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-Saldol_EP00054_Salpingoeca_dolichothecata_P017623 | OG0 |
                              •Codhol_EP00042_Codosiga_hollandica_P050821 | OG0 |
                                                       -Pbac sb2648356 | OG0 |
 81
                               Salmac_EP00052_Salpingoeca_macrocollata_P028603 | OG0 |
                            ·Salurc_EP00051_Salpingoeca_urceolata_P031977 | OG0 |
   87
                                   •Salpun EP00053_Salpingoeca_punica_P010527 | OG0 |
     86
                                         •Microa_EP00043_Microstomoeca_roanoka_P002475 | OG0 |
                          100
                                         •Sros_EGD83695 | OG0 |
        91
                                          Salmac_EP00052_Salpingoeca_macrocollata_P028601 | OG0 |
                                   Mylflu EP00047 Mylnosiga fluctuans P023726 | OG0 |
                                         •Salkve EP00050_Salpingoeca_kvevrii_P015659 | OG0 |
                             -Pbac sb2659553 | OG1 |
                         <sup>100</sup> Horcal_Hcv1.av93.c12.g300.i1 | OG1 |
                      100
                            •Pbac_sb2657583 | OG1 |
                          <sup>100</sup>Ū-Pbac sb2660847 | OG1 |
                           Mertsp Mesp sb542199 | OG1 |
                                    -Eupdun Edun sb10659003 | OG1 |
                             Lampan Lpan 1046 01 | OG1 |
                          <sup>1001</sup>Lampan_Lpan_1045_01 | OG1 |
                                 •Eupdun_Edun_sb10659002 | OG1 |
                                     Vmul sbMO436046 | OG1 |
               100
                       49
                                  ·Vmul_sbMO436043 | OG1 |
                          100
                                   •Coemet_Cmet_scaffold22436_01 | OG1 |
                                      <sub>[</sub>Coeast_Cast_sb805424 | OG1 |
                                    <sup>97]</sup>•Coeast Cast sb805425 | OG1 |
                               100
                                         Coeast Cast sb805426 | OG1 |
                                         -Coeast_Cast_sb805427 | OG1 |
                                           "Coeast_Cast_sb805420 | OG1 |
                  45
                                         <sup>981</sup>Coeast_Cast_sb805421 | OG1 |
                                           Coeast_Cast_sb805423 | OG1 |
                                         <sup>75]</sup>Coeast_Cast_sb805422 | OG1 |
                             Bolinf_Binf_sb1225783 | OG1 |
                           <sup>991</sup>Bolinf_Binf_sb1225784 | OG1 |
                              Mertsp Mesp sb584133 | OG1 |
                           <sup>100</sup> Mertsp_Mesp_sb584134 | OG1 |
                                Drygla_Dgla_sb306666 | OG1 |
                                     Drygla_Dgla_sb306662 | OG1 |
                     79
                             99
                                 <sup>100</sup>Drygla Dgla sb306664 | OG1 |
                                     Drygla_Dgla_sb306663 | OG1 |
                                  <sup>100</sup>lDrygla_Dgla_sb306665 | OG1 |
                       99
                                Beraby_Baby_sb347928 | OG1 |
                             <sup>100</sup>•Beraby_Baby_sb347929 | OG1 |
                                 Bolinf_Binf_sb1253482 | OG1 |
                         99
                                   <sub>_</sub>Bolinf_Binf_sb1253483 | OG1 |
                                 <sup>99</sup> Mertsp_Mesp_sb573807 | OG1 |
                                    Bolinf_Binf_sb1217643 | OG1 |
                           100
                                 <sup>100</sup> Mertsp_Mesp_sb539388 | OG1 |
                                     Mlei_ML1658.g1.i1 | OG1 |
                                        •Vmul_sbMO427127 | OG1 |
                                           *Vmul sbMO362604 | OG1 |
                                        <sup>100</sup> Mlei_ML0590.g1.i1 | OG1 |
                           •Plespi_Pspi_scaffold15310_01 | OG1 |
                               -Xestes_snap_masked-XT_scaffold57209-processed-gene-0.2-mRNA-1 | OG1 |
                                 <sub>•</sub>Halamb_c53470_g2_i1mm.5220 | OG1 |
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        99
                              <sup>1001</sup>-Halamb_c54376_g1_i1m.38112 | OG1 |
                                  Aque Aqu2.1.20472_001 | OG1 |
                               <sup>1001</sup>•Aque_Aqu2.1.27567_001 | OG1 |
                             •Tetwil g19300.t1 | OG1 |
                               •Tetwil_g19303.t1 | OG1 |
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                    99
                              Stycar_maker-SC_scaffold6422-snap-gene-0.5-mRNA-1 | OG1 |
                                  •Emue Em0005g262a.t1 | OG1 |
                                 Spolac c103637 g1 i1 m.35303c103637-g1 | OG1 |
                           100
                               Spoiac_c103637_g1_i2_m.35304c103637-g1 | OG1 |
   44
                                                           -Amil_Amillepora20438-RA | OG1 |
                              <sub>]</sub>TrH2_TrispH2_009098-RA | OG1 |
                           Tadh TriadT56039 | OG1 |
                                  •HoiH23 PIH23 008204-RA | OG1 |
                         100
                               <sup>100</sup> -HoiH06_mRNA7742 | OG1 |
                                  Hhon_g01842.t1 | OG1 |
                               1001HoiH04 mRNA4928 | OG1 |
          92
                                  •Plajan Pjan scaffold23292 01 | OG1 |
                                   Opea_EP00118_Oscarella_pearsei_P004101 | OG1 |
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     66
                                <sup>100</sup>  Ocar_g3722 | OG1 |
                                  Leuniv_Lniv_C760374_01 | OG1 |
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                                    .com_lcpid23745 | OG1 |
                         100
                                  Leuniv Lniv C813946 01 | OG1 |
                                     Gracom_Gcom_C561062_01 | OG1 |
                                    •Scil scict024645.1 | OG1 |
                               Bralan BLAG02001334_1 | OG1 |
                           Hsap ENST00000620755 | OG1 |
        76
                             Hsap ENST00000381474 | OG1 |
                           99 Hsap_ENST00000263012 | OG1 |
                             •Skow_XM_006814047.1 | OG1 |
             93
                  99
                              Spur XM 030977562.1 | OG1 |
                                •Acagra_model.g7786.t1 | OG1 |
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                                 -Cgig_XM_011424286.3 | OG1 |
                 95
                                   •Cscu_XM_023386542.1 | OG1 |
                                            Dpul mRNA22395 | OG1 |
                                         <sup>100</sup> - Dpul_mRNA22402 | OG1 |
                         98
          73
                                                -Tcas_TC004746_001 | OG1 |
                                97
                                                          -Lcom_lcpid147454 | OG1 |
                                   •Hvul_g28188_1 | OG1 |
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                                •Chem TCONS 00016435 | OG1 |
                               •Morvir_scaffold2.g161.t1 | OG1 |
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                               Sanmal Sma 016858-T1 | OG1 |
                                     Aaur_scaffold469.g8.t1 | OG1 |
                       100
                                  •Nemnom_g699.t1 | OG1 |
                          100
                             <sup>100</sup><sup>L</sup>•Rhoesc_Res_012842-T1 | OG1 |
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                             Dgig_XM_028545376.1 | OG1 |
                       100
                              •Xesp_015751-T1 | OG1 |
                              •Nvec_NVE15349 | OG1 |
                     100
                            •Exapal_XM_028659648.1 | OG1 |
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                                •Exapal_XM_028659679.1 | OG1 |
                                <sub>!</sub>Nvec_v1g13339 | OG1 |
                       100
                  100
                                 -Nvec_NVE15350 | OG1 |
                                 •Gasp_gasp1.m3.15934.m1 | OG1 |
                              <sup>97</sup>-Fspp_ffun1.m4.369.m1 | OG1 |
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                                 •Spis_XP_022788925_1 | OG1 |
                                    <sub>-</sub>Pocdam_XM_027191501.1 | OG1 |
                                 <sup>100</sup> Spis XP 022788943_1 | OG1 |
                        100
                                •Gfas_gfas1.m1.12645.m1 | OG1 |
                                  •Adig_XM_015919500.1 | OG1 |
                                     •Amil_Amillepora37684-RA | OG1 |
                                       Adig XM 015919499.1 | OG1 |
                                    <sup>1001</sup>•Amil Amillepora37683-RA | OG1 |
0.416667
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