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Lcom lcpid231421 | OG0 |
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                    Saldol EP00054 Salpingoeca dolichothecata P014368 | OG0 |
     58
                Salurc EP00051 Salpingoeca urceolata P032721 | OG0 |
 60
               Mylflu EP00047 Mylnosiga fluctuans P009998 | OG0 |
       68
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                Salmac EP00052 Salpingoeca macrocollata P030577 | OG0 |
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                    Salpun EP00053 Salpingoeca punica P004682 | OG0 |
                                     -Plespi Pspi C581642 01 | OG0 |
                                         Diagra EP00040 Diaphanoeca grandis P032199 | OG0 |
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                                           Stedip EP00041 Stephanoeca diplocostata P037831 | OG0 |
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                                         Acaspe EP00035 Acanthoeca spectabilis P007742 | OG0 |
                                     100 Helnan EP00037 Helgoeca nana P013365 | OG0 |
                     Microa_EP00043_Microstomoeca_roanoka_P011008 | OG0 |
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                                •Sros EGD72484 | OG0 |
                       Salinf EP00049 Salpingoeca infusionum P018064 | OG0 |
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                       Mbre EDQ87614 | OG0 |
                      •Chofle EP00730 Choanoeca flexa P001459 | OG0 |
                      •Choper_EP00045_Choanoeca_perplexa_P000465 | OG0 |
                •Codhol EP00042 Codosiga hollandica P037010 | OG1 |
       100
                        Codhol EP00042 Codosiga hollandica P042734 | OG1 |
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                  Diagra EP00040 Diaphanoeca grandis P016524 | OG1 |
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                  Stedip EP00041 Stephanoeca diplocostata P035376 | OG1 |
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                    Acaspe EP00035 Acanthoeca spectabilis P039895 | OG1 |
                <sup>100</sup><sup>1</sup>-Helnan EP00037_Helgoeca_nana_P013055 | OG1 |
                •Lcom | lcpid190537 | OG1 |
                     Vmul SimC391988 | OG1 |
                       •Coemet_Cmet_C583470_01 | OG1 |
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     54
                          Coeast Cast sb787528 | OG1 |
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                             Coeast Cast sb787530 | OG1 |
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                          <sup>100</sup> Coeast Cast sb787529 | OG1 |
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                          •Eupdun_Edun_sb10637959 | OG1 |
                         Mertsp_Mesp_sb560210 | OG1 |
                              Beraby Baby sb328841 | OG1 |
                               Mlei ML0758.g1.i1 | OG1 |
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                            100 Vmul_sbMO372562 | OG1 |
                               <u>-</u>Drygla_Dgla_sb297189 | OG1 |
                            <sup>100</sup>-Drygla Dgla_sb297190 | OG1 |
                                   Horcal Hcv1.av93.c2.g1656.i1 | OG1 |
                                 <sup>63</sup> Pbac sb2656071 | OG1 |
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                                 <sup>100</sup><sup>1</sup>-Pbac sb2668893 | OG1 |
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                     •Cgig XM 034457272.1 | OG1 |
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                         •Cscu XM 023367228.1 | OG1 |
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                         •Skow_XM_002738150.2 | OG1 |
                        ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA183405 | OG1 |
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                49
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                                   -Xesp_013159-T1 | OG1 |
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                                      <sub>!</sub>Spis_XP_022802899_1 | OG1 |
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                                   <sup>100</sup> Pocdam XM 027201372.1 | OG1 |
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                                        Adig_XM_015895856.1 | OG1 |
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                               Bralan_BLAG03001362_1 | OG1 |
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                                     -Hsap ENST00000677851 | OG1 |
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                                        •HoiH23_PIH23_004927-RA | OG1 |
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                                      <sup>96</sup><sup>™</sup>-HoiH06_mRNA13620 | OG1 |
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                                         <sub>r</sub>Hhon_g07166.t1 | OG1 |
                                      <sup>100</sup> HoiH04_mRNA6044 | OG1 |
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                                         Opea_EP00118_Oscarella_pearsei_P025063 | OG1 |
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                                      <sup>100</sup> Ocar_g4434 | OG1 |
                              43
                                       •Clacor_Ccor_scaffold11568_01 | OG1 |
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                                         -Scil_scict014872.1 | OG1 |
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                                      <sup>74</sup>-Leuniv_Lniv_C796250_01 | OG1 |
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                                        •Tetwil_g8701.t1 | OG1 |
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                                               Stycar_maker-SC_scaffold143305-augustus-gene-0.31-mRNA-1 | OG1 |
                                                 Xestes maker-XT scaffold116143-snap-gene-0.34-mRNA-1 | OG1 |
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0.416667
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