

A One-page Poster Abstract MS Word Template

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Keywords: Electronic structure calculation, Ligand docking, Protein modeling

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[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/taikai15/index.html>