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Vmul sbMO455722 | OG0 |
       <sup>100</sup> Vmul_Simscaffold16612_01 | OG0 |
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                Coeast Cast sb794117 | OG0 |
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              <sup>92</sup> Coeast Cast_sb794118 | OG0 |
                Coeast Cast sb794120 | OG0 |
                -Coeast_Cast_sb794119 | OG0 |
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            Lampan_Lpan_14946_01 | OG0 |
            Mertsp Mesp sb599783 | OG0 |
               Bolinf Binf sb1275063 | OG0 |
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            <sup>100</sup> Bolinf_Binf_sb1275064 | OG0 |
             Beraby_Baby_sb351143 | OG0 |
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               Beraby_Baby_sb351142 | OG0 |
             <sup>87</sup> Beraby_Baby_sb351145 | OG0 |
              Vmul sbMO427114 | OG0 |
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                •Mlei ML0332.g37.i1 | OG0 |
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                 •Drygla_Dgla_sb305104 | OG0 |
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                     Drygla Dgla sb305107 | OG0 |
                    99 Drygla_Dgla_sb305106 | OG0 |
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                     Leuniv Lniv scaffold23896 01 | OG0 |
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                             Scil_scict008490.2 | OG0 |
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                                •Lcom_lcpid242651 | OG0 |
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      59
                              •Helnan_EP00037_Helgoeca_nana_P029143 | OG0 |
                             •Acaspe_EP00035_Acanthoeca_spectabilis_P020551 | OG0 |
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                         Salkve_EP00050_Salpingoeca_kvevrii_P015278 | OG0 |
                              Saldol EP00054 Salpingoeca dolichothecata P009129 | OG0 |
                            Codhol EP00042 Codosiga hollandica P052093 | OG0 |
            68
                               -Salurc_EP00051_Salpingoeca_urceolata_P028061 | OG0 |
                               •Mylflu_EP00047_Mylnosiga_fluctuans_P020508 | OG0 |
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                                 ·Salmac_EP00052_Salpingoeca_macrocollata_P009268 | OG0 |
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                                  Salpun EP00053 Salpingoeca punica P012179 | OG0 |
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                                Salinf EP00049 Salpingoeca infusionum P005541 | OG0 |
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                                Sros EGD73791 | OG0 |
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                                Mbre_EDQ91921 | OG0 |
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                               Chofle EP00730 Choanoeca flexa P003635 | OG0 |
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                    •Plajan_Pjan_C572052_01 | OG0 |
                    <sub>•</sub>TrH2_TrispH2_004843-RA | OG0 |
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                 <sup>1001</sup>Tadh_TriadT54976 | OG0 |
                     •HoiH11 mRNA9581 | OG0 |
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                         •HoiH23 PIH23 001298-RA | OG0 |
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                     <sup>1001</sup>•HoiH06 mRNA7088 | OG0 |
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                        <sub>e</sub>Hhon_g08634.t1 | OG0 |
                           HoiH04 mRNA14849 | OG0 |
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                        <sup>100</sup> HoiH04 mRNA14728 | OG0 |
                       Plespi_Pspi_scaffold2450 01 | OG0 |
                        Plespi_Pspi_scaffold2453_01 | OG0 |
                         Plespi_Pspi_scaffold2454_01 | OG0 |
                        <sup>11</sup>Plespi Pspi scaffold2451 01 | OG0 |
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                    •Tetwil g27138.t1 | OG0 |
                       Xestes maker-XT scaffold1492-augustus-gene-0.9-mRNA-1 | OG0 |
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                         -Aque_Aqu2.1.34594_001 | OG0 |
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                        •Tetwil g27141.t2 | OG0 |
                         Spolac c104653 g1 i2 m.42145c104653-g1 | OG0 |
                          <u>r</u>Emue_Em0014g773a.t1 | OG0 |
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                       <sup>100</sup>Lemue Em0014g784a.t1 | OG0 |
                        •Hsap_ENST00000366923 | OG0 |
                      Bralan_BLAG04102214_1 | OG0 |
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                       •Skow_XM_006812442.1 | OG0 |
                         Spur_XM_030998457.1 | OG0 |
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                           Cscu XM 023371128.1 | OG0 |
                            •Tcas TC033636 001 | OG0 |
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                         •Cgig_XM_034469491.1 | OG0 |
                         Acagra model.g15810.t1 | OG0 |
                      <sup>100</sup> Acagra model.g15810.t1.1.5d3b882d | OG0 |
        75
                            •Chem_TCONS_00058467 | OG0 |
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                       <sup>100</sup> Hvul_g13987_1 | OG0 |
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                      Sanmal_Sma_023821-T1 | OG0 |
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                         •Aaur_scaffold144.g9.t3 | OG0 |
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                            Nemnom_g9688.t1 | OG0 |
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                            -Rhoesc_Res_004475-T1 | OG0 |
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                         -Adig XM 015901526.1 | OG0 |
                      <sup>100</sup> Amil Amillepora27477-RA | OG0 |
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                              •Gasp_gasp1.m3.6784.m1 | OG0 |
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