Poster/Software Demonstration Abstract LaTeX2ε Template

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Keywords: aberrant splicing, database, point mutation, scanning model

1 Introduction

This section title is recommended for the first section.

The number of pages for poster/software demonstration abstract is **exactly TWO**. In case the number of pages still exceeds this limit after your efforts, please contact giw2006@ims.u-tokyo.ac.jp for advice.

Do not change the text width, text height, baseline, font size, etc., specified in LaTeX2e/LaTeX/MS Word templates.

2 Method and Results

Of course you can create more sections if you want, within the page limit.

2.1 Tables

Table 1 is an example of a table.

Table 1: Bioinformatics conferences in 2006.

Conference	Date	Site
RECOMB	April 2-5 1	Venice
ISMB	August 6-10	Fortaleza
GIW	December 18-20	Yokohama

2.2 Figures

When you use a figure (Fig. 1) with a special style file, please provide it.

Printed version of the abstract is monochromatic. The poster and software demonstration abstracts are compiled as "The Sixteenth International Conference on Genome Informatics, Poster and Software Demonstration Abstracts" and will be delivered at the conference. The electronic version (PDF) will be published at JSBi Home Page (http://www.jsbi.org/jsbi_new/gioinfo.html).

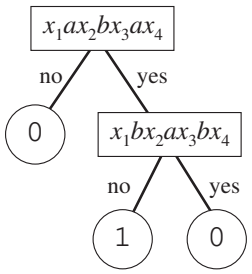

Figure 1: This is a sample figure.

2.3 Citation and References

Reference style and citation should strictly follow this template [1, 2, 3, 4, 5].

1. References should be arranged in the alphabetical order of authors and all references SHOULD be cited.
2. Style (see examples in **References** below).
   (a) Journal: author(s), title, journal name, vol(issue):startpage-endpage, year.
   (b) Proceedigns: author(s), title, Proc. conference name, publisher (if possible), startpage-endpage, year.
   (c) Book: author(s), title, publisher, year.
   (d) Other: author(s), title, any helpful information, year.
   (e) URL: http://(URL site). URLs should be listed in the last.
3. Provide all author names. If the number of authors is too large, you may reduce appropriately (such as Venter, J.C., Adams, M.D., Myers, E.W., Li, P.W., *et al.*, The sequence of the human genome, *Science*, 291(5507):1304–1351, 2001.)

3 Discussions

Authors are recommended to use spell checker program and grammar check program for improving your final version.

References

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[4] Smith, T.F. and Waterman, M.S., Identification of common molecular subsequences, *J. Mol. Biol.*, 147(1):195–197, 1981.

[5] http://www.jsbi.org/