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•Xestes_maker-XT_scaffold188124-augustus-gene-0.132-mRNA-1 | OG0 |
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                     •Lcom lcpid208072 | OG1 |
                        Lcom | lcpid185024 | OG1 |
                    Eupdun Edun sb10673820 | OG1 |
                       Lampan_Lpan_14978_01 | OG1 |
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                         •Vmul_SimC403202 | OG1 |
                  36
                          •Coemet_Cmet_C598371_01 | OG1 |
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             100
                             Coeast Cast sb808422 | OG1 |
                               rCoeast Cast sb808435 | OG1 |
                                 -Coeast_Cast_sb808425 | OG1 |
                               931 Coeast Cast sb808426 | OG1 |
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                       -Pbac_sb2649783 | OG1 |
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                         Pbac sb2645859 | OG1 |
                       99 Horcal Hcv1.av93.c2.g576.i1 | OG1 |
                          Beraby_Baby_sb332405 | OG1 |
                            Drygla Dgla sb275044 | OG1 |
                          67 Beraby Baby sb332406 | OG1 |
                           Drygla_Dgla_sb298876 | OG1 |
                        <sup>100</sup> Drygla_Dgla_sb298877 | OG1 |
                             Mertsp_Mesp_sb572111 | OG1 |
                                -Bolinf_Binf_sb1244278 | OG1 |
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                                <sup>54</sup> Mertsp_Mesp_sb572110 | OG1 |
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                              •Vmul sbMO465764 | OG1 |
                                    •Vmul sbMO465763 | OG1 |
                                  Bolinf Binf sb1234121 | OG1 |
                                   Bolinf_Binf_sb1234120 | OG1 |
                                 <sup>991</sup>Mertsp_Mesp_sb572112 | OG1 |
                    TrH2 TrispH2 007011-RA | OG2 |
                 <sup>100</sup> Tadh TriadT49733 | OG2 |
                    •HoiH11 mRNA4244 | OG2 |
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                       HoiH23 PIH23 000831-RA | OG2 |
                     <sup>91</sup> HoiH06 mRNA2112 | OG2 |
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                        Hhon g04051.t1 | OG2 |
                     <sup>1001</sup>HoiH04_mRNA3705 | OG2 |
                          •Morvir scaffold349.g16.t1 | OG2 |
                          •Cgiq XM 011428854.3 | OG2 |
                             Spur_XM_782442.5 | OG2 |
                           Bralan BLAG19000166 1 | OG2 |
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                             Clacor Ccor C540279 01 | OG2 |
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                              Lcom | lcpid67672 | OG2 |
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                                •Scil_scict022177.1 | OG2 |
                                 <sub>「</sub>-Leuniv_Lniv_scaffold15387_01 | OG2 |
                              98<sup>1</sup>-Gracom_Gcom_C548652_01 | OG2 |
                          •Acagra_model.g10793.t1 | OG2 |
                         -Lcom_lcpid255035 | OG2 |
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                              Hsap ENST00000261755 | OG2 |
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                                           Lcom | lcpid143536 | OG2 |
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                                         •Cscu_XM_023373272.1 | OG2 |
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                                     •Tcas_TC009325_001 | OG2 |
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                                       •Dpul_mRNA30433 | OG2 |
                                  <sup>100</sup> Dpul_mRNA16016 | OG2 |
                       •Nvec_v1g241460 | OG2 |
                       •Exapal_XM_021040150.1 | OG2 |
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                          <sub>•</sub>Amil_Amillepora26817-RA | OG2 |
                       <sup>100</sup> Adig XM_015911224.1 | OG2 |
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                          •Fspp ffun1.m4.22962.m1 | OG2 |
                            3asp_gasp1.m3.18838.m1 | OG2 |
                             •Pocdam_XM_027189417.1 | OG2 |
                                 •Spis_XP_022778306_1 | OG2 |
                            <sup>1001</sup>Spis XP 022802454 1 | OG2 |
                          -Chem_TCONS_00071736 | OG2 |
                             Hvul g31925 1 | OG2 |
                               •Cscu_XM_023388979.1 | OG2 |
                            •Morvir_scaffold349.g14.t1 | OG2 |
                            Aaur scaffold93.g13.t1 | OG2 |
                              •Nemnom_g3133.t1 | OG2 |
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                           •Nvec_v1g157033 | OG2 |
                           •Dgig_XM_028547077.1 | OG2 |
                            •Xesp_021840-T1 | OG2 |
                               •Tetwil_g26745.t2 | OG2 |
                               Emue Em0010g72a.t1 | OG2 |
                                 <sub>|</sub>Spolac_c103187_g1_i1_m.32894c103187-g1 | OG2 |
                           100
                              <sup>1001</sup>Spolac_c103187_g1_i4_m.32895c103187-g1 | OG2 |
                                  Aque_Aqu2.1.24512_001 | OG2 |
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                                 •Plajan_Pjan_C560338_01 | OG2 |
                                  Ocar_g1827 | OG2 |
                              <sup>100</sup>-Opea_EP00118_Oscarella_pearsei_P025052 | OG2 |
0.416667
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