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Bralan BLAG01000592 1 | OG0 |
              -Spur_XM_779894.5 | OG0 |
              •Skow XM 002730971.2 | OG0 |
           •Hsap ENST00000290155 | OG0 |
                       -Hsap ENST00000300258 | OG0 |
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            <sup>100</sup> Hsap ENST00000673807 | OG0 |
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              Ctel gnl WGS AMQN_CAPTEDRAFT_mRNA179325 | OG0 |
              •Cgig XM 011455197.3 | OG0 |
              Acagra model.g4406.t1 | OG0 |
              Cscu XM 023355953.1 | OG0 |
                  •Tcas TC007236 001 | OG0 |
                   Horcal Hcv1.av93.c3.g1064.i1 | OG0 |
     49
                      -Pbac sb2665328 | OG0 |
                100
       53
                   <sup>100</sup> Pbac_sb2666005 | OG0 |
                      -Coeast_Cast_sb771547 | OG0 |
             100
                  <sup>100</sup><sup>1</sup>-Coemet_Cmet_scaffold162_01 | OG0 |
                         Beraby_Baby_sb326311 | OG0 |
                           Lampan Lpan C405435 01 | OG0 |
                89
                            Mlei ML1688.g3.i1 | OG0 |
                         <sup>100</sup> Vmul_sbMO399664 | OG0 |
                            Mertsp Mesp sb564313 | OG0 |
                  86
                              Bolinf_Binf_sb1224487 | OG0 |
                                Bolinf Binf sb1224485 | OG0 |
                              <sup>54</sup> Bolinf Binf sb1224486 | OG0 |
                              Eupdun_Edun_sb10651733 | OG0 |
                           <sup>1001</sup>•Eupdun_Edun_sb10651734 | OG0 |
                              -Drygla_Dgla_sb288776 | OG0 |
                      92
                           <sup>100</sup> Drygla_Dgla_sb288775 | OG0 |
                             Mertsp_Mesp_sb572611 | OG0 |
                              Bolinf Binf sb1231204 | OG0 |
                             <sup>43</sup> Mertsp_Mesp_sb572612 | OG0 |
               Salurc EP00051 Salpingoeca urceolata P026910 | OG0 |
                  Saldol_EP00054_Salpingoeca_dolichothecata_P033065 | OG0 |
                 | Salkve_EP00050_Salpingoeca_kvevrii_P001663
                     •Mylflu_EP00047_Mylnosiga_fluctuans_P004212 | OG0 |
         80
                    •Salpun_EP00053_Salpingoeca_punica_P005807 | OG0 |
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                     Salmac EP00052 Salpingoeca macrocollata P017921 | OG0 |
                        Salmac_EP00052_Salpingoeca_macrocollata_P017920 | OG0 |
                     Mbre_EDQ85604 | OG0 |
                      •Chofle EP00730 Choanoeca flexa P000168 | OG0 |
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                       •Choper_EP00045_Choanoeca_perplexa_P008822 | OG0 |
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                      •Microa EP00043 Microstomoeca roanoka P007761 | OG0 |
                          Salinf EP00049 Salpingoeca infusionum P015941 | OG0 |
                         <sub>'</sub>Salinf_EP00049_Salpingoeca_infusionum_P015940 | OG0 |
                      67 Salinf EP00049 Salpingoeca infusionum P015949 | OG0 |
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                          Salkve_EP00050_Salpingoeca_kvevrii_P001662 | OG0 |
                  Salkve_EP00050_Salpingoeca_kvevrii_P001657 | OG0 |
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                       Salkve EP00050 Salpingoeca kvevrii P001659 | OG0 |
                68
                         -Salkve_EP00050_Salpingoeca_kvevrii_P001658 | OG0 |
           84
                      Codhol_EP00042_Codosiga_hollandica_P033990 | OG0 |
                         Didcos EP00039 Didymoeca costata P016004 | OG0 |
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                           •Acaspe_EP00035_Acanthoeca_spectabilis_P023477 | OG0 |
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                            •Helnan_EP00037_Helgoeca_nana_P008381 | OG0 |
                   100
                            Stedip_EP00041_Stephanoeca_diplocostata_P024106 | OG0 |
                            Diagra EP00040_Diaphanoeca_grandis_P027509 | OG0 |
                •Morvir scaffold438.g10.t1 | OG0 |
                    Chem TCONS 00022211 | OG0 |
          93
                   •Hvul g3654 1 | OG0 |
                •Nemnom_g11288.t1 | OG0 |
             <sup>l00</sup><sup>1</sup>•Rhoesc Res_008723-T1 | OG0 |
                  •Dgig_XM_028557368.1 | OG0 |
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                 •Xesp_010577-11 | OG0 |
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                 •Exapal_XM_021056531.2 | OG0 |
                 •Nvec_v1g216057 | OG0 |
                •Spis_XP_022784802_1 | OG0 |
             <sup>100</sup><sup>1</sup>-Pocdam_XM_027196537.1 | OG0 |
               •Gasp_gasp1.m3.11441.m1 | OG0 |
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                    Fspp ffun1.m4.32564.m1 | OG0 |
                      Fspp_ffun1.m4.32563.m1 | OG0 |
                    451-Fspp_ffun1.m4.32565.m1 | OG0 |
                     •Gfas_gfas1.m1.17746.m1 | OG0 |
   75
                       Amil_Amillepora19281-RA | OG0 |
                    100 Adig XM 015919109.1 | OG0 |
               •Plajan_Pjan_C568868_01 | OG0 |
                   POpea_EP00118_Oscarella_pearsei_P002014 | OG0 |
                100 Ocar_g2035 | OG0 |
                Clacor_Ccor_scaffold10712_01 | OG0 |
                  Lcom_lcpid93707 | OG0 |
                      Leuniv_Lniv_C794034_01 | OG0 |
                         •Gracom_Gcom_scaffold15834_01 | OG0 |
                         •Scil scict002181.1 | OG0 |
     69
                   -HoiH23_PIH23_009289-RA | OG0 |
                <sup>100</sup><sup>1</sup>•Hhon_g07571.t1 | OG0 |
                    -HoiH11_mRNA6759 | OG0 |
             100
                     TrH2_TrispH2_005867-RA | OG0 |
                  100 Tadh_TriadT25469 | OG0 |
       85
                •Plespi_Pspi_scaffold17936_01 | OG0 |
                     -Xestes_maker-XT_scaffold513-augustus-gene-0.6-mRNA-1 | OG0 |
                          •Xestes_maker-XT_scaffold45004-augustus-gene-0.2-mRNA-1 | OG0 |
              99
         98
                      •Aque_Aqu2.1.23842_001 | OG0 |
                       Halamb_c42385_g1_i2mm.1098 | OG0 |
                     801-Halamb_c42524_g2_i1mm.1116 | OG0 |
           98
                      •Tetwil_g22133.t2 | OG0 |
                      Stycar_augustus_masked-SC_scaffold113405-processed-gene-0.1-mRNA-1 | OG0 |
                   <sup>100]</sup>•Stycar_maker-SC_scaffold4362-augustus-gene-0.81-mRNA-1 | OG0 |
                       Spolac c100810 g1 i1 m.23588c100810-g1 | OG0 |
                    <sup>100</sup>Ū-Emue Em0023g709a.t1 | OG0 |
0.416667
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