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Saldol EP00054 Salpingoeca dolichothecata P004205 | OG0 |
                  •Didcos_EP00039_Didymoeca_costata_P001350 | OG0 |
              100 Didcos_EP00039_Didymoeca_costata_P001351 | OG0 |
                  Diagra_EP00040_Diaphanoeca_grandis_P005996 | OG0 |
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 100
                     Stedip EP00041 Stephanoeca diplocostata P023523 | OG0 |
          96
                     Acaspe EP00035 Acanthoeca spectabilis P011651 | OG0 |
                    •Helnan_EP00037_Helgoeca_nana_P006132 | OG0 |
    97
                  Salmac EP00052 Salpingoeca macrocollata P003972 | OG0 |
        98
                    -Salpun_EP00053_Salpingoeca_punica_P023487 | OG0 |
                         -Salkve_EP00050_Salpingoeca_kvevrii P009759 | OG0 |
                       Codhol EP00042 Codosiga hollandica P045840 | OG0 |
      86
                                        •Codhol_EP00042_Codosiga_hollandica_P045838 | OG0 |
                       100 Codhol_EP00042_Codosiga_hollandica_P045839 | OG0 |
        35
                     ·Salurc_EP00051_Salpingoeca_urceolata_P022934 | OG0 |
                           Mylflu EP00047 Mylnosiga fluctuans P005626 | OG0 |
          41
                                            -Lcom | lcpid244760 | OG0 |
            35
                                 Salinf EP00049 Salpingoeca infusionum P001226 | OG0 |
                             •Microa_EP00043_Microstomoeca_roanoka_P026840 | OG0 |
              92
                   98
                             •Sros_EGD77734 | OG0 |
                81
                               Mbre EDQ87922 | OG0 |
                                Chofle EP00730 Choanoeca flexa P003459 | OG0 |
                                -Choper_EP00045_Choanoeca_perplexa_P005896 | OG0 |
                  •Plespi_Pspi_C608950_01 | OG1 |
                •Tetwil_g11943.t1 | OG1 |
              •Stycar_maker-SC_scaffold5989-augustus-gene-0.28-mRNA-1 | OG1 |
                Spolac c98512 g1 i1 m.17392c98512-g1 | OG1 |
    100
                  Emue Em0016g935a.t1 | OG1 |
               <sup>100</sup> Emue_Em0582g7a.t1 | OG1 |
                   -Halamb_c55018_g1_i1m.39791 | OG1 |
       69
                    Aque Aqu2.1.12802_001 | OG1 |
                  -Aque Aqu2.1.04151 001 | OG1 |
            100
                     •Aque_Aqu2.1.01006_001 | OG1 |
                 <sup>97</sup> Aque_Aqu2.1.16125_001 | OG1 |
                  •Morvir scaffold895.g1.t1 | OG1 |
                       •Hvul_g13225_1 | OG1 |
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                      Chem TCONS 00071212 | OG1 |
                      •Morvir_scaffold315.g20.t1 | OG1 |
          92
                      Sanmal_Sma_021601-T1 | OG1 |
                   <sup>100]</sup>Sanmal_Sma_021602-T1 | OG1 |
 100
                      •Aaur scaffold347.g14.t1 | OG1 |
               100
      95
                         •Nemnom g18705.t1 | OG1 |
                     <sup>100</sup><sup>1</sup>-Rhoesc_Res_015335-T1 | OG1 |
                 •Dgig_XM_028555163.1 | OG1 |
                  •Xesp_008066-T1 | OG1 |
                    •Exapal_XM_021050984.1 | OG1 |
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                    •Nvec v1g241701 | OG1 |
                    •Spis_XP_022779280_1 | OG1 |
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                 <sup>1001</sup>•Pocdam_XM_027196948.1 | OG1 |
                        Fspp_ffun1.m4.4261.m1 | OG1 |
              100
                      Gasp_gasp1.m3.13647.m1 | OG1 |
                       Gfas gfas1.m1.18912.m1 | OG1 |
                         •Amil_Amillepora15393-RA | OG1 |
                          •Adig_XM_015918724.1 | OG1 |
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                            Adig_XM_015893575.1 | OG1 |
                               •Amil Amillepora15401-RA | OG1 |
                                 Amil Amillepora15397-RA | OG1 |
                                  -Adig XM 015918731.1 | OG1 |
                    •Clacor_Ccor_scaffold1129_01 | OG1 |
                     •Lcom_lcpid74798 | OG1 |
                       Scil scict011077.1 | OG1 |
                          Leuniv Lniv scaffold17052 01 | OG1 |
                          •Gracom Gcom scaffold12534 01 | OG1 |
                    •Plajan_Pjan_scaffold17452_01 | OG1 |
                 <sup>100</sup>Ū-•Plajan_Pjan_scaffold17453_01 | OG1 |
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                    Opea_EP00118_Oscarella_pearsei_P007230 | OG1 |
                              Opea EP00118 Oscarella pearsei P007229 | OG1 |
                    <sup>58</sup> Ocar g6650 | OG1 |
                         -Spur_XM_030997867.1 | OG1 |
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                        Skow XM 006825716.1 | OG1 |
                           •Hsap ENST00000374199 | OG1 |
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                        Bralan BLAG08000356 1 | OG1 |
                           Cscu_XM_023356375.1 | OG1 |
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                              •Tcas_TC000271_001 | OG1 |
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                      100
                                 •Dpul_mRNA312 | OG1 |
                             Cgig XM 011450792.3 | OG1 |
                  97
                      100
                                    -Acagra model.g20932.t1 | OG1 |
                                          •Acagra_model.g20934.t1 | OG1 |
                              Ctel gnl WGS AMQN CAPTEDRAFT mRNA226300 | OG1 |
                            1001 Ctel _gnl_WGS_AMQN_CAPTEDRAFT_mRNA166413 | OG1 |
          76
                            -TrH2_TrispH2 003014-RA | OG1 |
                          <sup>99</sup>Ū-Tadh TriadT21736 | OG1 |
                              •HoiH11 mRNA3938 | OG1 |
                       100
                                   -HoiH23_PIH23_000056-RA | OG1 |
                                <sup>[100]</sup>-HoiH06_mRNA6769 | OG1 |
                             100
                                   Hhon g03747.t1 | OG1 |
                                <sup>1001</sup>HoiH04 mRNA14897 | OG1 |
                                 -Lampan_Lpan_13437_01 | OG1 |
                                    •Vmul_sbMO435135 | OG1 |
            70
                             99
                                        Coemet_Cmet_scaffold13746_01 | OG1 |
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                                     <sup>100</sup> Coemet Cmet_scaffold13747_01 | OG1 |
                                           Coeast Cast sb787778 | OG1 |
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                                        <sup>100</sup> Coeast Cast sb787780 | OG1 |
                                          Coeast_Cast_sb787779 | OG1 |
                                        99 Coeast_Cast_sb787777 | OG1 |
                          100
                                  •Vmul sbMO404441 | OG1 |
                                      Drygla Dgla sb289450 | OG1 |
                               53
                                        Pbac sb2669009 | OG1 |
                                 58
                                            Pbac_sb2657445 | OG1 |
                                          <sup>93]</sup>Horcal _Hcv1.av93.c8.g356.i1 | OG1 |
                                            Pbac sb2640596 | OG1 |
                                          <sup>981</sup>-Pbac_sb2657786 | OG1 |
                                    Beraby_Baby_sb338910 | OG1 |
                                    Beraby_Baby_sb338909 | OG1 |
                                    <sup>1</sup>•Beraby_Baby_sb338907 | OG1 |
                                      Beraby_Baby_sb338912 | OG1 |
                               100
                                    | Beraby Baby sb338911 | OG1 |
                                      1-Beraby_Baby_sb338908 | OG1 |
                                      Beraby_Baby_sb338915 | OG1 |
                                    <sup>11</sup>Beraby Baby sb338917 | OG1 |
0.416667
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