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•Mylflu_EP00047_Mylnosiga_fluctuans_P016343 | OG0 |
                <sup>100</sup> Mylflu_EP00047_Mylnosiga_fluctuans_P016344 | OG0 |
                         Codhol EP00042 Codosiga hollandica P024877 | OG0 |
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  100
                              Codhol EP00042 Codosiga hollandica P024875 | OG0 |
                        Salpun_EP00053_Salpingoeca_punica_P003442 | OG0 |
     94
                         -Salmac_EP00052_Salpingoeca_macrocollata_P004534 | OG0 |
                                   •Mbre EDQ85331 | OG0 |
                                 Choper EP00045 Choanoeca perplexa P012991 | OG0 |
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                                  Chofle_EP00730_Choanoeca_flexa_P000043 | OG0 |
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                               100 Chofle EP00730_Choanoeca_flexa_P000042 | OG0 |
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                                            Salinf_EP00049_Salpingoeca_infusionum_P000830 | OG0 |
                                    Sros EGD75385 | OG0 |
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                                       Microa EP00043 Microstomoeca roanoka P017044 | OG0 |
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                                    1001-Microa_EP00043_Microstomoeca_roanoka_P017046 | OG0 |
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                             Saldol_EP00054_Salpingoeca_dolichothecata_P037665 | OG0 |
                             Salurc EP00051 Salpingoeca urceolata P010551 | OG0 |
                          Didcos EP00039 Didymoeca costata P018948 | OG0 |
                           Stedip_EP00041_Stephanoeca_diplocostata_P004662 | OG0 |
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                           Diagra_EP00040_Diaphanoeca_grandis_P024313 | OG0 |
                             Acaspe_EP00035_Acanthoeca_spectabilis_P003742 | OG0 |
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                         <sup>100</sup>-Helnan_EP00037_Helgoeca_nana_P012368 | OG0 |
                 Stycar maker-SC scaffold3321-augustus-gene-0.6-mRNA-1 | OG0 |
                        -Aque_Aqu2.1.39810_001 | OG0 |
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                    <sup>100</sup><sup>1</sup>-Halamb_c42430_g1_i2m.14924 | OG0 |
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                         •Spolac_c100473_g1_i1_m.22536c100473-g1 | OG0 |
                           Emue Em0015g154a.t1 | OG0 |
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                        <sup>1001</sup>Emue_Em0015g157a.t1 | OG0 |
                 •HoiH11_mRNA4802 | OG0 |
                      •TrH2_TrispH2_000319-RA | OG0 |
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                   <sup>100</sup><sup>-</sup>Tadh_TriadT30201 | OG0 |
                        HoiH04 mRNA17306 | OG0 |
                     <sup>100</sup> Hhon g06985.t1 | OG0 |
                        HoiH06 mRNA13213 | OG0 |
                     <sup>100</sup> -HoiH23 PIH23 000374-RA | OG0 |
              •Tcas TC006653 001 | OG0 |
                    •Dpul mRNA809 | OG0 |
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                              •Scil scict014417.1 | OG0 |
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                            <sup>100</sup> Gracom_Gcom_scaffold3563_01 | OG0 |
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                                      Leuniv_Lniv_scaffold11476_01 | OG0 |
                                    <sup>711</sup>Leuniv_Lniv_scaffold11475_01 | OG0 |
                 •Bralan_BLAG01001280_1 | OG0 |
                     •Cscu_XM_023380077.1 | OG0 |
                    ·Skow XM 002735948.2 | OG0 |
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                          •Acagra_model.g14584.t1 | OG0 |
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                            •Cgig_XM_011437162.3 | OG0 |
                   •Xesp 005101-T1 | OG0 |
                   •Dgig XM 028540815.1 | OG0 |
                    Nvec v1g179634 | OG0 |
                     •Exapal_XM_021049461.2 | OG0 |
                      Adig_XM_015906968.1 | OG0 |
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                        <sup>971</sup>-Pocdam_XM_027202812.1 | OG0 |
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                             •Hvul_g24915_1 | OG0 |
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                          •Hvul_g24909_1 | OG0 |
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                           •Aaur_scaffold978.g3.t3 | OG0 |
                           Rhoesc Res 006891-T1 | OG0 |
                        <sup>1001</sup>•Nemnom_g7746.t1 | OG0 |
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                          Coemet_Cmet_scaffold678_01 | OG0 |
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                                Coeast Cast sb784525 | OG0 |
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                             <sup>100</sup><sup>1</sup>-Coeast_Cast_sb784524 | OG0 |
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                               •Beraby_Baby_sb340065 | OG0 |
                                  Horcal Hcv1.av93.c3.g1247.i1 | OG0 |
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