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Lcom_lcpid97917 | OG0 |
               •Leuniv_Lniv_scaffold6024_01 | OG0 |
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                 Scil_scict025005.1 | OG0 |
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                   Gracom_Gcom_scaffold3900_01 | OG0 |
                <sup>100</sup> Gracom_Gcom_scaffold3899_01 | OG0 |
 80

    Salmac EP00052 Salpingoeca macrocollata P033070 | OG0 |

           98
                      Salpun_EP00053_Salpingoeca_punica_P014356 | OG0 |
                     •Mylflu_EP00047_Mylnosiga_fluctuans_P004122 | OG0 |
                                 Salinf_EP00049_Salpingoeca_infusionum_P018555 | OG0 |
     79
                                     Microa EP00043 Microstomoeca roanoka P030046 | OG0 |
                          100
                                Sros EGD83299 | OG0 |
       69
                100
                                  •Mbre_EDQ88874 | OG0 |
                                Chofle_EP00730_Choanoeca_flexa_P019831 | OG0 |
                      100
                                 ·Choper_EP00045_Choanoeca_perplexa_P010922 | OG0 |
         58
                               Didcos EP00039 Didymoeca costata P019438 | OG0 |
                            Salurc EP00051 Salpingoeca urceolata P027838 | OG0 |
                  66
                               Salkve_EP00050_Salpingoeca_kvevrii_P010936 | OG0 |
            57
                            Diagra EP00040 Diaphanoeca grandis P014718 | OG0 |
                          Stedip EP00041 Stephanoeca diplocostata P007894 | OG0 |
                                  •Acaspe_EP00035_Acanthoeca_spectabilis_P022519 | OG0 |
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                                   •Helnan EP00037 Helgoeca nana P026266 | OG0 |
                    -HoiH23_PIH23_001047-RA | OG0 |
                 <sup>96</sup> HoiH06_mRNA3193 | OG0 |
                      <sub>f</sub>Hhon_g01534.t1 | OG0 |
                    <sup>1001</sup>HoiH04 _mRNA5199 | OG0 |
                       •HoiH11 mRNA2292 | OG0 |
                           TrH2_TrispH2_002306-RA | OG0 |
                        •Plajan_Pjan_scaffold17344_01 | OG0 |
                      <sub>"</sub>Ocar_g8988 | OG0 |
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                        62<sup>1</sup>Opea EP00118 Oscarella pearsei P017882 | OG0 |
                        Xestes maker-XT scaffold2388-augustus-gene-0.58-mRNA-1 | OG0 |
     92
 80
                     •Aque_Aqu2.1.37546_001 | OG0 |
                      •Halamb_c39597_g1_i1mm.722 | OG0 |
       99
                ·Stycar_maker-SC_scaffold5271-snap-gene-0.8-mRNA-1 | OG0 |
                    Tetwil g2944.t1 | OG0 |
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                                   •Emue Em0007g680a.t1 | OG0 |
                         -Spolac_c102478_g1_i1_m.29505c102478-g1 | OG0 |
                                     •Emue Em0007g679a.t1 | OG0 |
                   •Hsap_ENST00000292314 | OG0 |
                                   <sub>!</sub>Horcal_Hcv1.av93.c8.g221.i1 | OG0 |
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                      Bralan_BLAG12000656_1 | OG0 |
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                               •Tcas_TC004360_001 | OG0 |
                        Spur_XM_785778.5 | OG0 |
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                     Skow_XM_002740622.2 | OG0 |
                          •Cscu_XM_023384426.1 | OG0 |
                            •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA179668 | OG0 |
                          Acagra model.g5990.t1 | OG0 |
                            •Cgig_XM_011420357.3 | OG0 |
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                                     •Dpul mRNA28600 | OG0 |
                              •Hvul g31068 1 | OG0 |
                        <sup>100</sup> Hvul_g31073_1 | OG0 |
                     Morvir scaffold12.g31.t1 | OG0 |
                       Sanmal Sma 005582-T1 | OG0 |
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                          Aaur scaffold58.g18.t1 | OG0 |
                100
                          •Nemnom_g16739.t1 | OG0 |
                      <sup>100</sup><sup>1</sup>-Rhoesc_Res_005559-T1 | OG0 |
        94
                  •Nvec v1g238072 | OG0 |
                    •Exapal XM 021049797.1 | OG0 |
                       •Dgig_XM_028543844.1 | OG0 |
                        Xesp_009484-T1|OG0|
                            --Xesp_004085-T1 | OG0 |
                      •Fspp_ffun1.m4.1397.m1 | OG0 |
             86
                        •Gasp gasp1.m3.23735.m1 | OG0 |
                          ┌•Spis_XP_022788496_1 | OG0 |
                       <sup>100</sup><sup>-</sup>Pocdam_XM_027184425.1 | OG0 |
                98
                     •Gfas_gfas1.m1.6289.m1 | OG0 |
                       Amil_Amillepora14731-RA | OG0 |
                     100 Adig_XM_015900540.1 | OG0 |
                       <sup>97</sup> Adig XM 015900541.1 | OG0 |
0.416667
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