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•Mylflu EP00047 Mylnosiga fluctuans P002950 | OG0 |
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                      •Codhol_EP00042_Codosiga_hollandica_P014796 | OG0 |
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                 ·Salurc_EP00051_Salpingoeca_urceolata_P032025 | OG0 |
                 Saldol EP00054 Salpingoeca dolichothecata P003247 | OG0 |
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                  Salkve_EP00050_Salpingoeca_kvevrii_P005978 | OG0 |
                                  Didcos EP00039 Didymoeca costata P023771 | OG0 |
                           Diagra EP00040 Diaphanoeca grandis P040451 | OG0 |
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                           Stedip EP00041 Stephanoeca diplocostata P011079 | OG0 |
                          Acaspe EP00035 Acanthoeca spectabilis P022232 | OG0 |
                           •Helnan_EP00037_Helgoeca_nana_P008700 | OG0 |
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                             •Microa EP00043 Microstomoeca roanoka P012128 | OG0 |
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                              •Sros EGD74314 | OG0 |
                         Salinf EP00049 Salpingoeca infusionum P018149 | OG0 |
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                               •Mbre EDQ92197 | OG0 |
                              ·Chofle_EP00730_Choanoeca_flexa_P016269 | OG0 |
                             Choper EP00045 Choanoeca perplexa P003025 | OG0 |
                                  Lcom | lcpid63014 | OG1 |
                                   •Gracom_Gcom_scaffold18318_01 | OG1 |
                                     -Scil_scict015397.4 | OG1 |
                                 88 Leuniv_Lniv_scaffold17056_01 | OG1 |
      91
                  Clacor Ccor scaffold11931 01 | OG1 |
                  Lcom Icpid66414 | OG1 |
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                  •Scil scict010513.2 | OG1 |
                  •Leuniv Lniv scaffold20074 01 | OG1 |
                   Gracom Gcom C553572 01 | OG1 |
             Plajan Pjan_C553331_01 | OG1 |
                    Opea EP00118 Oscarella pearsei P000109 | OG1 |
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               Opea EP00118 Oscarella pearsei P000108 | OG1 |
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                    Eupdun Edun sb10654741 | OG1 |
                  <sup>98</sup> Eupdun Edun_sb10654742 | OG1 |
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                       Coemet Cmet scaffold22477 01 | OG1 |
                    <sup>100</sup><sup>1</sup>•Vmul_Simscaffold6365_01 | OG1 |
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                       Drygla Dgla sb285294 | OG1 |
                         Horcal_Hcv1.av93.c8.g735.i1 | OG1 |
                      <sup>100</sup> Pbac_sb2650064 | OG1 |
                   <sub>"</sub>HoiH11_mRNA5293 | OG1 |
                        TrH2_TrispH2_008255-RA | OG1 |
                     <sup>100</sup>Tadh TriadT52100 | OG1 |
                       HoiH23 PIH23 007302-RA | OG1 |
                          Hhon_g09296.t1 | OG1 |
                       <sup>100</sup> HoiH04_mRNA509 | OG1 |
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                    •Hsap ENST00000431926 | OG1 |
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                        •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA158276 | OG1 |
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                64
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                       Skow XM 002742356.2 | OG1 |
                          Spur XM 030993376.1 | OG1 |
                               -Cscu XM 023358304.1 | OG1 |
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                           <sup>100</sup> Cscu XM 023378920.1 | OG1 |
                                Bralan BLAG02000665 1 | OG1 |
                                 •Tcas TC012028 001 | OG1 |
                                -Dpul mRNA3235 | OG1 |
                           •Plespi Pspi C605416 01 | OG1 |
                                     →Stycar_maker-SC_scaffold5915-snap-gene-0.93-mRNA-1 | OG1 |
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                              •Tetwil g7549.t2 | OG1 |
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                              Xestes maker-XT scaffold384-snap-gene-0.14-mRNA-1 | OG1 |
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                                  rague_Aqu2.1.12952_001 | OG1 |
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                          •Chem_TCONS_00072549 | OG1 |
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                                Nemnom g16998.t1 | OG1 |
                             <sup>100</sup>-Rhoesc Res 013695-T1 | OG1 |
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                            Spis_XP_022804356_1 | OG1 |
                           23 Pocdam_XM_027182750.1 | OG1 |
                             <sup>12</sup> Nvec NVE3993 | OG1 |
0.416667
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