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•Plajan_Pjan_C568898_01 | OG0 |
                    Opea_EP00118_Oscarella_pearsei_P026854 | OG0 |
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   60
                 ·Clacor_Ccor_scaffold3858_01 | OG0 |
                    Lcom | lcpid41502 | OG0 |
                 <sup>100</sup> Lcom_lcpid35838 | OG0 |
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                      -Leuniv_Lniv_C814650_01 | OG0 |
                        Gracom_Gcom_scaffold18866_01 | OG0 |
                             Xestes_maker-XT_scaffold4123-augustus-gene-0.16-mRNA-1 | OG0 |
                     Xestes_maker-XT_scaffold3620-snap-gene-0.14-mRNA-1 | OG0 |
                       •Halamb_c49503_g1_i1mm.3104 | OG0 |
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                          -Halamb_c49929_g1_i9mm.3278 | OG0 |
                        •Aque_Aqu2.1.00389_001 | OG0 |
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                         -Aque_Aqu2.1.10839_001 | OG0 |
                            Aque_Aqu2.1.24517_001 | OG0 |
                              --Aque_Aqu2.1.03689_001 | OG0 |
     94
                  Stycar maker-SC scaffold503-snap-gene-0.2-mRNA-1 | OG0 |
                   •Tetwil_g24232.t2 | OG0 |
            85
                         •Stycar_maker-SC_scaffold7000-augustus-gene-0.5-mRNA-1 | OG0 |
                                            -Xestes_maker-XT_scaffold4123-snap-gene-0.18-mRNA-1 | OG0 |
                  Spolac c101933 g1_i4_m.27303c101933-g1 | OG0 |
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                      •Spolac_c101933_g1_i2_m.27296c101933-g1 | OG0 |
                       •Spolac_c101933_g1_i1_m.27293c101933-g1 | OG0 |
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                      Emue Em0010g447a.t1 | OG0 |
                   <sup>100</sup> Emue_Em0481g3a.t1 | OG0 |
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                                      -Spolac_c96622_g1_i1_m.13424c101933-g1 | OG0 |
                                         Emue_Em0010g448a.t1 | OG0 |
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                           •Vmul_Simscaffold1237_01 | OG1 |
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                           Coemet Cmet scaffold7742 01 | OG1 |
                           •Lampan_Lpan_C409539_01 | OG1 |
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                                  -Eupdun_Edun_sb10690832 | OG1 |
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                              Beraby_Baby_sb326091 | OG1 |
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                                     -Lampan_Lpan_C401553_01 | OG1 |
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                                     eHorcal_Hcv1.av93.c2.g1500.i1 | OG1 |
                                       Pbac_sb2655140 | OG1 |
                           70
                                     <sup>98</sup> Pbac_sb2658162 | OG1 |
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                                      •Drygla Dgla sb288445 | OG1 |
                                         •Beraby_Baby_sb326092 | OG1 |
                                        •Bolinf_Binf_sb1314492 | OG1 |
                                    <sup>100</sup><sup>1</sup>•Mlei_ML0001.g19.i1 | OG1 |
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                       Spur_NM_001163807.1 | OG1 |
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                     Skow_XM_002738718.2 | OG1 |
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                        Bralan_BLAG03001341_1 | OG1 |
                            Lcom_lcpid247974 | OG1 |
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                        •Cgig_XM_011433419.3 | OG1 |
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                        •Xesp_013593-T1 | OG1 |
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                       rAmil_Amillepora05415-RA | OG1 |
                    <sup>1001</sup>-Adig XM_015892496.1 | OG1 |
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                        -Spis_XP_022779050_1 | OG1 |
                    <sup>100</sup><sup>1</sup>-Pocdam_XM_027183306.1 | OG1 |
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                                             •Fspp_ffun1.m4.17485.m1 | OG1 |
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                               <sup>100</sup>Tadh_TriadT19077 | OG1 |
                                    -HoiH23 PIH23 002633-RA | OG1 |
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                                  <sup>98</sup>-HoiH06 mRNA11335 | OG1 |
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                                  <sup>100</sup> HoiH04_mRNA3279 | OG1 |
                          •Hvul_g23684_1 | OG1 |
                         Chem TCONS 00011010 | OG1 |
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                            •Hvul_g23691_1 | OG1 |
            99
                          Morvir scaffold59.g60.t1 | OG1 |
                          Sanmal Sma 009044-T1 | OG1 |
              90
                         -Aaur_scaffold455.g1.t1 | OG1 |
                 100
                                •Aaur_scaffold7380.g1.t1 | OG1 |
                    98
                             •Nemnom g13535.t1 | OG1 |
                         <sup>991</sup>--Rhoesc_Res_003097-T1 | OG1 |
0.416667
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