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·Clacor_Ccor_scaffold12056_01 | OG0 |
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                    •Gracom_Gcom_scaffold13349_01 | OG0 |
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                  •Tetwil g23146.t1 | OG0 |
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            Stycar_maker-SC_scaffold6530-augustus-gene-0.6-mRNA-1 | OG0 |
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                Emue Em0009g1132a.t1 | OG0 |
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                  -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>100</sup>•Aque_Aqu2.1.16370_001 | OG0 |
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                               -Aque_Aqu2.1.08468_001 | OG0 |
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          •Morvir_scaffold75.g5.t1 | OG1 |
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                •Hvul g16704 1 | OG1 |
                •Sanmal_Sma_018553-T1 | OG1 |
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              Aaur_scaffold6.g7.t2 | OG1 |
             •Nemnom_g12291.t1 | OG1 |
              Rhoesc Res 009146-T1 | OG1 |
                       Eupdun Edun sb10650758 | OG2 |
                          •Lampan_Lpan_3477_01 | OG2 |
                            Vmul SimC398608 | OG2 |
                              ·Coeast_Cast_sb788608 | OG2 |
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                                Coemet_Cmet_scaffold1424_01 | OG2 |
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                             -Horcal_Hcv1.av93.c8.g676.i1 | OG2 |
                         <sup>100</sup> Pbac_sb2665461 | OG2 |
                              -Bolinf Binf sb1238858 | OG2 |
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                              <sup>59</sup> Mertsp Mesp_sb578712 | OG2 |
                                •Beraby_Baby_sb342050 | OG2 |
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                                  <sup>100</sup>l<sub>•</sub>Drygla Dgla_sb292957 | OG2 |
                                •Mlei_ML0769.g9.i1 | OG2 |
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                                   Bolinf_Binf_sb1267127 | OG2 |
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                    POpea EP00118 Oscarella pearsei P002731 | OG2 |
                  <sup>100</sup>-Ocar _g6974 | OG2 |
                   <sub>f</sub>Hhon_g01214.t1 | OG2 |
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                 <sup>991</sup>-HoiH04_mRNA1416 | OG2 |
                       TrH2_TrispH2_001344-RA | OG2 |
                   <sup>1001</sup>Tadh_TriadT54615 | OG2 |
                        •HoiH11 mRNA6106 | OG2 |
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                         <sub>e</sub>HoiH23_PIH23_000509-RA | OG2 |
                      <sup>100</sup> HoiH06_mRNA11289 | OG2 |
                    •Nvec NVE14766 | OG2 |
                    •Nvec NVE14769 | OG2 |
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                      •Exapal XM 021060325.2 | OG2 |
                          •Xesp 002185-T1 | OG2 |
                          Dgig_XM_028543161.1 | OG2 |
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                        <sup>1001</sup>•Dgig XM_028560670.1 | OG2 |
                  -Spis_XP_022795520 1 | OG2 |
               <sup>100</sup><sup>1</sup>-Pocdam XM_027197810.1 | OG2 |
                       •Gfas_gfas1.m1.23062.m1 | OG2 |
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                          p_gasp1.m3.26055.m1 | OG2 |
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                          •Gfas_gfas1.m1.17433.m1 | OG2 |
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                   Lcom | lcpid144899 | OG2 |
                     •Spur_XM_787412.5 | OG2 |
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                     •Skow_XM_002731321.2 | OG2 |
                        •Bralan_BLAG02000217_1 | OG2 |
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                           •Hsap ENST00000299518 | OG2 |
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                           Ctel gnl WGS AMQN CAPTEDRAFT_mRNA149799 | OG2 |
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                                  Cscu_XM_023387728.1 | OG2 |
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