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Salkve EP00050 Salpingoeca kvevrii P023073 | OG0 |
                   Didcos_EP00039_Didymoeca_costata_P033258 | OG0 |
                      Diagra_EP00040_Diaphanoeca_grandis_P037482 | OG0 |
       100
                    Stedip_EP00041_Stephanoeca_diplocostata_P033440 | OG0 |
 100
          100
                         Acaspe EP00035 Acanthoeca spectabilis P021433 | OG0 |
                          •Helnan EP00037 Helgoeca nana P030585 | OG0 |
                           •Helnan_EP00037_Helgoeca_nana_P030583 | OG0 |
    32
                       Saldol EP00054 Salpingoeca dolichothecata P016909 | OG0 |
        84
                         •Codhol_EP00042_Codosiga_hollandica_P055310 | OG0 |
                    Mylflu EP00047 Mylnosiga fluctuans P011844 | OG0 |
      32
                       Salmac EP00052 Salpingoeca macrocollata P017915 | OG0 |
              100
                      -Salpun_EP00053_Salpingoeca_punica_P026267 | OG0 |
        60
                     Salurc_EP00051_Salpingoeca_urceolata_P002079 | OG0 |
                       •Microa_EP00043_Microstomoeca_roanoka_P013840 | OG0 |
                        Sros EGD82369 | OG0 |
              100
                          Choper_EP00045_Choanoeca_perplexa_P018135 | OG0 |
                 100
                         •Mbre EDQ86561 | OG0 |
              •Gracom_Gcom_C536754_01 | OG0 |
               •Leuniv_Lniv_C747273_01 | OG0 |
         100
                      -Leuniv_Lniv_C808902 01 | OG0 |
                      Gracom Gcom C561570 01 | OG0 |
                    Lcom_lcpid87611 | OG0 |
              76
                  <sup>941</sup>Lcom_lcpid120591 | OG0 |
                    Lcom_lcpid138214 | OG0 |
                  891 Lcom | lcpid137655 | OG0 |
                   •Beraby_Baby_sb355803 | OG0 |
                      Pbac sb2664948 | OG0 |
                99
                    •Pbac_sb2652512 | OG0 |
                        Horcal Hcv1.av93.c5.g137.i1 | OG0 |
 100
                          Horcal Hcv1.av93.c5.g137.i2 | OG0 |
             100
                        89 Horcal_Hcv1.av93.c5.g137.i3 | OG0 |
                      Vmul sbMO473587 | OG0 |
                   <sup>1001</sup>-Vmul_sbMO473611 | OG0 |
                      Coeast Cast sb785634 | OG0 |
                100
                          Coemet Cmet scaffold4075 01 | OG0 |
                   100
                          Coemet_Cmet_scaffold4076_01 | OG0 |
                           •Coeast_Cast_sb785635 | OG0 |
                             Coeast Cast sb785633 | OG0 |
                           <sup>84</sup> Coeast_Cast_sb785632 | OG0 |
    39
                   Xestes maker-XT_scaffold3865-augustus-gene-0.5-mRNA-1 | OG0 |
                     •Aque_Aqu2.1.41625_001 | OG0 |
             100
                     •Halamb_c42345_g1_i2mm.1089 | OG0 |
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                   •Tetwil_g25155.t2 | OG0 |
                    Spolac_c104554_g3_i1_m.41362c104554-g3 | OG0 |
                 <sup>100</sup> Spolac c104554 g3_i2_m.41364c104554-g3 | OG0 |
                    •Plajan_Pjan_scaffold23791_01 | OG0 |
                       Opea_EP00118_Oscarella_pearsei_P019122 | OG0 |
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                        <sub>!</sub>Opea_EP00118_Oscarella_pearsei_P019121 | OG0 |
      49
                   100
                      <sup>981</sup>-Opea EP00118_Oscarella_pearsei_P019120 | OG0 |
                         r-TrH2 TrispH2 001325-RA | OG0 |
                      <sup>100</sup> Tadh_TriadT22916 | OG0 |
                          •HoiH11_mRNA7118 | OG0 |
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                               <sub>!</sub>HoiH23_PIH23_000529-RA | OG0 |
                      100
                            <sup>[100]</sup> HoiH06 _mRNA5886 | OG0 |
                         100
                               Hhon g01234.t1 | OG0 |
                            <sup>100</sup> HoiH04_mRNA10842 | OG0 |
                          •Bralan_BLAG18000900_1 | OG0 |
                   100
                            •Hsap_ENST00000338432 | OG0 |
                           •Hsap ENST00000616317 | OG0 |
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                            •Spur XM 030988059.1 | OG0 |
               100
                               •Cscu_XM_023367420.1 | OG0 |
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                                  •icas_iC015613_001 | OG0 |
                          100
                                 •Dpul_mRNA26376 | OG0 |
                     83
                                  •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA150877 | OG0 |
                                Acagra_model.g6517.t1 | OG0 |
                                 •Cgig_XM_034477561.1 | OG0 |
                              <sup>100</sup> Cgig_XM_034477509.1 | OG0 |
             93
                               •Chem_TCONS_00008959 | OG0 |
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                               •Hvul_g4455_1 | OG0 |
                             •Morvir scaffold56.g15.t1 | OG0 |
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                             Sanmal Sma 017520-T1 | OG0 |
                                Aaur scaffold5321.g1.t1 | OG0 |
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                              <sup>96</sup> Aaur_scaffold359.g16.t1 | OG0 |
                                 •Rhoesc_Res_002397-T1 | OG0 |
                                   Nemnom g7074.t1 | OG0 |
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                                <sup>1001</sup>Nemnom_g8392.t1 | OG0 |
                           •Dgig XM 028552439.1 | OG0 |
                            •Xesp_008415-T1 | OG0 |
                               Exapal_XM_021047163.2 | OG0 |
                  97
                       100
                              •Nvec_v1g20079 | OG0 |
                              Gfas_gfas1.m1.18636.m1 | OG0 |
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                                -Amil_Amillepora05958-RA | OG0 |
                                  •Adig_XM_015898567.1 | OG0 |
                        100
                              Fspp_ffun1.m4.29687.m1 | OG0 |
                                Gasp_gasp1.m3.12808.m1 | OG0 |
                                  •Spis_XP_022806358_1 | OG0 |
                               <sup>100</sup>  Pocdam XM 027191210.1 | OG0 |
0.416667
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