# Full Paper LaTeX2ε Template for Genome Informatics 2000

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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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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 with any section titles.

### 2.1 Tables

Table 1 is an example of a table.

Table 1: Bioinformatics conferences in 2000.

| Conference | Date | Site |
|---|---|---|
| PSB | January 5-9 | Hawaii |
| RECOMB | April 8-11 | Tokyo |
| ISMB | August 19-23 | San Diego |
| GIW | December 18-19 | Tokyo |

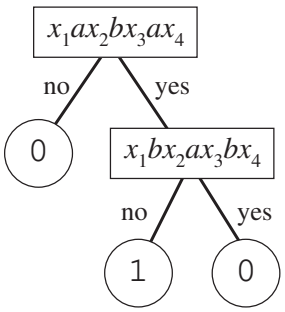


Figure 1: This is a sample figure.

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.

   (e) URL: http://(URL site). URLs should be listed in the last.

## 3 Discussions

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

## References

[1] Akutsu, T., Kuhara, S., Maruyama, O., and Miyano, S. Identification of gene regulatory networks by strategic gene disruptions and gene overexpressions, *Proc. 9th ACM-SIAM Symp. Discrete Algorithms*, 695–702, 1998.

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