**SUPPLEMENTARY FIGURES for the manuscript**

**Integrated taxonomy and DNA barcoding of Alpine midges (Diptera: Chironomidae)**

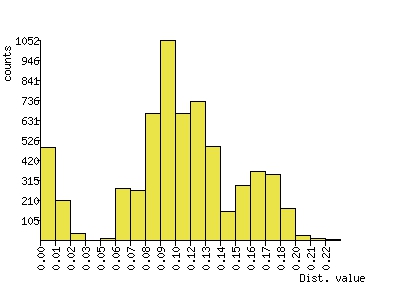
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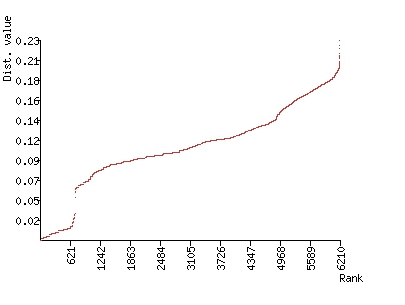
This file includes:

Supplementary Figures A, B, C

**Figure A.** Histogram of pairwise nucleotide distances. Graph obtained through the web-based interface available at (http://wwwabi.snv.jussieu.fr/public/abgd); model of nucleotide evolution settled to Kimura-2-parameter [40].



**Figure B.** Plot of ranked pairwise nucleotide distances. Graph obtained through the web-based interface available at (http://wwwabi.snv.jussieu.fr/public/abgd); model of nucleotide evolution settled to Kimura-2-parameter [40].



**Figure C.** Automatic partition of the analyzed data set. Graph obtained through the web-based interface available at (http://wwwabi.snv.jussieu.fr/public/abgd). Thenumber of groups composing the partitions (initial and recursive) are reported as a function of the prior limit between intra- and interspecies divergence.

