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Opea EP00118 Oscarella pearsei P003881 | OG0 |
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                       Clacor_Ccor_scaffold10749 01 | OG0 |
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        93
                                          -Lcom | lcpid53227 | OG0 |
                                                     -Gracom_Gcom_scaffold10048_01 | OG0 |
                                         •Leuniv_Lniv_scaffold14722_01 | OG0 |
   76
          •Plespi_Pspi_scaffold18850_01 | OG0 |
                Xestes maker-XT scaffold6506-augustus-gene-0.7-mRNA-1 | OG0 |
                  •Aque Aqu2.1.36570 001 | OG0 |
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                 -Halamb_c25287_g1_i1mm.106 | OG0 |
                 •Tetwil_g4349.t2 | OG0 |
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                     Stycar_maker-SC_scaffold815-snap-gene-0.28-mRNA-1 | OG0 |
                  Stycar maker-SC scaffold7120-snap-gene-0.47-mRNA-1 | OG0 |
           98
                   •Spolac c101819 g1 i2 m.26885c101819-g1 | OG0 |
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                    Spolac_c101819_g1_i3_m.26886c101819-g1 | OG0 |
                  981-Emue Em0015g716a.t1 | OG0 |
             •Coemet_Cmet_scaffold10637_01 | OG0 |
          <sup>joj</sup>'--Vmul_SimC393064 | OG0 |
               Lampan Lpan 3735 01 | OG0 |
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                 Lampan_Lpan_3732_01 | OG0 |
               66 Lampan_Lpan_3733_01 | OG0 |
                 Horcal_Hcv1.av93.c1.g1632.i1 | OG0 |
          95
              <sup>1001</sup>•Horcal_Hcv1.av93.c1.g1632.i2 | OG0 |
                 •Mlei_ML1906.g5.i1 | OG0 |
                    Drygla_Dgla_sb288858 | OG0 |
                     Beraby Baby sb333348 | OG0 |
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                  <sup>1001</sup>•Beraby_Baby_sb333349 | OG0 |
                            Eupdun_Edun_sb10673548 | OG0 |
                         <sup>100</sup> Eupdun Edun sb10673550 | OG0 |
                           Eupdun_Edun_sb10673549 | OG0 |
                         <sup>73</sup> Eupdun_Edun_sb10673553 | OG0 |
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                              Mertsp Mesp sb563784 | OG0 |
                               Bolinf Binf sb1229138 | OG0 |
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                                    Mertsp_Mesp_sb545474 | OG0 |
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                                      Bolinf_Binf_sb1181886 | OG0 |
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                                       Beraby_Baby_sb338474 | OG0 |
                                      Beraby_Baby_sb338475 | OG0 |
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                                  Drygla Dgla sb293963 | OG0 |
                                   Drygla Dgla sb293965 | OG0 |
                                   Drygla_Dgla_sb293966 | OG0 |
                                   <sup>1</sup>•Drygla_Dgla_sb293964 | OG0 |
                        •Codhol_EP00042_Codosiga_hollandica_P017355 | OG0 |
                               -Salkve_EP00050_Salpingoeca_kvevrii_P011312 | OG0 |
                               Salurc EP00051 Salpingoeca urceolata P006977 | OG0 |
                            •Mylflu EP00047 Mylnosiga fluctuans P010617 | OG0 |
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                              -Saldol_EP00054_Salpingoeca_dolichothecata_P015227 | OG0 |
                                  Salmac_EP00052_Salpingoeca_macrocollata_P020570 | OG0 |
                                      Salpun EP00053 Salpingoeca punica P001965 | OG0 |
                                 Didcos EP00039 Didymoeca costata P030570 | OG0 |
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                                       •Diagra_EP00040_Diaphanoeca_grandis_P004065 | OG0 |
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                                        Stedip_EP00041_Stephanoeca_diplocostata_P010998 | OG0 |
                        100
                                        -Acaspe_EP00035_Acanthoeca_spectabilis_P023993 | OG0 |
                                     <sup>100</sup>Ū-Helnan EP00037_Helgoeca_nana_P002521 | OG0 |
               55
                                     •Mbre_EDQ93073 | OG0 |
                                   •Chofle_EP00730_Choanoeca_flexa_P019837 | OG0 |
                              100 Choper_EP00045_Choanoeca_perplexa_P004864 | OG0 |
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                                  <sub>!</sub>Salinf_EP00049_Salpingoeca_infusionum_P010341 | OG0 |
   76
                                      Salinf EP00049 Salpingoeca infusionum P010342 | OG0 |
                                Microa EP00043 Microstomoeca roanoka P001611 | OG0 |
                                Sros EGD76761 | OG0 |
                            •Hsap_ENST00000293845 | OG0 |
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                                     •Cscu_XM_023355566.1 | OG0 |
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                                                        -Tcas_TC015011_001 | OG0 |
              •Morvir scaffold542.g1.t1 | OG0 |
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                    •Sanmal_Sma_018826-T1 | OG0 |
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                  1001-Rhoesc_Res_006628-T1 | OG0 |
                    TrH2_TrispH2_005266-RA | OG0 |
                 <sup>[100]</sup>Tadh_TriadT51517 | OG0 |
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                     •HoiH06_mRNA10222 | OG0 |
       88
                      •HoiH23 PIH23 008551-RA | OG0 |
                       •HoiH04_mRNA6530 | OG0 |
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                       |Spur_XM_778978.5 | OG0
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                      Acagra_model.g14394.t1 | OG0 |
                       Cgig_XM_011425788.3 | OG0 |
                          <sup>1001</sup>Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA93836 | OG0 |
                 •Dgig XM 028544742.1 | OG0 |
                   •Xesp 004431-T1 | OG0 |
                   •Exapal_XM_021040413.2 | OG0 |
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                    •Nvec_v1g187067 | OG0 |
                       •Spis XP 022780526 1 | OG0 |
                    <sup>100</sup> Pocdam XM 027203325.1 | OG0 |
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                      Gasp_gasp1.m3.7485.m1 | OG0 |
                    •Gfas_gfas1.m1.239.m1 | OG0 |
                       <sub>*</sub>Amil_Amillepora30044-RA | OG0 |
                         Amil Amillepora30066-RA | OG0 |
                       <sup>51</sup> Adig XM 015922491.1 | OG0 |
0.416667
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