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-Lampan_Lpan_16257 01 | OG0 |
                              -Bolinf Binf sb1219004 | OG0 |
                          <sup>100</sup><sup>L</sup>•Mlei ML1169.g17.i1 | OG0 |
                                  •Drygla_Dgla_sb278924 | OG0 |
                        96
                                     •Pbac sb2665225 | OG0 |
                                   Horcal Hcv1.av93.c6.g585.i1 | OG0 |
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                                 <sup>79</sup> Pbac sb2665227 | OG0 |
                             Vmul_Simscaffold15231_01 | OG0 |
                                100
                             <sup>1001</sup>Coemet Cmet_scaffold759_01 | OG0 |
                                 Coeast Cast sb777155 | OG0 |
                          100
                                 Coeast Cast sb777157 | OG0 |
                                 Coeast Cast sb777156 | OG0 |
 83
                               <sup>931</sup>Coeast_Cast_sb777158 | OG0 |
                         Plajan Pjan C572080 01 | OG0 |
                          Opea EP00118 Oscarella pearsei P007211 | OG0 |
            100
                       100 Ocar g8429 | OG0 |
                                    •Mylflu_EP00047_Mylnosiga_fluctuans_P018586 | OG0 |
                                               Salurc_EP00051_Salpingoeca_urceolata_P024083 | OG0 |
                                           <sup>100</sup> Salurc EP00051 Salpingoeca_urceolata_P024084 | OG0 |
              50
   84
                                               Codhol EP00042 Codosiga hollandica P058899 | OG0 |
                61
                                               •Mbre EDQ91249 | OG0 |
                                                 •Choper_EP00045_Choanoeca_perplexa_P020906 | OG0 |
                                    100
                                                Chofle_EP00730_Choanoeca_flexa_P017091 | OG0 |
                         99
                                                Salinf EP00049 Salpingoeca infusionum P002629 | OG0 |
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                                                   •Microa EP00043 Microstomoeca roanoka P006887 | OG0 |
                           60
                                           100
                                                   •Sros EGD80612 | OG0 |
                                  Salmac_EP00052_Salpingoeca_macrocollata_P014461 | OG0 |
                                     Salmac EP00052_Salpingoeca_macrocollata_P014462 | OG0 |
                    92
                                             Salpun EP00053 Salpingoeca punica P005089 | OG0 |
                                Saldol EP00054 Salpingoeca dolichothecata P025753 | OG0 |
                  83
                                      Codhol EP00042 Codosiga hollandica P058898 | OG0 |
                                    Salkve EP00050 Salpingoeca kvevrii P021408 | OG0 |
                78
                                               Diagra EP00040 Diaphanoeca grandis P020608 | OG0 |
                  84
                                                 Acaspe EP00035 Acanthoeca spectabilis P010662 | OG0 |
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                                                    •Helnan EP00037 Helgoeca nana P024691 | OG0 |
                                  100
                                             Stedip_EP00041_Stephanoeca_diplocostata_P037459 | OG0 |
                                               -Stedip_EP00041_Stephanoeca_diplocostata_P037460 | OG0 |
                       •Plespi_Pspi_scaffold6309_01 | OG0 |
                           •Emue Em0010g935a.t1 | OG0 |
                          •Spolac c89420 g1 i1 m.5639c89420-g1 | OG0 |
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                          •Tetwil_g26272.t1 | OG0 |
                 86
                       Stycar_maker-SC_scaffold2878-augustus-gene-0.19-mRNA-1 | OG0 |
           99
                                Halamb_c54650_g2_i2mm.6052 | OG0 |
                             <sup>100</sup>lHalamb_c54255_g1_i1mm.5785 | OG0 |
                              Aque Aqu2.1.08819 001 | OG0 |
                       100
                                 Aque_Aqu2.1.03268_001 | OG0 |
                           98
                                   Aque_Aqu2.1.17973_001 | OG0 |
                                  <sup>75</sup> Aque_Aqu2.1.02479_001 | OG0 |
                              •Cscu XM 023353662.1 | OG0 |
                      •Bralan BLAG17000184 1 | OG0 |
        87
            93
                          Spur_XM_031000529.1 | OG0 |
                           •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA225863 | OG0 |
          90
                         -Acagra_model.g11274.t1.1.5d3b865a_model.g11274.t1_model.g11275.t1 | OG0 |
     32
                          Cgig XM 034475100.1 | OG0 |
                                                  •Tcas TC032187 001 | OG0 |
                                 -•Tadh_TriadT19228 | OG0 |
                              <sup>100</sup> TrH2_TrispH2_001032-RA | OG0 |
                                  •HoiH11_mRNA8567 | OG0 |
                                       <sub>-</sub>HoiH23_PIH23_008578-RA | OG0 |
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                                    <sup>100</sup><sup>1</sup>•HoiH06_mRNA7544 | OG0 |
                                       •Hhon_g02888.t1 | OG0 |
                                    <sup>100]</sup>•HoiH04_mRNA1512 | OG0 |
                        •Morvir_scaffold293.g11.t1 | OG0 |
                             •Chem TCONS 00017114 | OG0 |
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                    100
                                 •Hvul_g30520_1 | OG0 |
                          Sanmal Sma 026215-T1 | OG0 |
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                             •Aaur scaffold1476.g4.t1 | OG0 |
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                                   •Aaur_scaffold1476.g3.t1 | OG0 |
                     100
                        100
                                Nemnom q4369.t1 | OG0 |
      97
                           <sup>100</sup> Rhoesc_Res_013092-T1 | OG0 |
                         •Xesp_020337-T1 | OG0 |
                          Dgig_XM_028561068.1 | OG0 |
                       <sup>100</sup> Dgig_XM_028546582.1 | OG0 |
                     Exapal XM 021044886.2 | OG0 |
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             100
                   •Nvec v1g89190 | OG0 |
                      •Gfas gfas1.m1.10849.m1 | OG0 |
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                       Amil_Amillepora04486-RA | OG0 |
                    100 Adig_XM_015911079.1 | OG0 |
                        -Spis_XP_022783585_1 | OG0 |
               100
                     <sup>100</sup><sup>1</sup>-Pocdam_XM_027187111.1 | OG0 |
                        -Fspp_ffun1.m4.3101.m1 | OG0 |
                     <sup>100</sup>└-Gasp_gasp1.m3.24741.m1 | OG0 |
0.416667
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