A One-page Poster Abstract MS Word Template

FirstName1 Middle Name LastName1¹ lastname1@ims.u-tokyo.ac.jp **FirstName2 LastName2**² lastname2@cbi-society.org

FirstName3 LastName3³ Lastname3@jsbi.org

- ¹ Human Genome Center, Institute of Medical Science, University of Tokyo, 4-6-1 Shirokanedai, Minato-ku, Tokyo 108-8639, Japan
- ² Chem-Bio Informatics Society (CBI), Tokyo Institute of Technology, G3-50, 4256 Nagatsuta, Midori-ku, Yokohama, Kanagawa 226-8502, Japan
- ³ The Japanese Society for Bioinformatics Tokyo Office, University of Tokyo, 4-6-1 Shirokanedai, Tokyo 108-8639, Japan

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

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.

- [1] Alon, U., An Introduction to Systems Biology: Design Principles of Biological Circuits, CRC Press, 2006.
- [2] Altschul, S. F., Gish, W., Miller, W., Myers, E.W., and Lipman, D.J., Basic local alignment search tool, *Journal of Molecular Biology*, 215:403-410, 1990.
- [3] Friedman, N., Linial, M., Nachman, I., and Pe'er, D., Using Bayesian networks to analyze expression data, *Proc. 4th Annual International Conference on Computational Molecular Biology*, ACM, 127-135, 2000.
- [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