

A One-page Poster Abstract MS Word Template

FirstName1 Middle Name LastName1¹
lastname1@ims.u-tokyo.ac.jp

FirstName2 LastName2²
lastname2@cbi-society.org

FirstName3 LastName3²
Lastname3@cbi-society.org

¹ Human Genome Center, Institute of Medical Science, University of Tokyo, 4-6-1
Shirokanedai, Minato-ku, Tokyo 108-8639, Japan

² Tokyo Institute of Technology, J3-25, 4259, Nagatsuta, Midori-ku, Yokohama, Kanagawa
226-8503, Japan

Keywords: Electronic structure calculation, Ligand docking, Protein modeling

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

The poster abstracts are compiled as “CBI Annual Meeting 2017 Poster Abstracts” and will be published on 2 Oct. on CBI website Conference page.

The one page format requires only abstract. The oral speakers will be selected from the poster abstracts reviewed by the program committee members. The authors can check the Acceptable or Unacceptable (to be an oral speaker) box when submitting 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/taikai/taikai17/index.html>