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Salinf EP00049 Salpingoeca infusionum P004592 | OG0 |
               •Microa_EP00043_Microstomoeca_roanoka_P017935 | OG0 |
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                                                                 -Adig_XM_015893939.1 | OG0 |
                          •Mbre EDQ88298 | OG0 |
                            Choper EP00045 Choanoeca perplexa P013353 | OG0 |
                               •Chofle_EP00730_Choanoeca_flexa_P015177 | OG0 |
           Salpun EP00053 Salpingoeca punica P018060 | OG1 |
                     •Didcos_EP00039_Didymoeca_costata_P008029 | OG1 |
                         Diagra EP00040 Diaphanoeca grandis P031773 | OG1 |
                       Diagra EP00040 Diaphanoeca grandis P031772 | OG1 |
                         Stedip EP00041 Stephanoeca diplocostata P035693 | OG1 |
                              -Diagra EP00040 Diaphanoeca grandis P031774 | OG1 |
                            Helnan EP00037 Helgoeca nana P001736 | OG1 |
                            Acaspe EP00035 Acanthoeca spectabilis P034609 | OG1 |
                 Salurc EP00051 Salpingoeca urceolata P019401 | OG2 |
                 •Salkve_EP00050_Salpingoeca_kvevrii_P008097 | OG2 |
                   •Codhol_EP00042_Codosiga_hollandica_P037017 | OG2 |
                    Codhol EP00042 Codosiga hollandica P037018 | OG2 |
                     Codhol EP00042 Codosiga hollandica P055003 | OG2 |
                    Codhol EP00042 Codosiga hollandica P055001 | OG2 |
                    •Mylflu_EP00047_Mylnosiga_fluctuans_P020709 | OG2 |
                     Salpun EP00053 Salpingoeca punica P017748 | OG2 |
                      Salmac EP00052 Salpingoeca macrocollata P024111 | OG2 |
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                    Salpun EP00053 Salpingoeca punica P017746 | OG2 |
                  Saldol_EP00054_Salpingoeca_dolichothecata_P029611 | OG2 |
                                                       Tetwil g6029.t1 | OG2 |
                                                            Stycar_maker-SC_scaffold137542-snap-gene-0.7-mRNA-1 | OG2 |
                                                             •Stycar maker-SC scaffold16866-snap-gene-0.24-mRNA-1 | OG2 |
                                                                •Pbac sb2645430 | OG2 |
                                Eupdun_Edun sb10609234 | OG2 |
                                                                   •Pbac sb2652622 | OG2 |
                                  Eupdun Edun sb10633704 | OG2 |
                                     -Bolinf_Binf_sb1183726 | OG2 |
                                               -Pbac sb2639343 | OG2 |
                                        Horcal_Hcv1.av93.c4.g428.i1 | OG2 |
                                         Horcal Hcv1.av93.c4.g428.i2 | OG2 |
                                                        •Pbac sb2652625 | OG2 |
                                       Drygla Dgla sb290313 | OG2 |
                                      📶 Drygla Dgla sb290311 | OG2 |
                                        <sup>1</sup>•Drygla_Dgla_sb290312 | OG2 |
                                           Drygla Dgla sb290308 | OG2 |
                                          •Mlei_ML0651.g9.i1 | OG2 |
                                                       •Mlei ML1132.g2.i1 | OG2 |
                                          Beraby Baby sb339141 | OG2 |
                                             Beraby_Baby_sb339137 | OG2 |
                                            <sup>11</sup>Beraby Baby sb339138 | OG2 |
                                             Beraby Baby sb339142 | OG2 |
                                              Beraby_Baby_sb339139 | OG2 |
                                             <sup>1</sup>Beraby_Baby_sb339140 | OG2 |
                                             Lampan Lpan 7655 01 | OG2 |
                                            📶 <sub>"</sub>Lampan_Lpan_7656_01 | OG2 |
                                             <sup>1</sup>-Lampan_Lpan_7657_01 | OG2 |
                                          •Vmul_sbMO451764 | OG2 |
                                           -Coeast_Cast_sb766612 | OG2 |
                                             •Vmul_sbMO451752 | OG2 |
                                              rVmul_sbMO420346 | OG2 |
                                             1 Vmul sbMO420347 | OG2 |
                                              •Coemet_Cmet_scaffold17324 01 | OG2 |
                                                Coemet Cmet scaffold17322 01 | OG2 |
                                                Coemet_Cmet_scaffold17323_01 | OG2 |
                                                 Coeast_Cast_sb820107 | OG2 |
                                                 Coeast_Cast_sb820106 | OG2 |
                                                   rCoeast_Cast_sb820104 | OG2 |
                                                     Coeast_Cast_sb820108 | OG2 |
                                                      Coeast Cast sb820109 | OG2 |
                                                        Coeast Cast sb820112 | OG2 |
                                                       <sup>11</sup>-Coeast_Cast_sb820120 | OG2 |
                                                      Coeast Cast sb820102 | OG2 |
                                                       Coeast Cast sb820103 | OG2 |
                                                       Coeast Cast sb820105 | OG2 |
                                                      Coeast Cast sb820113 | OG2 |
                                                         rCoeast_Cast_sb820110 | OG2 |
                                                        <sup>1</sup> Coeast_Cast_sb820111 | OG2 |
                                                          Coeast_Cast_sb820115 | OG2 |
                                                         <sup>1</sup> Coeast_Cast_sb820116 | OG2 |
                                                          Coeast Cast sb820117 | OG2 |
                                                          <sup>1]</sup>Coeast Cast sb820118 | OG2 |
                                Morvir scaffold537.g2.t1 | OG2 |
                                  •Hvul_g27135 1 | OG2 |
                                 •Chem_TCONS_00004772 | OG2 |
                                Sanmal Sma 024158-T1 | OG2 |
                                Aaur scaffold17.g29.t1 | OG2 |
                                 -Aaur scaffold17.g27.t1 | OG2 |
                                 •Nemnom_g13268.t1 | OG2 |
                                 -Rhoesc Res 016912-T1 | OG2 |
                                <sup>1</sup><sup>_</sup>Nemnom_g13267.t1 | OG2 |
                               •Xesp_024501-T1 | OG2 |
                              •Dgig_XM_028563351.1 | OG2 |
                             <sup>11</sup>Daia XM 028549048.1 | OG2 |
                              *Xesp 024500-T1 | OG2 |
                              -Xesp 024483-T1 | OG2 |
                              <sup>1</sup>Xesp 024498-T1 | OG2 |
                                •Exapal_XM_028659882.1 | OG2 |
                            •Nvec_v1g183698 | OG2 |
                            -Spis XP 022781710 1 | OG2 |
                           <sup>11</sup>•Pocdam XM 027192007.1 | OG2 |
                            <sub>-</sub>-Gasp_gasp1.m3.300.m1 | OG2 |
                           <sup>1</sup><sup>_</sup>•Fspp_ffun1.m4.13170.m1 | OG2 |
                            •Gfas_gfas1.m1.1739.m1 | OG2 |
                            Amil Amillepora13276-RA | OG2 |
                              -Adig XM 015893937.1 | OG2 |
                               -Adig_XM_015914488.1 | OG2 |
                              •Bralan_BLAG03000634_2 | OG2 |
                              Spur XM 030989304.1 | OG2 |
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                              •Hsap_ENST00000529230 | OG2 |
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                                   •Cscu_XM_023383957.1 | OG2 |
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                                    •Dpul mRNA22301 | OG2 |
                                 •Ctel gnl WGS AMQN CAPTEDRAFT mRNA154948 | OG2 |
                                   •Cgig_XM_034481775.1 | OG2 |
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                                 Acagra_model.g1733.t1.1.5d3b8334 | OG2 |
                                 <sup>11</sup>Acagra model.g1733.t1 | OG2 |
                                TrH2 TrispH2 003154-RA | OG2 |
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                                 •HoiH11_mRNA8608 | OG2 |
                                  <sub>-</sub>HoiH23_PIH23_002115-RA | OG2 |
                                 <sup>1</sup>-HoiH06 mRNA9191 | OG2 |
                                  Hhon g03713.t1 | OG2 |
                                 <sup>1</sup> HoiH04_mRNA16820 | OG2 |
                                 <sub>!</sub>Plajan_Pjan_scaffold4018_01 | OG2 |
                                <sup>1</sup>Plajan_Pjan_scaffold4019_01 | OG2 |
                                    Ocar g3043 | OG2 |
                                   1 Opea EP00118 Oscarella pearsei P028343 | OG2 |
                                      •Clacor Ccor scaffold9979 01 | OG2 |
                                  Lcom | lcpid48281 | OG2 |
                                  Leuniv Lniv C771959 01 | OG2 |
                                     Lcom | lcpid81417 | OG2 |
                                     Leuniv Lniv C744609 01 | OG2 |
                                       •Gracom_Gcom_scaffold18751_01 | OG2 |
                                             Lcom | lcpid46950 | OG2 |
                                      Leuniv Lniv scaffold25971_01 | OG2 |
                                Plespi Pspi scaffold343 01 | OG2 |
                               <sup>1</sup>Plespi Pspi scaffold346 01 | OG2 |
                                             •Tetwil q21226.t2 | OG2 |
                                            11-Tetwil g11458.t1 | OG2 |
                                  Spolac c100223 g1 i1 m.21852c100223-g1 | OG2 |
                                   Spolac c100223 g1 i3 m.21857c100223-g1 | OG2 |
                                        Emue Em0003g63a.t1 | OG2 |
                                  Spolac_c100223_g1_i3_m.21859c100223-g1 | OG2 |
                                   Spolac c100223 g1 i2 m.21855c100223-g1 | OG2 |
                                                 —Emue_Em0018g6a.t1 | OG2 |
                                                   -Emue Em0004g1629a.t1 | OG2 |
                                                 11-Emue Em0188g4a.t1 | OG2 |
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                                  11-Xestes maker-XT scaffold1110-snap-gene-0.29-mRNA-1 | OG2 |
                                    •Halamb c54418 g1 i3m.38234 | OG2 |
                                      -Aque Aqu2.1.36420 001 | OG2 |
                                       •Halamb c49686 g2 i1m.26666 | OG2 |
                                         -Xestes maker-XT scaffold119083-snap-gene-0.6-mRNA-1 | OG2 |
                                    Stycar augustus masked-SC scaffold4752-processed-gene-0.1-mRNA-1 | OG2 |
                                          •Tetwil g6027.t2 | OG2 |
                                                  -Tetwil g20971.t2 | OG2 |
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                                         1 Spolac c103208 g1 i1 m.32965c100223-g1 | OG2 |
                                          <sup>1]</sup>Spolac c103208 g1 i2 m.32967c100223-g1 | OG2 |
0.416667
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