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Lcom lcpid77576 | OG3 |
                Saldol EP00054 Salpingoeca dolichothecata P001315 | OG0 |
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 54
                      Diagra_EP00040_Diaphanoeca_grandis_P038441 | OG0 |
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                           -Acaspe_EP00035_Acanthoeca_spectabilis_P034938 | OG0 |
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   46
                           -Helnan_EP00037_Helgoeca_nana_P032175 | OG0 |
              92
                     -Stedip_EP00041_Stephanoeca_diplocostata_P010259 | OG0 |
                                             -Lcom_lcpid255161 | OG0 |
                        Salurc_EP00051_Salpingoeca_urceolata_P027372 | OG0 |
     38
                                    •Microa EP00043 Microstomoeca roanoka P005800 | OG0 |
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                                      Sros_EGD76223 | OG0 |
                              •Mbre_EDQ86753 | OG0 |
            81
                             -Chofle_EP00730_Choanoeca_flexa_P005823 | OG0 |
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                             Choper_EP00045_Choanoeca_perplexa_P013535 | OG0 |
                      -Salkve_EP00050_Salpingoeca_kvevrii_P006653 | OG0 |
                        •Lcom | lcpid195601 | OG0 |
                           Salmac EP00052 Salpingoeca macrocollata P004432 | OG0 |
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                          Codhol EP00042 Codosiga hollandica P038330 | OG0 |
                                                Salinf_EP00049_Salpingoeca_infusionum P024094 | OG0 |
                          •Lcom_lcpid237631 | OG1 |
                    rHorcal_Hcv1.av93.c6.g673.i1 | OG1 |
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                      <sub>!</sub>Eupdun_Edun_sb10668102 | OG1 |
    54
                   <sup>100</sup>•Eupdun_Edun_sb10668103 | OG1 |
                         •Vmul_Simscaffold11011_01 | OG1 |
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                             •Coemet_Cmet_C604810_01 | OG1 |
                                -Coeast_Cast_sb803001 | OG1 |
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                                  Coeast_Cast_sb803000 | OG1 |
                                94 Coeast Cast sb803004 | OG1 |
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                        100°
                               Coeast_Cast_sb803002 | OG1 |
                                 "Coeast_Cast_sb803003 | OG1 |
                               <sup>21</sup> Coeast_Cast_sb803005 | OG1 |
                               Coeast Cast sb802997 | OG1 |
                                 Coeast_Cast_sb802996 | OG1 |
                  53
                                     Coeast_Cast_sb802998 | OG1 |
                                   <sup>57</sup> Coeast_Cast_sb802999 | OG1 |
                        •Lampan_Lpan_3045_01 | OG1 |
                             Mertsp Mesp sb592764 | OG1 |
                                Bolinf_Binf_sb1259692 | OG1 |
                             66 Mertsp_Mesp_sb592762 | OG1 |
                             Bolinf_Binf_sb1259693 | OG1 |
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                               Mertsp_Mesp_sb592763 | OG1 |
                             80 | Mertsp_Mesp_sb592765 | OG1 |
                                Mlei_ML4094.g9.i1 | OG1 |
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                                <sup>55</sup> Vmul_sbMO470525 | OG1 |
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                                Bolinf_Binf_sb1253085 | OG1 |
                                  Bolinf Binf sb1253084 | OG1 |
                                     Bolinf_Binf_sb1253086 | OG1 |
                                       Mertsp_Mesp_sb590965 | OG1 |
                                        Bolinf_Binf_sb1253087 | OG1 |
                                       <sup>57</sup><sup>1</sup>-Mertsp_Mesp_sb590966 | OG1 |
 54
                                Bolinf Binf sb1171902 | OG1 |
                                  Bolinf Binf sb1310807 | OG1 |
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                                      Beraby_Baby_sb345143 | OG1 |
                                    <sup>92</sup> Beraby_Baby_sb345144 | OG1 |
                                       <sub>•</sub>Beraby_Baby_sb345145 | OG1 |
                          81
                                    <sup>100</sup> Beraby_Baby_sb345146 | OG1 |
                                Drygla Dgla sb302990 | OG1 |
                                  Drygla_Dgla_sb302991 | OG1 |
                                    Drygla_Dgla_sb302993 | OG1 |
                                      Drygla Dgla sb302994 | OG1 |
                                        Drygla_Dgla_sb302997 | OG1 |
                                          Drygla_Dgla_sb302995 | OG1 |
                                            Drygla Dgla sb302996 | OG1 |
                                          <sup>57</sup> Drygla_Dgla_sb302992 | OG1 |
                   Opea EP00118 Oscarella pearsei P025467 | OG2 |
                <sup>100</sup> Ocar_g2092 | OG2 |
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                Plajan_Pjan_scaffold4594_01 | OG2 |
             100 Plajan_Pjan_scaffold4596_01 | OG2 |
                Plajan_Pjan_scaffold4595_01 | OG2 |
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                      <sub>5</sub>TrH2_TrispH2_009290-RA | OG2 |
                   <sup>[100]</sup>Tadh TriadT21889 | OG2 |
                         •HoiH23_PIH23_002914-RA | OG2 |
                100
                      <sup>100</sup> -HoiH06_mRNA2907 | OG2 |
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                      100 HoiH04 mRNA7203 | OG2 |
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                      •Dpul_mRNA12564 | OG2 |
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                      •Hsap ENST00000360876 | OG2 |
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                       Spur_XM_030983906.1 | OG2 |
                      •Cgig_XM_011434667.3 | OG2 |
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                        Lcom Icpid128186 | OG2 |
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                           -Scil scict002767.2 | OG2 |
                             Leuniv_Lniv_scaffold14581_01 | OG2 |
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     47
                          <sup>1001</sup>Leuniv _Lniv_scaffold14583_01 | OG2 |
                         -Lcom_lcpid54042 | OG2 |
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                      <sup>100</sup>1•Lcom | lcpid50254 | OG2 |
                    99
                           •Gracom Gcom scaffold16719 01 | OG2 |
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                             •Scil_scict013273.1 | OG2 |
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                          rTetwil_g1402.t1 | OG2 |
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                             •Aque Aqu2.1.43324 001 | OG2 |
                             <sub>r</sub>Halamb_c33541_g1_i1m.7706 | OG2 |
                           100 Halamb_c31636_g1_i2mm.234 | OG2 |
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                         •Rhoesc Res 013961-T1 | OG2 |
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                             -Nemnom g7429.t1 | OG2 |
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                   -Xesp_024030-T1 | OG2 |
                   •Nvec_v1g113484 | OG2 |
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                       _•Adig_XM_015922559.1 | OG2 |
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                     Gfas_gfas1.m1.22242.m1 | OG2 |
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                           Gasp_gasp1.m3.15438.m1 | OG2 |
                        <sup>100</sup>-Gasp_gasp1.m3.15439.m1 | OG2 |
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0.416667
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