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•Tetwil_g27180.t1 | OG0 |
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      92
                          •Emue_Em0033g11a.t1 | OG0 |
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                       <sup>100</sup>-Emue_Em0033g23a.t1 | OG0 |
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                   -Xestes_maker-XT_scaffold30178-augustus-gene-0.5-mRNA-1 | OG0 |
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                        Eupdun Edun sb10632129 | OG0 |
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                           Coemet_Cmet_scaffold11644_01 | OG0 |
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                            <sub>[</sub>•Bolinf_Binf_sb1219366 | OG0 |
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                                  Pbac sb2657878 | OG0 |
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                                    Horcal Hcv1.av93.c1.g287.i1 | OG0 |
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                                 Drygla_Dgla_sb279012 | OG0 |
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                                  -Beraby_Baby_sb337547 | OG0 |
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                         TrH2 TrispH2 006337-RA | OG0 |
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                             -HoiH23 PIH23 002271-RA | OG0 |
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                         Ocar g2416 | OG0 |
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                            Popea EP00118 Oscarella pearsei P007276 | OG0 |
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                           Cscu XM 023387998.1 | OG0 |
                              •Dpul mRNA10738 | OG0 |
                               •Tcas TC010010 001 | OG0 |
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                         •Spis_XP_022780047_1 | OG0 |
                      <sup>100</sup> Pocdam XM 027202330.1 | OG0 |
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                          •Gasp_gasp1.m3.22450.m1 | OG0 |
                          •Gfas_gfas1.m1.6170.m1 | OG0 |
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0.416667
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