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•Plespi_Pspi_C611762 01 | OG0 |
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                                   100
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                                                   <sub>|</sub>-Aque_Aqu2.1.29110_001 | OG0 |
                                                <sup>100</sup> Ague Aqu2.1.01412_001 | OG0 |
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                                  Tetwil g12889.t2 | OG0 |
                                 -Stycar_maker-SC_scaffold142519-snap-gene-0.105-mRNA-1 | OG0 |
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                                       •Emue_Em0019g153a.t1 | OG0 |
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85
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                 Clacor Ccor C515565 01 | OG0 |
                             Leuniv Lniv scaffold25336 01 | OG0 |
     63
                                •Scil_scict019292.1 | OG0 |
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                                        •Lcom_lcpid112989 | OG0 |
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                                       Gracom_Gcom_C557508_01 | OG0 |
   63
                      Leuniv Lniv scaffold25723 01 | OG0 |
                         Lcom | lcpid46833 | OG0 |
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                           -Lcom_lcpid124333 | OG0 |
                         65 Lcom_lcpid27915 | OG0 |
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                        Scil scict013153.1 | OG0 |
                               Gracom Gcom scaffold8744 01 | OG0 |
                            <sup>100</sup> Gracom_Gcom_scaffold8745_01 | OG0 |
                Popea_EP00118_Oscarella_pearsei_P005178 | OG1 |
             <sup>100</sup> Ocar_g3253 | OG1 |
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              Plajan_Pjan_scaffold21201_01 | OG1 |
           <sup>1001</sup>Plajan Pjan scaffold21202 01 | OG1 |
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                               •Lcom_lcpid139665 | OG1 |
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                                Saldol EP00054 Salpingoeca dolichothecata P004541 | OG1 |
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                                   Salmac EP00052 Salpingoeca macrocollata P008565 | OG1 |
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                                   Salpun EP00053 Salpingoeca punica P010776 | OG1 |
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                                    •Mylflu_EP00047_Mylnosiga_fluctuans_P024063 | OG1 |
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                                      •Codhol EP00042 Codosiga hollandica P043526 | OG1 |
                                          •Codhol EP00042 Codosiga hollandica P048558 | OG1 |
                                       •Didcos_EP00039_Didymoeca_costata_P025622 | OG1 |
                                             Stedip_EP00041_Stephanoeca_diplocostata_P020808 | OG1 |
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                                100
                                           Diagra_EP00040_Diaphanoeca_grandis_P024884 | OG1 |
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                                         -Acaspe EP00035 Acanthoeca spectabilis P020792 | OG1 |
                                          •Helnan EP00037 Helgoeca nana P012361 | OG1 |
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                                                         Salurc_EP00051_Salpingoeca_urceolata_P002672 | OG1 |
                                       •Mbre_EDQ88732 | OG1 |
                                       •Chofle_EP00730_Choanoeca_flexa_P020235 | OG1 |
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                                       Choper EP00045 Choanoeca perplexa P009657 | OG1 |
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                                           Salinf EP00049 Salpingoeca infusionum P018059 | OG1 |
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                                   •Coemet_Cmet_scaffold19617_01 | OG1 |
                              <sup>100</sup> Coeast Cast sb782254 | OG1 |
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                                 <sup>97</sup> Beraby_Baby_sb343774 | OG1 |
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                     Spur XM 030999425.1 | OG1 |
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                            •Hsap ENST00000316924 | OG1 |
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                            •Hsap_ENST00000259708 | OG1 |
                                           Hsap ENST00000470402 | OG1 |
                          Lcom | lcpid170713 | OG1 |
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                         •Cgig_XM_034468117.1 | OG1 |
                         Lcom | lcpid240950 | OG1 |
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                       ·Cscu XM 023357625.1 | OG1 |
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                           •Cscu_XM_023387069.1 | OG1 |
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                                              •Dpul mRNA1018 | OG1 |
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                            <sub>*</sub>TrH2_TrispH2_007657-RA | OG1 |
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                         <sup>100</sup> Tadh TriadT64331 | OG1 |
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                           <sup>100</sup> HoiH06 mRNA861 | OG1 |
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                       •Hvul_g4113_1 | OG1 |
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                        •Chem_TCONS_00011432 | OG1 |
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                       •Sanmal_Sma_027127-T1 | OG1 |
                          -Aaur_scaffold1201.g5.t1 | OG1 |
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                             -Nemnom_g18289.t1 | OG1 |
                         <sup>100</sup> Rhoesc_Res_001843-T1 | OG1 |
                 ⊢•Nvec_v1g164652 | OG1 |
                    •Exapal_XM_028659486.1 | OG1 |
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                              -Gasp gasp1.m3.29715.m1 | OG1 |
           100
                     Amil Amillepora15142-RA | OG1 |
                  <sup>100</sup>Ū-Adig XM_015903356.1 | OG1 |
              92
                     •Spis_XP_022810576_1 | OG1 |
                  <sup>1001</sup>-Pocdam_XM_027194327.1 | OG1 |
                    Gfas gfas1.m1.2167.m1 | OG1 |
                        •Gasp_gasp1.m3.4578.m1 | OG1 |
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                       <sup>100</sup> Fspp ffun1.m4.11414.m1 | OG1 |
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