Full Paper LaTeX2ε Template for Genome Informatics 2002

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Abstract

This document is a LaTeX2ε template file for preparing a full paper. Please read the instructions carefully and prepare your manuscript.

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

1 Introduction

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2 Method and Results

Of course you can create more sections with any section titles.

2.1 Tables

Table 1 is an example of a table.

Table 1: Bioinformatics conferences in 2002.

Conference	Date	Site
PSB	January 3-7	Hawaii
RECOMB	April 10-13	Berlin
ISMB	August 3-7	Edmonton
GIW	December 16-18	Tokyo

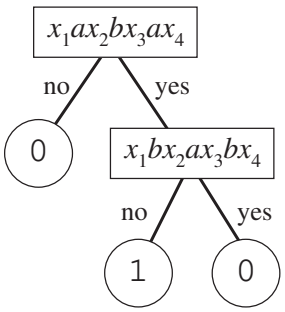

Figure 1: This is a sample figure.

Figure 2: This is a color sample figure.

2.2 Figures

When you use figures (Fig. 1, Fig. 2) with a special style file, please provide it.

Printed version of "Genome Informatics" is monochromatic and has a page limit. But, electronic version (PDF, then HTML, XML) does not have such restrictions. Therefore, we encourage the author(s) to send any hypermedia files which can be attached to your paper. The electronic version will be open to public at JSBi Home Page (http://www.jsbi.org/journal.html).

2.3 Citation and References

Here are examples of citation [1, 2, 3, 4, 6]. Reference style and citation should strictly follow this template.

1. References should be arranged in the alphabetical order of authors and all references should be cited in the paper.

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.

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3 Discussions

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

References

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