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•Clacor_Ccor_scaffold12056_01 | OG0 |
                      •Lcom_lcpid92593 | OG0 |
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                     •Gracom_Gcom_scaffold13349_01 | OG0 |
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82
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            Stycar_maker-SC_scaffold6530-augustus-gene-0.6-mRNA-1 | OG0 |
                  -Xestes_maker-XT_scaffold122047-augustus-gene-0.24-mRNA-1 | OG0 |
                 •Halamb_c35697_g1_i1m.8878 | OG0 |
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                   -Aque Aqu2.1.02913_001 | OG0 |
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                <sup>1001</sup>Aque_Aqu2.1.16370_001 | OG0 |
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                               -Aque_Aqu2.1.08468_001 | OG0 |
                           <sup>1001</sup>Aque _Aqu2.1.17781_001 | OG0 |
                    Stycar_snap_masked-SC_scaffold6252-processed-gene-0.4-mRNA-1 | OG0 |
                   -Tetwil g26152.t1 | OG0 |
                 Emue Em0009g1132a.t1 | OG0 |
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               <sup>99</sup> Emue Em0009g1143a.t1 | OG0 |
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             •Morvir scaffold75.g5.t1 | OG1 |
               -Chem_TCONS_00012269 | OG1 |
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             -Nemnom_g12291.t1 | OG1 |
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             -Rhoesc_Res_009146-T1 | OG1 |
             Aaur scaffold6.g7.t2 | OG1 |
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            <sup>1001</sup>Sanmal_Sma_018553-T1 | OG1 |
                   •Plajan_Pjan_scaffold21449_01 | OG1 |
                     Opea EP00118 Oscarella pearsei P002731 | OG1 |
                  <sup>100</sup> Ocar _g6974 | OG1 |
                     <sub>f</sub>Hhon_g01214.t1 | OG1 |
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                  <sup>1001</sup>•HoiH04_mRNA1416 | OG1 |
                       <sub>"</sub>TrH2_TrispH2_001344-RA | OG1 |
                    <sup>100</sup> Tadh_TriadT54615 | OG1 |
                         •HoiH11 mRNA6106 | OG1 |
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                         HoiH23 PIH23 000509-RA | OG1 |
                      <sup>100</sup> HoiH06_mRNA11289 | OG1 |
                    •Nvec NVE14766 | OG1 |
                     •Nvec NVE14769 | OG1 |
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                      •Exapal XM 021060325.2 | OG1 |
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                        <sup>100</sup> Dgig_XM_028543161.1 | OG1 |
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                     •Spis XP 022795520 1 | OG1 |
                     -Pocdam XM 027197810.1 | OG1 |
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                        •Bralan BLAG02000217 1 | OG1 |
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                            •Hsap_ENST00000299518 | OG1 |
                          •Cgig XM 034470040.1 | OG1 |
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                        Saldol_EP00054_Salpingoeca_dolichothecata_P036845 | OG1 |
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                            •Codhol EP00042 Codosiga hollandica P039471 | OG1 |
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                        Mbre EDQ92233 | OG1 |
                         Chofle EP00730 Choanoeca flexa P013913 | OG1 |
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                         Choper EP00045 Choanoeca_perplexa_P011148 | OG1 |
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                              •Coeast_Cast_sb788608 | OG1 |
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                           100
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                             -Horcal Hcv1.av93.c8.g676.i1 | OG1 |
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                          <sup>100</sup> Pbac_sb2665461 | OG1 |
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                               <sup>53</sup> Mertsp_Mesp_sb578711 | OG1 |
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                                 <sup>86</sup> Bolinf Binf sb1238859 | OG1 |
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                                 98 Mertsp_Mesp_sb578713 | OG1 |
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                                 1 Beraby_Baby_sb342052 | OG1 |
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