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Xestes_maker-XT_scaffold55704-snap-gene-0.1-mRNA-1 | OG0 |
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                          Aque Aqu2.1.12631 001 | OG0 |
                         <sup>98</sup> - Aque_Aqu2.1.07695_001 | OG0 |
   Aque_Aqu2.1.29871_001 | OG1 |
      Xestes_maker-XT_scaffold138286-augustus-gene-0.3-mRNA-1 | OG1 |
        •Halamb c56277 g1 i1m.42918 | OG1 |
                    •Xestes augustus masked-XT scaffold10102-processed-gene-0.0-mRNA-1 | OG1 |
                Xestes_maker-XT_scaffold69728-augustus-gene-0.1-mRNA-1 | OG1 |
      85
                   Xestes_maker-XT_scaffold74568-augustus-gene-0.3-mRNA-1 | OG1 |
               Lcom_lcpid196673 | OG1 |
                  •Tetwil g25919.t1 | OG1 |
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                      -Tetwil g5725.t1 | OG1 |
                   <sup>100</sup>1-Tetwil_g5724.t1 | OG1 |
                     retwil_g5716.t1 | OG1 |
                 83
                          Emue Em0015g1228a.t1 | OG1 |
                           -Spolac c104942 g1 i1 m.44448c104942-g1 | OG1 |
                         <sup>74</sup> Spolac_c104942_g1_i2_m.44449c104942-g1 | OG1 |
                            •Emue_Em0015g1227a.t1 | OG1 |
            64
                           r-Spolac_c104942_g1_i2_m.44450c104942-g1 | OG1 |
                             Emue_Em0015g1226a.t1 | OG1 |
                           <sup>87</sup> Emue Em0015g1248a.t1 | OG1 |
                 Stycar maker-SC scaffold20685-augustus-gene-0.7-mRNA-1 | OG1 |
                     •Plespi_Pspi_scaffold5476_01 | OG1 |
                        -Tetwil_g5719.t2 | OG1 |
                              -Plespi_Pspi_scaffold371_01 | OG1 |
                     69
                                           *Tetwil g26533.t1 | OG1 |
                                             *Tetwil_g26535.t1 | OG1 |
                 48
                                           981 Tetwil g26528.t1 | OG1 |
                          •Plespi_Pspi_scaffold370_01 | OG1 |
                               -Plajan_Pjan_scaffold23776_01 | OG1 |
                                 Popea EP00118 Oscarella pearsei P024990 | OG1 |
                              <sup>100</sup>¹•Ocar_g7036 | OG1 |
                                 •Lcom_lcpid236355 | OG1 |
                                -Clacor_Ccor_C513541_01 | OG1 |
                                    Clacor Ccor C526167 01 | OG1 |
                                  <sup>991</sup>Clacor_Ccor_C528983_01 | OG1 |
                                    Clacor Ccor C514947 01 | OG1 |
                                       Gracom_Gcom_scaffold3448_01 | OG1 |
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                                          -Leuniv_Lniv_C808644_01 | OG1 |
                                            _Lcom_lcpid6836 | OG1 |
                                                     -Lcom |cpid186226 | OG1 |
                                 <sub>[</sub>•HoiH11_mRNA7324 | OG1 |
                                  rTadh_TriadT59099 | OG1 |
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                                    HoiH23 PIH23 000641-RA | OG1 |
                                100 HoiH06 mRNA12664 | OG1 |
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                                 92 HoiH04_mRNA15798 | OG1 |
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                                     Bralan BLAG01002703 1 | OG1 |
                                       →Spur_XM_030991507.1 | OG1 |
                                    99 Skow_XM_006812043.1 | OG1 |
                                       •Cscu_XM_023373178.1 | OG1 |
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                                    <sup>91</sup>_-Tcas_TC010565_001 | OG1 |
                                        -Mertsp Mesp sb586521 | OG1 |
                                    <sup>70</sup> Dpul mRNA941 | OG1 |
                                        •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA151544 | OG1 |
                                    <sup>91</sup>-Clacor_Ccor_C515103_01 | OG1 |
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                                       <sup>70</sup><sup>1</sup>-Cgig_XM_034482090.1 | OG1 |
                                  Tetwil_g25913.t1 | OG1 |
                                       <sub>-</sub>-Exapal_XM_021061474.2 | OG1 |
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                                           •Exapal XM 028661896.1 | OG1 |
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                                             <sub>_</sub>-Hvul_g10705_1 | OG1 |
                                           <sup>26</sup>—Chem_TCONS_00048224 | OG1 |
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                                           -Nemnom_g7740.t1 | OG1 |
                                             Rhoesc_Res_006885-T1 | OG1 |
                                               Nemnom_g7739.t1 | OG1 |
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                                                 Plespi_Pspi_scaffold5477_01 | OG1 |
                                                          -Xestes_maker-XT_scaffold62426-snap-gene-0.4-mRNA-1 | OG1 |
                                      -Eupdun_Edun_sb10639856 | OG1 |
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                                            •Vmul_Simscaffold12777_01 | OG1 |
                                              •Coemet_Cmet_scaffold6098_01 | OG1 |
                                                -Coeast_Cast_sb822358 | OG1 |
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                                                991Coeast_Cast_sb822346 | OG1 |
                                           <sub>-</sub>Lampan_Lpan_2827_01 | OG1 |
                                         <sup>88</sup><sup>™</sup>-Eupdun_Edun_sb10613655 | OG1 |
                                              <sub>!</sub>Eupdun_Edun_sb10652739 | OG1 |
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                                                 Drygla_Dgla_sb287056 | OG1 |
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                                                    Drygla_Dgla_sb308837 | OG1 |
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                                                        _Bolinf_Binf_sb1281191 | OG1 |
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                                                         Pbac_sb2657029 | OG1 |
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├────
0.416667
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