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Salpun_EP00053_Salpingoeca_punica_P011879 | OG1 |
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                   Diagra EP00040 Diaphanoeca grandis P029378 | OG2 |
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                   Diagra EP00040_Diaphanoeca_grandis_P029379 | OG2 |
                    Acaspe EP00035 Acanthoeca spectabilis P019744 | OG2 |
                    •Helnan_EP00037_Helgoeca_nana_P027783 | OG2 |
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                          Acaspe EP00035 Acanthoeca spectabilis P019751 | OG2 |
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                           Helnan_EP00037_Helgoeca_nana_P027787 | OG2 |
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                                   -Acaspe EP00035 Acanthoeca spectabilis P034990 | OG2 |
                 86
                       Stedip EP00041 Stephanoeca diplocostata P021487 | OG2 |
                           Stedip EP00041 Stephanoeca diplocostata P021485 | OG2 |
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                         <sup>57]</sup>Stedip EP00041 Stephanoeca diplocostata P021489 | OG2 |
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                              Stedip_EP00041_Stephanoeca_diplocostata_P021488 | OG2 |
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                    Salinf_EP00049_Salpingoeca_infusionum_P012421 | OG2 |
          91
                     •Microa_EP00043_Microstomoeca_roanoka_P012699 | OG2 |
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                  <sup>100</sup>Ū-Sros EGD82248 | OG2 |
                   Mbre EDQ91661 | OG2 |
                      Chofle EP00730 Choanoeca flexa P005609 | OG2 |
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                     Codhol EP00042 Codosiga hollandica P057523 | OG2 |
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                   Lcom | lcpid132186 | OG2 |
                   Lcom | lcpid132837 | OG2 |
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                      TrH2_TrispH2_001839-RA | OG0 |
                   <sup>100</sup> Tadh_TriadT25995 | OG0 |
                      •HoiH11 mRNA4025 | OG0 |
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                           +HoiH23 PIH23_000338-RA | OG0 |
                   99
                        <sup>100</sup> HoiH06_mRNA13438 | OG0 |
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                     100
                           Hhon g05848.t1 | OG0 |
                        <sup>100</sup> HoiH04 mRNA9616 | OG0 |
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                             Pbac_sb2662641 | OG0 |
                               Horcal Hcv1.av93.c3.g644.i1 | OG0 |
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                        <sup>100</sup><sup>_</sup>•Adig_XM_015900598.1 | OG0 |
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