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·Codhol_EP00042_Codosiga_hollandica_P022840 | OG0 |
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                                   •Mylflu_EP00047_Mylnosiga_fluctuans_P005566 | OG0 |
    59
                                  -Salpun_EP00053_Salpingoeca_punica_P003635 | OG0 |
                                   Salmac_EP00052_Salpingoeca_macrocollata_P017866 | OG0 |
                           Salinf EP00049 Salpingoeca infusionum P011315 | OG0 |
                               Salkve EP00050 Salpingoeca kvevrii P021315 | OG0 |
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     56
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                                      Didcos EP00039 Didymoeca costata P008955 | OG0 |
                         Sros EGD78673 | OG0 |
                          •Microa_EP00043_Microstomoeca_roanoka_P007977 | OG0 |
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                             Diagra EP00040 Diaphanoeca grandis P042906 | OG0 |
                             Salurc EP00051 Salpingoeca urceolata P026437 | OG0 |
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                86
                                          -Lcom_lcpid245256 | OG0 |
                                   •Mbre EDQ87701 | OG0 |
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                                         •Diagra_EP00040_Diaphanoeca_grandis_P011681 | OG0 |
                                            Stedip_EP00041_Stephanoeca_diplocostata_P025057 | OG0 |
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                                               -Stedip EP00041 Stephanoeca diplocostata P025059 | OG0 |
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    Saldol EP00054 Salpingoeca dolichothecata P021925 | OG1 |

                                •Hvul_g23095_1 | OG1 |
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                           •Sanmal Sma 020657-T1|OG1|
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                                      -Rhoesc_Res_009698-T1 | OG1 |
                                        •Nemnom_g3369.t1 | OG1 |
                         Morvir scaffold35.g26.t1 | OG1 |
           63
                                             Vmul Simscaffold12731 01 | OG1 |
                                               Coemet Cmet scaffold614 01 | OG1 |
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                                               Coeast Cast sb805693 | OG1 |
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                                                 Coeast_Cast_sb805685 | OG1 |
                                                    "Coeast_Cast_sb805686 | OG1 |
                                                 <sup>100</sup> Coeast_Cast_sb805694 | OG1 |
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                                                 •Lampan_Lpan_C409109_01 | OG1 |
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                                                     •Pbac_sb2649196 | OG1 |
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                                                  <sup>100</sup> Horcal_Hcv1.av93.c5.g868.i1 | OG1 |
                                                     •Beraby_Baby_sb332905 | OG1 |
                                         95
                                                       •Mlei ML2395.g21.i1 | OG1 |
                                                        Mertsp Mesp sb559269 | OG1 |
                                                     <sup>100</sup> Mertsp_Mesp_sb559266 | OG1 |
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                                                       Drygla_Dgla_sb284405 | OG1 |
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                                                       Drygla_Dgla_sb284404 | OG1 |
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                           Plajan Pjan scaffold10475_01 | OG1 |
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                        <sup>1001</sup>-Plajan_Pjan_scaffold10477_01 | OG1 |
                               Opea EP00118 Oscarella pearsei P005188 | OG1 |
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                                  Ocar g11134 | OG1 |
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                              Exapal XM 028656959.1 | OG1 |
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                           Gasp gasp1.m3.12728.m1 | OG1 |
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                                 Skow XM 002734686.2 | OG1 |
                                        •Dpul mRNA29810 | OG1 |
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                                    Lcom lcpid58037 | OG1 |
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                                       Scil scict003161.2 | OG1 |
                                              •Gracom Gcom scaffold9074 01 | OG1 |
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                                          Cscu XM 023358268.1 | OG1 |
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                                    •Plespi Pspi scaffold4377 01 | OG1 |
                                 Xestes_maker-XT_scaffold56383-augustus-gene-0.5-mRNA-1 | OG1 |
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                                      Stycar_maker-SC_scaffold4590-snap-gene-0.13-mRNA-1 | OG1 |
                        68
                             100
                                          •Tetwil g11934.t1 | OG1 |
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