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Salurc EP00051 Salpingoeca urceolata P004905 | OG0 |
                     •Microa EP00043 Microstomoeca roanoka P002476 | OG0 |
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                                   •Lcom_lcpid229415 | OG0 |
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                   Salinf EP00049 Salpingoeca infusionum P003773 | OG0 |
                    •Mbre EDQ84393 | OG0 |
     39
                  Chofle_EP00730_Choanoeca_flexa P019110 | OG0 |
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                  •Choper_EP00045_Choanoeca_perplexa_P011945 | OG0 |
                    •Mylflu_EP00047_Mylnosiga_fluctuans_P002424 | OG1 |
                     Salkve_EP00050_Salpingoeca kvevrii P015921 | OG1 |
         60
                      -Salpun EP00053_Salpingoeca_punica_P014564 | OG1 |
       38
                              Lcom | lcpid155225 | OG1 |
                      Salmac EP00052 Salpingoeca macrocollata P014341 | OG1 |
   54
                1001 Salmac_EP00052 Salpingoeca macrocollata P014343 | OG1 |
               Saldol EP00054 Salpingoeca dolichothecata P021593 | OG1 |
                  Saldol EP00054 Salpingoeca dolichothecata P021590 | OG1 |
                    Didcos_EP00039_Didymoeca_costata_P019594 | OG1 |
                       Stedip EP00041 Stephanoeca diplocostata P038488 | OG1 |
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                  <sup>1001</sup>Stedip_EP00041_Stephanoeca_diplocostata_P038489 | OG1 |
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                       Diagra EP00040 Diaphanoeca grandis P016829 | OG1 |
                           •Helnan EP00037 Helgoeca nana P033493 | OG1 |
                            •Acaspe_EP00035_Acanthoeca_spectabilis_P023069 | OG1 |
                   Codhol EP00042 Codosiga hollandica P045354 | OG2 |
                    Codhol_EP00042_Codosiga_hollandica_P045353 | OG2 |
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                       Codhol EP00042 Codosiga hollandica P045355 | OG2 |
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                             Gracom Gcom C554430 01 | OG2 |
                             •Scil_scict005167.1 | OG2 |
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                             Eupdun_Edun_sb10610486 | OG2 |
                                     •Coemet Cmet scaffold21872 01 | OG2 |
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   85
                      100
                                     Vmul_Simscaffold10236_01 | OG2 |
                                     Horcal Hcv1.av93.c10.g45.i1 | OG2 |
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                                   <sup>100</sup>-Pbac_sb2655333 | OG2 |
                                    Nmul sbMO399898 | OG2 |
                           93
                                 100 Mlei_ML1427.g2.i1 | OG2 |
                                       Drygla Dgla sb275329 | OG2 |
                                        <sub>f</sub>Beraby_Baby_sb324204 | OG2 |
                                     <sup>1001</sup>-Beraby_Baby_sb324205 | OG2 |
     80
                      Plespi_Pspi_scaffold6664_01 | OG2 |
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                100
                       •Halamb_c54245_g1_i1m.37756 | OG2 |
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                         •Spolac c98542 g1 i1 m.17470c98542-g1 | OG2 |
                           rEmue_Em0001g1243a.t1 | OG2 |
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                     100
                        <sup>100</sup>Emue Em0001g1158a.t1 | OG2 |
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                         r-Tetwil_g24843.t1 | OG2 |
                      <sup>100</sup> Tetwil_g24847.t1 | OG2 |
                      Stycar_snap_masked-SC_scaffold59892-processed-gene-0.1-mRNA-1 | OG2 |
                    <sup>981</sup>Stycar_snap_masked-SC_scaffold49621-processed-gene-0.2-mRNA-1 | OG2 |
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                    •Plajan_Pjan_C564538_01 | OG2 |
                     Opea EP00118 Oscarella_pearsei_P002212 | OG2 |
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                  100 Ocar_g3104 | OG2 |
                         •Dgig_XM_028550531.1 | OG2 |
                              •Xesp_000227-T1 | OG2 |
                      •Exapal XM 021061013.2 | OG2 |
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                        •Nvec_v1g182319 | OG2 |
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                           Amil_Amillepora22008-RA | OG2 |
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                        <sup>100</sup><sup>1</sup>•Adig_XM_015898942.1 | OG2 |
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                            <sub>[</sub>•Spis_XP_022789745_1 | OG2 |
                          <sup>891</sup>-Pocdam XM 027190146.1 | OG2 |
                                •Gfas_gfas1.m1.1071.m1 | OG2 |
                            <sup>70</sup>-Fspp ffun1.m4.6834.m1 | OG2 |
                               •Gasp_gasp1.m3.24044.m1 | OG2 |
            83
                            40 Gfas_gfas1.m1.22298.m1 | OG2 |
                         Hsap_ENST00000359087 | OG2 |
                             <sub>[</sub>TrH2_TrispH2_004504-RA | OG2 |
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                          <sup>100</sup> Tadh_TriadT27487 | OG2 |
                              •HoiH11_mRNA9340 | OG2 |
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                                 <sub>|</sub>•HoiH23_PIH23_005562-RA | OG2 |
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                               <sup>991</sup>•HoiH06_mRNA12964 | OG2 |
                                  Hhon_g07424.t1 | OG2 |
                               <sup>1001</sup>HoiH04_mRNA16195 | OG2 |
                          •Bralan_BLAG07000434_1 | OG2 |
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                          -Spur_XM_003728760.3 | OG2 |
                          Skow_XM_006821407.1 | OG2 |
                                   •Dpul_mRNA3747 | OG2 |
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                                      •Tcas_TC003419_001 | OG2 |
                                  Cscu_XM_023384512.1 | OG2 |
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                                 -Cscu_XM_023381675.1 | OG2 |
                               <sup>97</sup> Cscu XM_023381684.1 | OG2 |
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                                     Acagra_model.g30730.t1_model.g30731.t1 | OG2 |
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                                   <sup>99</sup>Acagra_model.g30730.t1_model.g30731.t1.1.5d401217 | OG2 |
                                              •Morvir scaffold58.g48.t1 | OG2 |
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                                                •Hvul_g10720_1 | OG2 |
                                        Sanmal_Sma_023002-T1 | OG2 |
                             100
                                        •Aaur_scaffold1697.g5.t1 | OG2 |
                                           •Nemnom_g2742.t1 | OG2 |
                                      <sup>100</sup><sup>L</sup>-Rhoesc_Res_003849-T1 | OG2 |
0.416667
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