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Author Information: Complete names and affiliation of all authors, including contact details of corresponding author (Telephone, Fax and E-mail address).

Abstract: The abstract should be informative and completely self-explanatory, briefly present the topic, state the scope of the experiments, indicate significant data, and point out major findings and conclusions. The abstract should summarize the manuscript content in 300 words or less. Standard nomenclature should be used and abbreviations should be avoided. The preferable format should accommodate a description of the study background, methods, results and conclusion. Following the abstract, a list of keywords (3-10) and abbreviations should be included.

Text:

Introduction: The introduction should set the tone of the paper by providing a clear statement of the study, the relevant literature on the study subject and the proposed approach or solution. The introduction should be general enough to attract a reader's attention from a broad range of scientific disciplines.

Materials and Methods: This section should provide a complete overview of the design of the study. Detailed descriptions of materials or participants, comparisons, interventions and types of analysis should be mentioned. However, only new procedures should be described in detail; previously published procedures should be cited and important modifications of published procedures should be mentioned briefly. Capitalize trade names and include the manufacturer's name and address.

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Acknowledgement: This section includes acknowledgment of people, grant details, funds, etc.

Note: If an author fails to submit his/her work as per the above instructions, they are pleased to maintain clear titles namely headings, subheading.

References:

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Published Papers:

Laemmli UK (1970) Cleavage of structural proteins during the assembly of the head of bacteriophage T4. *Nature* 227: 680-685.

Brusic V, Rudy G, Honeyman G, Hammer J, Harrison L (1998) Prediction of MHC Class II binding peptides using an evolutionary algorithm and artificial neural network. *Bioinformatics* 14: 121-130.

Doroshenko V, Airich L, Vitushkina M, Kolokolova A, Livshits V, et al. (2007) YddG from *Escherichia coli* promotes export of aromatic amino acids. *FEMS Microbiol Lett* 275: 312-318.

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Electronic Journal Articles Entrez Programming Utilities

http://eutils.ncbi.nlm.nih.gov/entrez/query/static/eutils_help.html

Books:

Baggot JD (1999) Principles of drug disposition in domestic animals: The Basis of Veterinary Clinical Pharmacology. (1st edn), W.B. Saunders Company, Philadelphia, London, Toronto.

Zhang Z (2006) Bioinformatics tools for differential analysis of proteomic expression profiling data from clinical samples. Taylor & Francis, CRC Press.

Conferences:

Hofmann T (1999) The Cluster-Abstraction Model: unsupervised learning of topic hierarchies from text data. Proceedings of the International Joint Conference on Artificial Intelligence.

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All images must be at or above intended display size, with the following image resolutions: Line Art 800 dpi, Combination (Line Art + Halftone) 600 dpi, Halftone 300 dpi. See the Image quality specifications chart for details. Image files also must be cropped as close to the actual image as possible.

Use Arabic numerals to designate figures and upper case letters for their parts (Figure 1). Begin each legend with a title and include sufficient description so that the figure is understandable without reading the text of the manuscript. Information given in legends should not be repeated in the text.

Figure legends: These should be typed in numerical order on a separate sheet

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Suggested Equation Extraction Method

Table Specifications

Equation Specifications

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Summary diagram/figure included as part of the Supplementary Information (optional).

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