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•Tetwil_g23951.t1 | OG0 |
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                                               •Stycar maker-SC scaffold137857-snap-gene-0.23-mRNA-1 | OG0 |
                   Xestes maker-XT scaffold4896-augustus-gene-0.38-mRNA-1 | OG0 |
  99
                    •Aque Aqu2.1.32105 001 | OG0 |
                    •Halamb c30230 g1 i1m.6196 | OG0 |
                    •Halamb_c248_g1_i1m.28 | OG0 |
     62
                              Emue Em0021g309a.t1 | OG0 |
                         <sup>100</sup><sup>1</sup> - Spolac_c96616_g1_i2_m.13411c96616-g1 | OG0 |
                        •Emue_Em0021g308a.t1 | OG0 |
                <sup>99</sup> Spolac_c104471_g5_i1_m.40782c104471-g5 | OG0 |
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                •Cgig XM 011450457.3 | OG0 |
                      -Bralan_BLAG18000191_1 | OG0 |
                      •Cscu_XM_023380420.1 | OG0 |
                Acagra_model.g3248.t1 | OG0 |
        68
             <sup>100</sup> Acagra_model.g3248.t1.1.5d3b824d | OG0 |
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                      •Skow_XM_002736989.2 | OG0 |
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                           •Ctel gnl WGS AMQN CAPTEDRAFT mRNA163376 | OG0 |
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                         <sub>!</sub>Opea_EP00118_Oscarella_pearsei_P003197 | OG0 |
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                         <sup>94</sup> Gracom Gcom scaffold15856 01 | OG0 |
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                            -Scil_scict004913.1 | OG0 |
                                         •Dpul mRNA20108 | OG0 |
                          •Dgig XM 028553387.1 | OG0 |
                    <sup>100</sup> -Xesp 008403-T1 | OG0 |
                                            •Codhol_EP00042_Codosiga_hollandica_P038503 | OG0 |
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                             <sup>97</sup> Pocdam_XM_027198565.1 | OG0 |
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                                 <sup>⁻</sup>-Gasp_gasp1.m3.2668.m1 | OG0 |
                                  •Gfas_gfas1.m1.16774.m1 | OG0 |
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                           ·Salkve_EP00050_Salpingoeca_kvevrii_P017949 | OG0 |
                               •Mylflu_EP00047_Mylnosiga_fluctuans_P002041 | OG0 |
                            ·Salurc_EP00051_Salpingoeca_urceolata_P026916 | OG0 |
                               Lcom_lcpid119301 | OG0 |
                                  Saldol EP00054 Salpingoeca dolichothecata P031968 | OG0 |
                                         Diagra EP00040 Diaphanoeca grandis P029435 | OG0 |
                                      •Acaspe_EP00035_Acanthoeca_spectabilis_P035717 | OG0 |
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                                 <sup>100</sup><sup>1</sup>-Helnan_EP00037_Helgoeca_nana_P032334 | OG0 |
                                             •Mbre_EDQ84003 | OG0 |
                                  •Didcos_EP00039_Didymoeca_costata_P006208 | OG0 |
                                 Salmac_EP00052_Salpingoeca_macrocollata_P030725 | OG0 |
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                                 ·Salinf_EP00049_Salpingoeca_infusionum_P017915 | OG0 |
                                       •Microa EP00043 Microstomoeca roanoka P019128 | OG0 |
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                                         Sros EGD77802 | OG0 |
                                    Choper_EP00045_Choanoeca_perplexa_P002121 | OG0 |
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                              98
                                          •Chofle_EP00730_Choanoeca_flexa_P013071 | OG0 |
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                             TrH2_TrispH2_005535-RA | OG0 |
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                          <sup>100</sup> Tadh_TriadT61818 | OG0 |
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                                -HoiH06_mRNA2420 | OG0 |
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                             <sup>100</sup> Mertsp_Mesp_sb556961 | OG0 |
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                                    •Coemet Cmet scaffold22141 01 | OG0 |
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                                       Pbac_sb2658571 | OG0 |
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