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Sanmal Sma 005683-T1 | OG0 |
            Nemnom g16727.t1 | OG0 |
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                •Morvir_scaffold70.g16.t1 | OG0 |
                     -Mylflu_EP00047_Mylnosiga_fluctuans_P009289 | OG0 |
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   16
                        <sup>97</sup>Codhol_EP00042_Codosiga_hollandica_P054461 | OG0 |
                       Salkve EP00050 Salpingoeca kvevrii P006866 | OG0 |
                          Diagra EP00040 Diaphanoeca grandis P018996 | OG0 |
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              18
                         Acaspe EP00035 Acanthoeca_spectabilis_P023455 | OG0 |
                        •Helnan_EP00037_Helgoeca_nana_P002520 | OG0 |
                              Stedip_EP00041_Stephanoeca_diplocostata_P020904 | OG0 |
                                 Stedip EP00041 Stephanoeca diplocostata P020902 | OG0 |
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                                   Stedip EP00041 Stephanoeca diplocostata P020905 | OG0 |
                           Didcos EP00039 Didymoeca costata P030767 | OG0 |
                         ·Salmac_EP00052_Salpingoeca_macrocollata_P019855 | OG0 |
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                            •Salpun_EP00053_Salpingoeca_punica_P009149 | OG0 |
                        Saldol EP00054 Salpingoeca dolichothecata P010205 | OG0 |
                           Salurc EP00051 Salpingoeca urceolata P027550 | OG0 |
                             Salinf EP00049 Salpingoeca infusionum P023677 | OG0 |
                                •Chofle_EP00730_Choanoeca_flexa_P017553 | OG0 |
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                                 Choper_EP00045_Choanoeca_perplexa_P012218 | OG0 |
                              Sros EGD75991 | OG0 |
                                  •Microa EP00043 Microstomoeca roanoka P016802 | OG0 |
                              Microa EP00043 Microstomoeca roanoka P016803 | OG0 |
               Clacor_Ccor_scaffold5881_01 | OG0 |
               -Scil scict013984.1 | OG0 |
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                  •Lcom_lcpid37431 | OG0 |
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                 Gracom Gcom C547868 01 | OG0 |
                  Chem TCONS 00061891 | OG0 |
   41
                    TrH2_TrispH2_011520-RA | OG0 |
                 <sup>100</sup><sup>L</sup>•Tadh_TriadT50087 | OG0 |
                    •HoiH11_mRNA9347 | OG0 |
                       -HoiH23 PIH23 002723-RA | OG0 |
                     98 -HoiH06 mRNA9099 | OG0 |
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                      100 HoiH04 mRNA7721 | OG0 |
                 •Plespi Pspi C599576 01 | OG0 |
                        •Aque_Aqu2.1.23241_001 | OG0 |
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                       Tetwil g4286.t2 | OG0 |
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                               •Stycar_maker-SC_scaffold6519-snap-gene-0.87-mRNA-1 | OG0 |
                       -Spolac c99242 g1 i1 m.19157c99242-g1 | OG0 |
                   <sup>100</sup> -Emue_Em0015g1264a.t1 | OG0 |
                 •Plajan_Pjan_scaffold22454_01 | OG0 |
        42
                              •Cscu XM 023384677.1 | OG0 |
                           •Hsap ENST00000395384 | OG0 |
      86
                  Opea EP00118 Oscarella pearsei P027244 | OG0 |
               <sup>100</sup> Ocar_g1680 | OG0 |
                -Acagra_model.g23699.t1 | OG0 |
                   Cgig XM 011452562.3 | OG0 |
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                    Spur XM 030991768.1 | OG0 |
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                    Bralan BLAG01001507 1 | OG0 |
                    •Skow XM 006813455.1 | OG0 |
     58
               •Dgig_XM_028546774.1 | OG0 |
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                   •Nvec v1g117116 | OG0 |
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