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Salmac_EP00052_Salpingoeca_macrocollata_P004433 | OG0 |
                                        Didcos EP00039 Didymoeca costata P029127 | OG0 |
    84
         85
                             Salpun_EP00053_Salpingoeca_punica_P004539 | OG0 |
 88
                           ·Salurc_EP00051_Salpingoeca_urceolata_P024814 | OG0 |
                                   •Acaspe EP00035 Acanthoeca_spectabilis_P037281 | OG0 |
      91
                                     •Helnan EP00037 Helgoeca nana P029174 | OG0 |
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                                 Stedip_EP00041_Stephanoeca_diplocostata_P025106 | OG0 |
                                  Diagra_EP00040_Diaphanoeca_grandis_P016858 | OG0 |
                       •Salkve_EP00050_Salpingoeca_kvevrii_P013840 | OG1 |
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                      -Salkve EP00050 Salpingoeca kvevrii P013841 | OG1 |
                                    Codhol EP00042 Codosiga hollandica P022836 | OG2 |
                                       •Codhol_EP00042_Codosiga_hollandica_P022757 | OG2 |
                 79
                                 <sup>1001</sup>-Codhol_EP00042_Codosiga_hollandica_P022760 | OG2 |
                                 Salinf_EP00049_Salpingoeca_infusionum_P009212 | OG2 |
              84
                                    •Sros EGD73446 | OG2 |
                                   Microa EP00043 Microstomoeca roanoka P010537 | OG2 |
                69
                            100
                                     Microa EP00043 Microstomoeca roanoka P010538 | OG2 |
                   96
                                        •Mbre EDQ92329 | OG2 |
                                      Chofle_EP00730_Choanoeca_flexa_P008506 | OG2 |
                                      Choper EP00045 Choanoeca perplexa P013455 | OG2 |
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           96
                                        Choper EP00045 Choanoeca perplexa P013458 | OG2 |
 88
                            -Salmac_EP00052_Salpingoeca_macrocollata_P016535 | OG2 |
                  50
                                •Salpun_EP00053_Salpingoeca_punica_P011423 | OG2 |
                                 Didcos_EP00039_Didymoeca_costata_P029431 | OG2 |
               59
                     93
                                    Mylflu EP00047 Mylnosiga fluctuans P005325 | OG2 |

    Saldol EP00054 Salpingoeca dolichothecata P021594 | OG2 |

                            1001-Saldol_EP00054_Salpingoeca_dolichothecata_P021597 | OG2 |
             92
                             Salurc_EP00051_Salpingoeca_urceolata_P015805 | OG2 |
                  91
                                 •Salkve_EP00050_Salpingoeca_kvevrii_P006565 | OG2 |
                                Acaspe EP00035 Acanthoeca spectabilis P034473 | OG2 |
                           100
                                   Helnan EP00037 Helgoeca nana P028932 | OG2 |
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                                     Diagra EP00040 Diaphanoeca grandis P031520 | OG2 |
                                   Stedip_EP00041_Stephanoeca_diplocostata_P008160 | OG2 |
                               Stedip_EP00041_Stephanoeca_diplocostata_P008158 | OG2 |
                              Aque Aqu2.1.39139 001 | OG2 |
                                 Xestes maker-XT scaffold71-augustus-gene-0.38-mRNA-1 | OG2 |
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                              Halamb_c53024_g2_i1m.34533 | OG2 |
                                   •Plespi_Pspi_C611192_01 | OG2 |
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                               •Tetwil_g17365.t2 | OG2 |
                                •Stycar_maker-SC_scaffold13185-augustus-gene-0.7-mRNA-1 | OG2 |
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                              Spolac c91558 g1 i1 m.7009c91558-g1 | OG2 |
                                          •Emue_Em0008g1226a.t1 | OG2 |
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                             <sup>94</sup> Emue Em0008g1221a.t1 | OG2 |
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                              Leuniv Lniv scaffold12776 01 | OG2 |
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                                •Scil_scict006027.2 | OG2 |
          100
                     100
                              Lcom | lcpid106866 | OG2 |
                           <sup>100</sup>-Lcom_lcpid102197 | OG2 |
                                -Pbac sb2650010 | OG2 |
                                  Horcal Hcv1.av93.c4.g27.i2 | OG2 |
                             100
                                   Horcal Hcv1.av93.c4.g27.i3 | OG2 |
                                    Horcal Hcv1.av93.c4.g27.i4 | OG2 |
                                     Horcal Hcv1.av93.c4.g27.i1 | OG2 |
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                                    1-Horcal_Hcv1.av93.c4.g27.i5 | OG2 |
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                                     Coemet Cmet scaffold22202 01 | OG2 |
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                                 100
                                     •Vmul_sbMO421934 | OG2 |
                                    Bolinf Binf sb1195398 | OG2 |
                             81
                                 <sup>100</sup> Mertsp_Mesp_sb540089 | OG2 |
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                                     Bolinf Binf sb1249578 | OG2 |
                                       Mlei ML0943.g35.i1 | OG2 |
                                    100 Vmul_sbMO426274 | OG2 |
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                                 01H11_MRNA2068 | OG2 |
                                  <sub>[</sub>TrH2_TrispH2_004564-RA | OG2 |
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                                         •Tadh_TriadT27042 | OG2 |
                                      HoiH23 PIH23 010287-RA | OG2 |
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                                        HoiH06_mRNA9152 | OG2 |
                                     Hhon g06913.t1 | OG2 |
                                   <sup>100</sup>-HoiH04 mRNA6011 | OG2 |
                               Plajan_Pjan_scaffold13640_01 | OG2 |
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                                    •Opea_EP00118_Oscarella_pearsei_P004128 | OG2 |
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                                 Opea_EP00118_Oscarella_pearsei_P004129 | OG2 |
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                                  •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA17752 | OG2 |
                                 •Acagra_model.g31130.t1 | OG2 |
                        76
                                  Cgig_XM_011446683.3 | OG2 |
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                                  •Skow_XM_002735232.2 | OG2 |
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                                      •Hsap_ENST00000400930 | OG2 |
                                            -Cscu_XM_023382598.1 | OG2 |
                                       Spur_XM_030979648.1 | OG2 |
                                    <sup>100</sup> Spur XM_030986978.1 | OG2 |
                                   Dgig_XM_028538267.1 | OG2 |
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                                     •Exapal_XM_021056958.1 | OG2 |
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                                               -Adig_XM_015909199.1 | OG2 |
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                                        Fspp_ffun1.m4.18029.m1 | OG2 |
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                                           -Spis XP 022809386 1 | OG2 |
                                        <sup>100</sup>-Pocdam XM 027197757.1 | OG2 |
                                       •Tcas TC013489 001 | OG2 |
                            72
                                        •Dpul_mRNA9968 | OG2 |
                                     Chem_TCONS_00059166 | OG2 |
                          32
                                      •Hvul_g2371_1 | OG2 |
                                       •Morvir scaffold159.g26.t1 | OG2 |
                            99
                                         •Hvul_g2382_1 | OG2 |
                                      •Sanmal_Sma_020059-T1 | OG2 |
                               82
                                   <sup>100</sup> Sanmal_Sma_004487-T1 | OG2 |
                                       •Aaur_scaffold209.g9.t1 | OG2 |
                                          Nemnom g4105.t1 | OG2 |
                                      <sup>100</sup> -Rhoesc Res 010058-T1 | OG2 |
0.416667
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