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Drygla_Dgla_sb292706 | OG0 |
              <sup>100</sup>Drygla Dgla sb292707 | OG0
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        58
              <sub>[</sub>•Horcal_Hcv1.av93.c6.g749.i1 | OG0 |
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                                 -Pbac sb2639674 | OG0 |
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                     Beraby_Baby_sb337239 | OG0 |
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                 Mertsp Mesp sb560369 | OG0 |
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                    Lcom | lcpid168613 | OG0 |
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                     •Cgig XM 011458472.3 | OG0 |
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      46
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                       <sup>95</sup>-HoiH06_mRNA2646 | OG0 |
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                         100 HoiH04 mRNA10642 | OG0 |
                             TrH2 TrispH2 001515-RA | OG0 |
                          <sup>100</sup> Tadh_TriadT23778 | OG0 |
                           •Clacor_Ccor_C529945_01 | OG0 |
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                           •Leuniv_Lniv_scaffold21979_01 | OG0 |
                             Gracom Gcom scaffold17851 01 | OG0 |
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                              -Lcom | lcpid98403 | OG0 |
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                                   •Mylflu EP00047 Mylnosiga fluctuans P002069 | OG0 |
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                                          Salinf EP00049 Salpingoeca infusionum P018599 | OG0 |
                                       Salmac_EP00052_Salpingoeca_macrocollata_P000548 | OG0 |
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                                      Salmac EP00052 Salpingoeca macrocollata P033684 | OG0 |
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                                           -Microa_EP00043_Microstomoeca_roanoka_P015198 | OG0 |
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                                             •Sros_EGD76288 | OG0 |
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                                             •Mbre EDQ87269 | OG0 |
                                             -Chofle EP00730 Choanoeca flexa P003353 | OG0 |
                                        99 Choper EP00045 Choanoeca_perplexa_P020438 | OG0 |
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                                 •Chem_TCONS_00061300 | OG0 |
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                                   •Sanmal_Sma_025180-T1 | OG0 |
                                   Aaur scaffold87.g15.t1 | OG0 |
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                                      •Rhoesc_Res_010809-T1 | OG0 |
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                                  •Pocdam_XM_027184820.1 | OG0 |
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                              <sup>100</sup><sup>-</sup>Spis_XP_022779650_1 | OG0 |
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                              99 Fspp ffun1.m4.12906.m1 | OG0 |
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                                             Chofle_EP00730_Choanoeca_flexa_P010170 | OG0 |
                                            Didcos EP00039 Didymoeca costata P018280 | OG0 |
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                                               r-Acaspe_EP00035_Acanthoeca_spectabilis_P021144 | OG0 |
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                                             <sup>57</sup> Helnan_EP00037_Helgoeca_nana_P002779 | OG0 |
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                                             Stycar_snap_masked-SC_scaffold71669-processed-gene-0.3-mRNA-1 | OG0 |
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                                                  -Xestes_augustus_masked-XT_scaffold166413-processed-gene-0.2-mRNA-1 | OG0 |
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                                             Xestes augustus masked-XT scaffold15810-processed-gene-0.4-mRNA-1 | OG0 |
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                                            Xestes_augustus_masked-XT_scaffold119503-processed-gene-0.2-mRNA-1 | OG0 |
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                                              Aaur scaffold10.g80.t1 | OG0 |
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    Xestes augustus masked-XT scaffold167838-processed-gene-0.0-mRNA-1 | OG0 |

0.416667
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•Eupdun Edun sb10624912 | OG0