Coeast_Cast_sb760941 | OG0 |
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                  Lampan_Lpan_15520_01 | OG0 |
                        •Mlei ML0828.g4.i1 | OG0 |
                            •Mlei ML0453.g1.i1 | OG0 |
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                         Mertsp_Mesp_sb590531 | OG0 |
                 91
                            Drygla Dgla sb301929 | OG0 |
                         <sup>1001</sup>•Drygla Dgla_sb301930 | OG0 |
                   81
                          -Mertsp Mesp sb590535 | OG0 |
                           Bolinf Binf sb1237479 | OG0 |
                         <sup>45</sup> Mertsp_Mesp_sb590532 | OG0 |
                           Beraby_Baby_sb345456 | OG0 |
                             Pbac sb2644882 | OG0 |
                          <sup>100</sup> Horcal_Hcv1.av93.c7.g310.i1 | OG0 |
        •Plajan_Pjan_C570664_01 | OG0 |
          •Ocar_g6858 | OG0 |
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           100
         <sup>68]</sup>Ocar g10876 | OG0 |
           Xestes maker-XT scaffold5663-augustus-gene-0.61-mRNA-1 | OG0 |
              •Aque_Aqu2.1.42135_001 | OG0 |
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              •Halamb_c55650_g1_i4mm.6715 | OG0 |
               •Tetwil_g19749.t1 | OG0 |
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        58
                •Plespi Pspi scaffold9374 01 | OG0 |
                 Plespi Pspi scaffold9375 01 | OG0 |
                -Stycar_maker-SC_scaffold1647-snap-gene-0.55-mRNA-1 | OG0 |
      58
             79
               •Tetwil_g19746.t2 | OG0 |
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                          •Tetwil g19743.t1 | OG0 |
               -Spolac_c101798_g1_i1_m.26805c101798-g1 | OG0 |
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                 93 Emue_Em0004g926a.t1 | OG0 |
                •Morvir scaffold26.g99.t1 | OG0 |
                   Chem TCONS 00025715 | OG0 |
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                   Aaur scaffold848.g4.t1 | OG0 |
                      •Nemnom g11465.t1 | OG0 |
                   <sup>100</sup><sup>1</sup>-Rhoesc_Res_012673-T1 | OG0 |
                  ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA158714 | OG0 |
                   •Acagra_model.g20237.t1 | OG0 |
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                   •Cgig XM 034455174.1 | OG0 |
                      •Tcas TC006019 001 | OG0 |
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                      •Dpul mRNA6212 | OG0 |
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                    Cscu_XM_023374880.1 | OG0 |
                 <sup>1001</sup>Cscu_XM_023374878.1 | OG0 |
                  Bralan BLAG10000876 1 | OG0 |
                     •Hsap ENST00000426333 | OG0 |
    54
                      •Skow XM 002732897.2 | OG0 |
                     --Spur_XM_030990508.1 | OG0 |
              <sub>!</sub>Stycar_maker-SC_scaffold94064-snap-gene-0.2-mRNA-1 | OG0 |
           <sup>100</sup> -Stycar_maker-SC_scaffold1647-augustus-gene-0.47-mRNA-1 | OG0 |
                   TrH2 TrispH2_008586-RA | OG0 |
        64
                <sup>100]</sup>Tadh_TriadT22861 | OG0 |
                   •HoiH11_mRNA2095 | OG0 |
                       <sub>e</sub>HoiH23_PIH23_005666-RA | OG0 |
                    <sup>100</sup> HoiH06 mRNA300 | OG0 |
                       Hhon g08387.t1 | OG0 |
      33
                    <sup>1001</sup>HoiH04 mRNA13665 | OG0 |
              <sub>"</sub>-Dgig_XM_028558462.1 | OG0 |
           <sup>100</sup><sup>1</sup>⊸Xesp_009363-T1 | OG0 |
                   <sub>!</sub>Exapal_XM_021039530.2 | OG0 |
                <sup>100</sup>L-Exapal_XM_028657614.1 | OG0 |
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                        •Nvec _NVE22910 | OG0 |
                 <sup>1001</sup>Nvec_v1g189775 | OG0 |
                    •Spis_XP_022797434_1 | OG0 |
                 <sup>100</sup> Pocdam XM 027196604.1 | OG0 |
                      <sub>-</sub>-Fspp_ffun1.m4.23207.m1 | OG0 |
                   <sup>99</sup> Gasp_gasp1.m3.13040.m1 | OG0 |
                       •Gfas_gfas1.m1.887.m1 | OG0 |
                         <sub>_</sub>Adig_XM_015908941.1 | OG0 |
                       <sup>100</sup>Amil Amillepora12457-RA | OG0 |
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