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Salinf_EP00049_Salpingoeca_infusionum_P026748 | OG0 |
                           Sros EGD77900 | OG0 |
                              •Microa_EP00043_Microstomoeca_roanoka_P010376 | OG0 |
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                 100
                          1001-Microa_EP00043_Microstomoeca_roanoka_P010375 | OG0 |
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                            Choper EP00045 Choanoeca perplexa P007724 | OG0 |
                             Chofle EP00730 Choanoeca flexa P000340 | OG0 |
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                          <sup>100</sup> Chofle_EP00730_Choanoeca_flexa_P000341 | OG0 |
                        Salkve_EP00050_Salpingoeca_kvevrii_P010210 | OG0 |
       63
                                                       -Pbac sb2647140 | OG0 |
                               Didcos EP00039 Didymoeca costata P023278 | OG0 |
                                             Codhol EP00042 Codosiga hollandica P057795 | OG0 |
                                   67 Codhol_EP00042_Codosiga_hollandica_P057800 | OG0 |
       46
                            •Mylflu_EP00047_Mylnosiga_fluctuans_P007609 | OG0 |
                           Diagra EP00040 Diaphanoeca_grandis_P021241 | OG0 |
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                                 -Acaspe EP00035 Acanthoeca spectabilis P008797 | OG0 |
                                  •Stedip_EP00041_Stephanoeca_diplocostata_P020764 | OG0 |
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                   Salmac EP00052 Salpingoeca macrocollata P007879 | OG1 |
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                     Salpun EP00053 Salpingoeca punica P008436 | OG1 |
                        Saldol EP00054 Salpingoeca dolichothecata P020366 | OG1 |
                              •Clacor_Ccor_C523415_01 | OG1 |
                                  Gracom_Gcom_C546874_01 | OG1 |
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                                     •Scil_scict004409.4 | OG1 |
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                                  Leuniv Lniv scaffold19137_01 | OG1 |
                                   Lcom | lcpid43358 | OG1 |
                                    •Plespi_Pspi_scaffold17950_01 | OG1 |
                                     •Tetwil_g22895.t1 | OG1 |
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                                        Stycar maker-SC scaffold4638-augustus-gene-0.5-mRNA-1 | OG1 |
                                        Xestes maker-XT scaffold6586-augustus-gene-0.8-mRNA-1 | OG1 |
                                          •Halamb c56609 g1 i1mm.7254 | OG1 |
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          92
                                          Aque_Aqu2.1.24904_001 | OG1 |
                                       <sup>100</sup><sup>1</sup>--Aque_Aqu2.1.11560_001 | OG1 |
   61
                                  Vmul sbMO436286 | OG1 |
                                    <sub>[</sub>Coemet_Cmet_scaffold4247_01 | OG1 |
                                 <sup>100</sup> Coemet Cmet_scaffold4248_01 | OG1 |
                                     -Pbac sb2647139 | OG1 |
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                                     Horcal_Hcv1.av93.c6.g998.i1 | OG1 |
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                                  -Beraby_Baby_sb335902 | OG1 |
                                    Bolinf Binf sb1241567 | OG1 |
                                 <sup>100</sup> Mertsp_Mesp_sb602087 | OG1 |
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                                       •Drygla_Dgla_sb300822 | OG1 |
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                                          •Lampan_Lpan_15591_01 | OG1 |
                                         •Mlei ML3814.g1.i1 | OG1 |
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                                        Bolinf Binf sb1225008 | OG1 |
                                      <sup>1001</sup>Mertsp_Mesp_sb531840 | OG1 |
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                             •Plajan_Pjan_C552085_01 | OG1 |
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                             Rhoesc_Res_015425-T1 | OG1 |
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                           •Xesp 021358-T1 | OG1 |
                            •Exapal_XM_021049542.2 | OG1 |
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                            •Nvec_v1g95124 | OG1 |
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                              Spis XP 022801508 1 | OG1 |
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                                     -Adig_XM_015892845.1 | OG1 |
                               -Amil Amillepora07231-RA | OG1 |
                             <sup>98</sup>-Adig XM _015899779.1 | OG1 |
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                              Spur XM 030998969.1 | OG1 |
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                                  Hsap ENST00000554271 | OG1 |
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                               •Skow XM 006815727.1 | OG1 |
                             Ctel gnl WGS AMQN CAPTEDRAFT_mRNA151190 | OG1 |
                               Acagra model.g16353.t1 | OG1 |
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                               Cscu XM 023360084.1 | OG1 |
                                 •Tcas TC009612 001 | OG1 |
                                      Dpul mRNA28313 | OG1 |
              86
                                   -HoiH23_PIH23_010491-RA | OG1 |
                                <sup>97</sup>-HoiH06_mRNA14586 | OG1 |
                                    •HoiH04_mRNA15801 | OG1 |
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                                    •Tadh_wf_g1504.t1 | OG1 |
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                                     Tadh TriadT52772 | OG1 |
                                    <sup>491</sup>•Tadh_TriadT62506 | OG1 |
0.416667
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