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-Pbac sb2646573 | OG0 |
                                 Horcal_Hcv1.av93.c4.g102.i1 | OG0 |
                            100
                               <sup>53</sup> Pbac sb2637757 | OG0 |
                        100
                                  •Eupdun_Edun_sb10628175 | OG0 |
                                     Beraby Baby sb343039 | OG0 |
                                     -Lampan Lpan 13912 01 | OG0 |
                                            •Coemet_Cmet_scaffold11249_01 | OG0 |
                                          -Vmul SimC396614 | OG0 |
                     Bralan_BLAG03000636_1 | OG0 |
          93
                        •Hsap ENST00000251993 | OG0 |
                     Skow XM 006824062.1 | OG0 |
 98
           98
                       •Spur XM 030989382.1 | OG0 |
                    •Acagra_model.g1891.t1 | OG0 |
      79
                      ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA228278 | OG0 |
             100
                        Cgig XM 011449875.2 | OG0 |
          100
                       •Tcas TC007350 001 | OG0 |
                  100
                          •Dpul mRNA23954 | OG0 |
              100
                         Cscu_XM_023358142.1 | OG0 |
                       Cscu_XM_023367496.1 | OG0 |
   54
                               •Morvir scaffold1626.g1.t1 | OG0 |
                               •Hvul g31971 1 | OG0 |
                           •Chem_TCONS_00023685 | OG0 |
             99
                          •Sanmal_Sma_024214-T1 | OG0 |
                           Aaur_scaffold24.g34.t2 | OG0 |
                           •Nemnom_g2262.t1 | OG0 |
                       <sup>100</sup><sup>1</sup>-Rhoesc Res_017315-T1 | OG0 |
     58
                •Exapal_XM_021048670.2 | OG0 |
           89
                 •Nvec_v1g85813 | OG0 |
                    •Amil_Amillepora13295-RA | OG0 |
        100
                    •Adig XM 015897466.1 | OG0 |
                    Gfas gfas1.m1.1735.m1 | OG0 |
             100
                      Spis_XP_022781754_1 | OG0 |
                       •Gasp gasp1.m3.288.m1 | OG0 |
                    <sup>96</sup>-Fspp_ffun1.m4.21044.m1 | OG0 |
                                     Salmac_EP00052_Salpingoeca_macrocollata_P024382 | OG0 |
               98
                                    Salpun_EP00053_Salpingoeca_punica_P018496 | OG0 |
                                     Mylflu EP00047 Mylnosiga fluctuans P014743 | OG0 |
                    60
                                              Salkve_EP00050_Salpingoeca_kvevrii_P008264 | OG0 |
                                  •Salurc_EP00051_Salpingoeca_urceolata_P014944 | OG0 |
         100
                                        •Codhol_EP00042_Codosiga_hollandica_P042057 | OG0 |
                     31
                                           Saldol EP00054 Salpingoeca dolichothecata P019978 | OG0 |
                                                  •Salinf_EP00049_Salpingoeca_infusionum_P011237 | OG0 |
             66
                                            •Microa EP00043 Microstomoeca roanoka P007167 | OG0 |
                         100
                                            •Sros EGD73281 | OG0 |
                                              •Chofle EP00730 Choanoeca flexa P012960 | OG0 |
                                                ·Choper_EP00045_Choanoeca_perplexa_P007358 | OG0 |
               73
                                  Didcos_EP00039_Didymoeca_costata_P019765 | OG0 |
                                      •Diagra_EP00040_Diaphanoeca_grandis_P012213 | OG0 |
                     99
                                         •Stedip_EP00041_Stephanoeca_diplocostata_P006835 | OG0 |
                                              Stedip EP00041 Stephanoeca diplocostata P006836 | OG0 |
                             100
                                         | Stedip EP00041 Stephanoeca_diplocostata_P006837 | OG0
                                97
                                         Acaspe EP00035 Acanthoeca spectabilis P038715 | OG0 |
                                           -Acaspe EP00035 Acanthoeca spectabilis P038716 | OG0 |
                                   100
                                          -Helnan_EP00037_Helgoeca_nana_P007162 | OG0 |
                                      <sup>100</sup> Helnan EP00037_Helgoeca_nana_P007163 | OG0 |
                        •Plajan_Pjan_scaffold15432 01 | OG0 |
                           •Ocar_g3923 | OG0 |
           99
                        Popea EP00118 Oscarella pearsei P000336 | OG0 |
                   100
                      <sup>99</sup> Ocar_g3924 | OG0 |
                            •Clacor_Ccor_scaffold6764_01 | OG0 |
                              Gracom Gcom C553632 01 | OG0 |
                100
   63
                                   •Lcom | lcpid53144 | OG0 |
                       100
                                         Lniv scaffold22440 01 | OG0 |
                                             -TrH2_TrispH2_005227-RA | OG0 |
                                          100 Tadh TriadT51482 | OG0 |
      69
                                                -HoiH23_PIH23_008457-RA | OG0 |
                                       100
                                            <sup>100</sup> HoiH06_mRNA12152 | OG0 |
                                               Hhon_g08133.t1 | OG0 |
                                            100 HoiH04 mRNA11676 | OG0 |
         75
                                    -Xestes_maker-XT_scaffold32448-snap-gene-0.12-mRNA-1 | OG0 |
                                               -Aque Aqu2.1.23776 001 | OG0 |
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                                  100
                                               •Halamb c45573 g1 i2m.19197 | OG0 |
                                    Stycar_maker-SC_scaffold1937-augustus-gene-0.23-mRNA-1 | OG0 |
                  100
                        99
                                        Tetwil g19678.t1 | OG0 |
                                        •Emue Em0003g722a.t1 | OG0 |
                      98
                                        •Spolac_c96802_g1_i1_m.13761c96802-g1 | OG0 |
                                     <sup>1001</sup>Spolac c96802 g1 i2 m.13762c96802-g1 | OG0 |
0.416667
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