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•Dgig XM 028552074.1 | OG0 |
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                100
                               Gracom_Gcom_C544390 01 | OG0 |
                                 Leuniv Lniv C804164 01 | OG0 |
                            •Emue_Em0001g1400a.t1 | OG0 |
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                              •Tetwil_g9969.t1 | OG0 |
                      100
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                              -Xestes snap masked-XT_scaffold3009-processed-gene-0.17-mRNA-1 | OG0 |
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                                   •Aque Aqu2.1.37435 001 | OG0 |
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              Exapal XM 021050660.2 | OG0 |
                 -Spis XP 022805583 1 | OG0 |
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             <sup>1001</sup>-Pocdam_XM_027195349.1 | OG0 |
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                  <sup>100</sup> Adig XM _015912476.1 | OG0 |
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              •Skow_XM_002740093.2 | OG0 |
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                                            •Hsap ENST00000609480 | OG0 |
        63
                                                 •Cscu XM 023375505.1 | OG0 |
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                                      ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA122410 | OG0 |
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              •Morvir scaffold19.g63.t1 | OG1 |
                   •Sanmal_Sma_009183-T1 | OG1 |
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                     Aaur scaffold2871.g2.t1 | OG1 |
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                     Nemnom g9452.t1 | OG1 |
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                          Skow XM 002740095.2 | OG1 |
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                          •Skow XM 002740092.2 | OG1 |
                                                 •Tcas TC011348 001 | OG1 |
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                          TrH2 TrispH2 000032-RA | OG1 |
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                       <sup>1001</sup>Tadh_TriadT52375 | OG1 |
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                             •HoiH23 PIH23 000880-RA | OG1 |
                              •HoiH06_mRNA1010 | OG1 |
                              <sub>e</sub>Hhon_g11557.t1 | OG1 |
                           <sup>100</sup> HoiH04_mRNA17388 | OG1 |
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                Opea_EP00118_Oscarella_pearsei_P025160 | OG1 |
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                                         •Vmul_Simscaffold9085_01 | OG1 |
                             100 Coemet Cmet_scaffold22355_01 | OG1 |
                              Drygla_Dgla_sb288537 | OG1 |
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                              Pbac sb2640684 | OG1 |
                           <sup>100</sup> Horcal_Hcv1.av93.c8.g708.i1 | OG1 |
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                               -Beraby_Baby_sb323258 | OG1 |
                                Mlei ML1498.g3.i1 | OG1 |
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                             <sup>100</sup><sup>1</sup>•Vmul sbMO432665 | OG1 |
                                Mertsp_Mesp_sb547432 | OG1 |
                             <sup>100</sup>1•Mertsp_Mesp_sb547433 | OG1 |
                           <sub>r</sub>Bolinf_Binf_sb1216023 | OG1 |
                        <sup>100</sup> Mertsp_Mesp_sb564434 | OG1 |
                             •Lampan_Lpan_14389_01 | OG1 |
                                   •Eupdun Edun sb10665179 | OG1 |
                        87
                                             •Lampan_Lpan_C404281_01 | OG1 |
                          100
                                           Bolinf Binf sb1211177 | OG1 |
                                        <sup>100</sup> Mertsp_Mesp_sb542912 | OG1 |
                                  100
                                                •Pbac sb2643501 | OG1 |
                                     100
                                                 •Mlei ML1498.g4.i1 | OG1 |
                                                    -Bolinf_Binf_sb1255196 | OG1 |
                                                 <sup>100</sup> Mertsp_Mesp_sb568253 | OG1 |
                                                   Drygla_Dgla_sb291177 | OG1 |
                                                   Bolinf_Binf_sb1187887 | OG1 |
                                                <sup>100</sup> Beraby Baby sb325649 | OG1 |
0.416667
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