

A Sample LaTeX Format for the Tenth Workshop on Genome Informatics

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Abstract

This is the abstract of a sample paper for the 8th Workshop on Genome Informatics. The workshop is intended to provide a forum for researchers working in interdisciplinary areas between biological sciences and computer science.

1 Introduction

Here is the introduction text.

The Workshop on Genome Informatics is organized by the Genome Science Research Project funded by a Grant-in-Aid for Scientific Research on Priority Areas from the Ministry of Education, Science, Sports and Culture of Japan.

2 Sample Section

Of course you can create other sections if you want.

2.1 Figures

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

Here are examples of citation [1, 3, 2, 4, 5, 6, 7]. References should be arranged in alphabetical order of authors and should be cited in the text.

Acknowledgments

The section for acknowledgements should not be numbered.

References

- [1] Bowie, J. U., Lüthy, R. and Eisenberg, D., A method to identify protein sequences that fold into a known three-dimensional structure, *Science*, 253:164–170, 1991.
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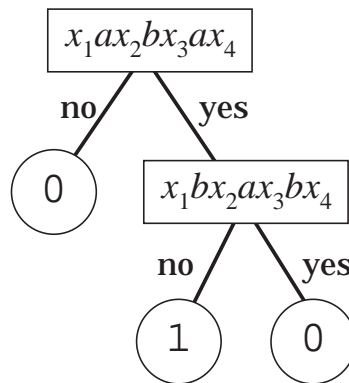


Figure 1: This is a sample figure.

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