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-Vmul SimC397414 | OG0 |
                           Coeast_Cast_sb794201 | OG0 |
               99
                        <sup>1001</sup>•Coeast_Cast_sb794204 | OG0 |
         100
                        Lampan_Lpan_C413251_01 | OG0 |
                            Mertsp Mesp sb541663 | OG0 |
                                  -Horcal_Hcv1.av93.c13.g128.i1 | OG0 |
                              <sup>100</sup><sup>1</sup>-Pbac sb2639793 | OG0 |
                  93
                               Beraby_Baby_sb333704 | OG0 |
                    96
                        61
                                    •Drygla_Dgla_sb282274 | OG0 |
                                Vmul sbMO371300 | OG0 |
                      69
                                 Vmul sbMO377223 | OG0 |
                            100
                               <sup>97</sup> Mlei_ML2057.g11.i1 | OG0 |
                         100
                               Bolinf_Binf_sb1252172 | OG0 |
                                 Bolinf Binf sb1252170 | OG0 |
                                   Mertsp Mesp sb553969 | OG0 |
                                 <sup>69</sup> Bolinf Binf sb1252171 | OG0 |
                         Clacor_Ccor_C515099_01 | OG0 |
                            •Lcom_lcpid67460 | OG0 |
              100
                              •Leuniv Lniv C765763 01 | OG0 |
                      100
                                 Gracom_Gcom_C557002 01 | OG0 |
                                Leuniv Lniv C765487 01 | OG0 |
                        •Plajan_Pjan_C561658_01 | OG0 |
              82
                              Opea_EP00118_Oscarella_pearsei_P027629 | OG0 |
                              Cscu XM 023371792.1 | OG0 |
         100
                                 Tcas TC010652 001 | OG0 |
                                             •Dpul_mRNA12258 | OG0 |
                                        •Tadh_TriadT61804 | OG0 |
                                     TrH2_TrispH2_005521-RA | OG0 |
                 46
                                         <sub>e</sub>HoiH04_mRNA15610 | OG0 |
                                 100
                                      <sup>100</sup> Hhon_g03375.t1 | OG0 |
            63
                                         •HoiH06 mRNA11004 | OG0 |
                                      <sup>100</sup><sup>1</sup>-HoiH23_PIH23_004630-RA | OG0 |
                    25
                                   Ctel gnl WGS AMQN CAPTEDRAFT mRNA150962 | OG0 |
                                  •Acagra model.g5085.t1 | OG0 |
                                  Cgig XM 034462185.1 | OG0 |
                                   Hsap ENST00000251588 | OG0 |
                           99
                                            •Hsap_ENST00000412079 | OG0 |
                                 Bralan_BLAG09000490_1 | OG0 |
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                                    Skow XM 002733202.2 | OG0 |
                                    •Spur_XM_030984079.1 | OG0 |
                             Sanmal Sma 020340-T1 | OG0 |
                               Nemnom g6244.t1 | OG0 |
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                              -Rhoesc_Res_010323-T1 | OG0 |
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                                Morvir scaffold108.g15.t1 | OG0 |
                                      Chem_TCONS_00022025 | OG0 |
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                                         •Hvul_g7628_1 | OG0 |
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                                •Dgig XM 028554370.1 | OG0 |
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                                 •Xesp 005392-T1 | OG0 |
                                 Exapal XM 021046117.2 | OG0 |
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                                  Nvec_v1g210509 | OG0 |
                                  •Amil Amillepora04979-RA | OG0 |
                   35
                        100
                                    Adig XM 015901719.1 | OG0 |
                           100
                                   Gfas gfas1.m1.4749.m1 | OG0 |
                                     •Gasp gasp1.m3.2653.m1 | OG0 |
                                      Fspp ffun1.m4.12594.m1 | OG0 |
                      33
                                    <sup>78</sup>Ū-Spis_XP_022798922_1 | OG0 |
                               Emue_Em0021g306a.t1 | OG0 |
                                 Spolac c88566 g1 i1 m.5121c88566-g1 | OG0 |
                           100
                              <sup>1001</sup>Spolac_c88566_g1_i3_m.5122c88566-g1 | OG0 |
                        93
                                  Tetwil g23954.t2 | OG0 |
                                       •Halamb_c43976_g1_i2mm.1378 | OG0 |
                                       •Aque_Aqu2.1.41518_001 | OG0 |
                            87
                                    Stycar maker-SC scaffold113779-snap-gene-0.11-mRNA-1 | OG0 |
                                     Stycar_maker-SC_scaffold113779-snap-gene-0.12-mRNA-1 | OG0 |
                                  <sup>100</sup>-Stycar_maker-SC_scaffold4431-snap-gene-0.3-mRNA-1 | OG0 |
0.416667
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