Poster Abstract MS Word Template

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

1 Introduction

The number of pages for a poster abstract must be either **one or two**. Do not change the text width, text height, baseline, font size, etc., specified in this template file.

The poster abstracts are compiled as "The Joint Conference on Informatics in Biology, Medicine and Pharmacology Poster Abstracts" and will be delivered at the conference in an electric form (e.g., USB memory).

The one page format requires only abstract, and the two page format requires main body divided into sections. Both one and two page formats are all right for oral presentation in the technical sessions. The oral speakers will be selected from the poster abstracts reviewed by the program committee members. The authors can request the oral speaker reviewing when submit their poster abstracts.

2 Method and Results

You can create more sections with any section titles, within the page limit.
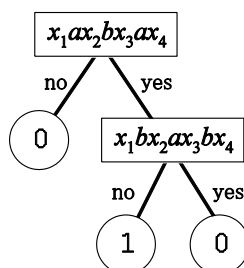
2.1 Figures

A sample figure is given in Fig. 1.

Figure 1: This is a sample figure.

2.2 Tables

Table 1 is an example of a table.

Table 1: Bioinformatics conferences in 2012.

Conference	Date	Site
RECOMB	March 28-31	Vancouver
ISMB/ECCB	July 17-July 19	Vienna
GIW	December 5-7	Busan

2.3   Citation and References

Reference style and citation should strictly follow this template [1,2,3,4,5,6].

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

2. Style (see examples in **References** below).

    (a) Journal: author(s), title, journal name, vol:startpage-endpage, year.
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    (c) Book: author(s), title, publisher, year.
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    (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., *et al.*, The sequence of the human genome, *Science*, 291:1304-1351, 2001.)

3   Discussions

Authors are recommended to use spell checker program and grammar check program for improving English presentation.

References

[1] Alon, U., *An Introduction to Systems Biology: Design Principles of Biological Circuits*, CRC Press, 2006.
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[4] Smith, T.F. and Waterman, M.S., Identification of common molecular subsequences, *Journal of Molecular Biology*, 147:195-197, 1981.
[5] Venter, J.C., *et al.*, The sequence of the human genome, *Science*, 291:1304-1351, 2001.
[6] http://cbi-society.org/cbi/taikai/taikai12/index_e.html