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√Vmul_sbMO414802 | OG0 |
                  Coemet_Cmet_scaffold19547_01 | OG0 |
               <sup>100]</sup>•Coeast_Cast_sb766571 | OG0 |
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                 -Lampan_Lpan_C412511_01 | OG0 |
                  •Pbac sb2662187 | OG0 |
                    Drygla Dgla sb283428 | OG0 |
                      •Mertsp Mesp sb526509 | OG0 |
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                       •Mlei_ML0789.g13.i1 | OG0 |
                        Bolinf_Binf_sb1188343 | OG0 |
                     <sup>1001</sup>-Mertsp_Mesp_sb607128 | OG0 |
                      •Cscu XM 023374440.1 | OG0 |
                     •HoiH11_mRNA293 | OG0 |
                        TrH2_TrispH2_008549-RA | OG0 |
                     <sup>100</sup> Tadh TriadT50572 | OG0 |
               100
                        -HoiH23 PIH23 008196-RA | OG0 |
       100
                    <sup>100</sup> HoiH06_mRNA7525 | OG0 |
                       <sub>[</sub>Hhon_g05690.t1 | OG0 |
                     <sup>100</sup> HoiH04_mRNA10062 | OG0 |
                     Spur_XM_030987833.1 | OG0 |
                           Opea_EP00118_Oscarella_pearsei_P004675 | OG0 |
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                           •Plajan_Pjan_scaffold632_01 | OG0 |
                            •Clacor Ccor C509121 01 | OG0 |
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                                 Lcom Icpid85252 | OG0 |
                                  •Gracom_Gcom_scaffold15533_01 | OG0 |
                            Plespi_Pspi_C607012_01 | OG0 |
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            69
                              •Tetwil g10141.t1 | OG0 |
                                Stycar_maker-SC_scaffold102045-augustus-gene-0.2-mRNA-1 | OG0 |
                    99
                           <sup>100</sup>lStycar_maker-SC_scaffold86453-snap-gene-0.6-mRNA-1 | OG0 |
                                      •Aque_Aqu2.1.43610_001 | OG0 |
                                    -Xestes maker-XT scaffold74549-augustus-gene-0.13-mRNA-1 | OG0 |
                                      Spolac_c85463_g1_i1_m.3732c85463-g1 | OG0 |
                                     •Spolac_c99146_g1_i3_m.18953c85463-g1 | OG0 |
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                                100
                                        rEmue_Em0674g10a.t1 | OG0 |
                                           <u>r</u>Emue_Em0010g39a.t1 | OG0 |
                                        <sup>100</sup>-Emue Em0010g80a.t1 | OG0 |
                         -Bralan_BLAG06001067_1 | OG0 |
                  23
                             -Hsap ENST00000368662 | OG0 |
                          Skow_XM_002739694.2 | OG0 |
                             •Nvec_v1g89613 | OG0 |
                              Exapal XM 021056311.2 | OG0 |
                               rAmil_Amillepora20385-RA | OG0 |
                       99
                            <sup>100</sup> Adig_XM_015901898.1 | OG0 |
                                Gfas_gfas1.m1.4732.m1 | OG0 |
                                  Fspp_ffun1.m4.8651.m1 | OG0 |
                                    •Gasp gasp1.m3.20196.m1 | OG0 |
                                    Pocdam XM 027185395.1 | OG0 |
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                                  <sup>97</sup> Spis_XP_022806330_1 | OG0 |
                              Acagra_model.g18563.t1 | OG0 |
                                •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA218781 | OG0 |
                                Ctel gnl WGS AMQN CAPTEDRAFT mRNA89314 | OG0 |
                                Cgig XM 011436043.3 | OG0 |
                                     •Acagra_model.g18565.t1 | OG0 |
                               Dgig XM 028549112.1 | OG0 |
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                                  •Xesp_002703-T1 | OG0 |
                                         Chem_TCONS_00028591 | OG0 |
                       87
                             100
                                          •Hvul_g24327_1 | OG0 |
                                       -Morvir_scaffold85.g68.t1 | OG0 |
                         99
                                     •Sanmal_Sma_016637-T1 | OG0 |
                                     •Nemnom_g11594.t1 | OG0 |
                                     •Rhoesc Res 012531-T1 | OG0 |
0.416667
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