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•Codhol_EP00042_Codosiga_hollandica_P042674 | OG0 |
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                     Stedip EP00041 Stephanoeca diplocostata P009132 | OG0 |
                           Acaspe EP00035 Acanthoeca spectabilis P007733 | OG0 |
                          Helnan EP00037 Helgoeca nana P006034 | OG0 |
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                    ·Saldol_EP00054_Salpingoeca_dolichothecata_P029555 | OG0 |
                              Salurc_EP00051_Salpingoeca_urceolata_P023296 | OG0 |
                                 Salurc EP00051 Salpingoeca urceolata P023295 | OG0 |
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                                           Salurc EP00051 Salpingoeca urceolata P023292 | OG0 |
                                      Salurc EP00051 Salpingoeca urceolata P023289 | OG0 |
     78
                                   ToolSalurc_EP00051_Salpingoeca_urceolata_P023294 | OG0 |
                  Salmac_EP00052_Salpingoeca_macrocollata_P007417 | OG0 |
         98
                       Salpun EP00053 Salpingoeca punica P014497 | OG0 |
                   Mylflu EP00047 Mylnosiga fluctuans P001972 | OG0 |
       92
                    Salkve EP00050 Salpingoeca kvevrii P017392 | OG0 |
                 Galkve EP00050 Salpingoeca_kvevrii_P017393 | OG0 |
         63
                                  Microa EP00043 Microstomoeca roanoka P025767 | OG0 |
           64
                                    •Sros EGD82809 | OG0 |
                              Salinf EP00049 Salpingoeca infusionum P002255 | OG0 |
                99
                               •Mbre EDQ92783 | OG0 |
                              •Chofle_EP00730_Choanoeca_flexa_P007534 | OG0 |
                               Choper EP00045 Choanoeca perplexa P007459 | OG0 |
               TrH2 TrispH2 006957-RA | OG0 |
            100 Tadh TriadT14661 | OG0 |
               •HoiH11 mRNA3459 | OG0 |
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                   -HoiH23_PIH23_004757-RA | OG0 |
                 <sup>991</sup>-HoiH06 mRNA4127 | OG0 |
                    Hhon g09460.t1 | OG0 |
                 <sup>100</sup> HoiH04 mRNA8486 | OG0 |
                     •Bralan_BLAG06000532_2 | OG0 |
                                Hsap ENST00000358337 | OG0 |
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                         Cscu XM 023362696.1 | OG0 |
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                         Skow XM 002739975.2 | OG0 |
                            •Spur XM_030980808.1 | OG0 |
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                        Cgig XM 011432005.2 | OG0 |
                         Acagra model.g13064.t1 | OG0 |
                      100 Acagra model.g13064.t1.1.5d3b8710 | OG0 |
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                              •Morvir_scaffold471.g4.t1 | OG0 |
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                            Hvul_g10336_1 | OG0 |
                              •Sanmal Sma 011135-T1 | OG0 |
                           Aaur_scaffold120.g15.t1 | OG0 |
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                            •Nemnom_g7335.t1 | OG0 |
                            •Rhoesc_Res_000351-T1 | OG0 |
                      Exapal_XM_028659038.1 | OG0 |
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                        •Dgig XM 028545862.1 | OG0 |
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                       Amil_Amillepora10581-RA | OG0 |
                     <sup>100</sup><sup>1</sup>-Adig XM 015895749.1 | OG0 |
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                       •Gasp_gasp1.m3.9289.m1 | OG0 |
                        •Spis_XP_022781098_1 | OG0 |
                           •Pocdam XM 027201124.1 | OG0 |
                         •Plespi Pspi scaffold6324 01 | OG0 |
                            -Xestes maker-XT scaffold4755-augustus-gene-0.8-mRNA-1 | OG0 |
                                -Aque_Aqu2.1.30485_001 | OG0 |
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                              •Tetwil_g10984.t2 | OG0 |
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                            •Emue_Em0022g216a.t1 | OG0 |
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                               Spolac_c96500_g1_i3_m.13192c96500-g1 | OG0 |
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                                <sub>s</sub>Spolac_c96500_g1_i1_m.13189c96500-g1 | OG0 |
                               83<sup>1</sup>-Spolac_c96500_g1_i2_m.13190c96500-g1 | OG0 |
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                               •Clacor Ccor_C536099_01 | OG0 |
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                100
                          <sup>100</sup> Lcom_lcpid105733 | OG0 |
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                                Gracom_Gcom_scaffold10138_01 | OG0 |
                                -Scil_scict000075.1 | OG0 |
                                 Leuniv Lniv scaffold15587 01 | OG0 |
                                Horcal_Hcv1.av93.c2.g1409.i1 | OG0 |
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                                   •Pbac_sb2665549 | OG0 |
                                <sup>100</sup> Pbac sb2642825 | OG0 |
                                      •Vmul_sbMO394691 | OG0 |
                                          -Eupdun_Edun_sb10635626 | OG0 |
                                     •Coeast_Cast_sb791151 | OG0 |
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                                    <sup>L</sup>-Coeast_Cast_sb791150 | OG0 |
                                       <sub>_</sub>Coeast_Cast_sb791147 | OG0 |
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                                     <sup>961</sup>-Coemet Cmet_scaffold21925_01 | OG0 |
                                       Coeast Cast sb791148 | OG0 |
                                    98 Coeast_Cast_sb791149 | OG0 |
                                   •Lampan_Lpan_7731_01 | OG0 |
                                    <sub>!</sub>Bolinf_Binf_sb1213135 | OG0 |
                                 <sup>100</sup> Mertsp_Mesp_sb554685 | OG0 |
                                       -Mlei ML0739.g7.i1 | OG0 |
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                                         Bolinf_Binf_sb1220904 | OG0 |
                                      <sup>100</sup> Bolinf_Binf_sb1220905 | OG0 |
                                         <sub>[</sub>•Bolinf_Binf_sb1191123 | OG0 |
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                                      <sup>1001</sup>•Beraby_Baby_sb337373 | OG0 |
                                          Drygla_Dgla_sb300545 | OG0 |
                                           Drygla_Dgla_sb300547 | OG0 |
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                                            Drygla_Dgla_sb300548 | OG0 |
                                            <sup>71</sup> Drygla_Dgla_sb300546 | OG0 |
0.416667
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