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-Xestes_maker-XT_scaffold134388-augustus-gene-0.2-mRNA-1 | OG0 |
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              -Aque_Aqu2.1.37906_001 | OG0 |
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             •Tetwil g12584.t1 | OG0 |
             Stycar maker-SC scaffold4650-snap-gene-0.7-mRNA-1 | OG0 |
                               -Emue_Em0018g1148a.t1 | OG0 |
   92
               Spolac c99490 g1 i1 m.19800c99490-g1 | OG0 |
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                 •Spolac c99490 g2 i1 m.19802c99490-g1 | OG0 |
                  •Emue Em0018g1149a.t1 | OG0 |
          •Plespi_Pspi_scaffold14063_01 | OG0 |
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                        Codhol_EP00042_Codosiga_hollandica_P032577 | OG0 |
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                     Saldol_EP00054_Salpingoeca_dolichothecata_P013885 | OG0 |
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                  77<sup>1</sup>-Saldol EP00054 Salpingoeca_dolichothecata_P013883 | OG0 |
                     •Mylflu EP00047 Mylnosiga fluctuans P007400 | OG0 |
                       Salpun EP00053 Salpingoeca punica P017139 | OG0 |
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        83
                         Salmac_EP00052_Salpingoeca_macrocollata_P013875 | OG0 |
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                     <sup>100</sup> Salmac_EP00052_Salpingoeca_macrocollata_P013876 | OG0 |
                         Didcos EP00039 Didymoeca costata P016971 | OG0 |
                          Diagra_EP00040_Diaphanoeca_grandis_P020142 | OG0 |
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                            -Acaspe_EP00035_Acanthoeca_spectabilis_P012710 | OG0 |
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                             •Helnan_EP00037_Helgoeca_nana_P037825 | OG0 |
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                             -Stedip EP00041 Stephanoeca diplocostata P003642 | OG0 |
                         1001 Stedip EP00041 Stephanoeca diplocostata P003643 | OG0 |
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                        <sub>'</sub>Salurc_EP00051_Salpingoeca_urceolata_P020697 | OG0 |
                           Salurc_EP00051_Salpingoeca_urceolata_P020696 | OG0
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                        <sup>1001</sup>--Salurc_EP00051_Salpingoeca_urceolata_P020698 | OG0 |
                            Salinf EP00049 Salpingoeca infusionum P015067 | OG0 |
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                               Sros EGD77881 | OG0 |
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                              Microa EP00043 Microstomoeca roanoka P012604 | OG0 |
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                           Mbre EDQ87782 | OG0 |
     100
                              •Chofle EP00730 Choanoeca flexa P006114 | OG0 |
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                          1001 Chofle EP00730 Choanoeca_flexa_P006116 | OG0 |
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                               •Choper_EP00045_Choanoeca_perplexa_P014622 | OG0 |
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                   POpea EP00118 Oscarella pearsei P010376 | OG0 |
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                <sup>100</sup> Gcar g7414 | OG0 |
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                               Gracom Gcom_C521494_01 | OG0 |
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                              Horcal_Hcv1.av93.c6.g690.i1 | OG0 |
                            <sup>97</sup>-Pbac sb2644228 | OG0 |
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                                Drygla Dgla sb288171 | OG0 |
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                                •Beraby_Baby_sb356401 | OG0 |
                                  •Lampan_Lpan_12831_01 | OG0 |
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                                      Bolinf Binf sb1284600 | OG0 |
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                                 Beraby Baby sb354396 | OG0 |
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                                           Coeast Cast sb776730 | OG0 |
                                       <sup>1001</sup>Coeast Cast_sb776731 | OG0 |
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                                       691Coeast_Cast_sb776733 | OG0 |
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                         •Bralan BLAG03000347 1 | OG0 |
                             Ctel_gnl_WGS_AMQN_CAPTEDRAFT mRNA182010 | OG0 |
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                             Spur XM 030989368.1 | OG0 |
                            Skow XM 006814824.1 | OG0 |
                              Cscu XM 023388518.1 | OG0 |
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                          Dgig XM 028539189.1 | OG0 |
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                            Amil Amillepora00748-RA | OG0 |
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                         <sup>100</sup> Adig XM 015897743.1 | OG0 |
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                               •Fspp_ffun1.m4.25323.m1 | OG0 |
                                 •Gasp_gasp1.m3.19397.m1 | OG0 |
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                                   <sub>-</sub>-Spis_XP_022795279_1 | OG0 |
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                                 <u>-Nemnom_g2289.t1 | OG0 |</u>
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                              <sup>1</sup>00<sup>1</sup>Nemnom_g12527.t1 | OG0 |
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                                <sup>1001</sup>Tadh TriadT49643 | OG0 |
                                    •HoiH11 mRNA6242 | OG0 |
                                      -HoiH04_mRNA9358 | OG0 |
                                         -HoiH23_PIH23_008730-RA | OG0 |
                                      <sup>100</sup>-HoiH06 mRNA13541 | OG0 |
0.416667
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