



Bioinformatics in Protein Evolution

Guest Editors:

Dr. Pablo Mier

Faculty of Biology, Institute of
Organismic and Molecular
Evolution, Johannes Gutenberg
University Mainz, 55128 Mainz,
Germany

Dr. Lisanna Paladin

Structural and Computational
Biology Unit, European Molecular
Biology Laboratory (EMBL), 69117
Heidelberg, Germany

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Message from the Guest Editors

Dear Colleagues,

The post-genomic era led to a huge and continuously increasing number of entries in protein databases. This, coupled with technologies allowing unprecedented annotation quality, provide the means to understand the evolution of protein families which are currently poorly described. For this Special Issue, we recruit manuscripts integrating bioinformatic analysis and discussing proteins' evolution, either as a process or focused on specific motifs/domains. The use of data from multiple species or strains is highly recommended. We will discuss the multiple faces of molecular evolution, from sequences to structures to functions, leveraging on the recent developments in protein features prediction and system biology.

Dr. Pablo Mier
Dr. Lisanna Paladin
Guest Editors





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Editors-in-Chief

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Department of Cellular and
Molecular Medicine, Faculty of
Health and Medical Sciences,
University of Copenhagen,
Blegdamsvej 3C, DK-2200
Copenhagen, Denmark

Prof. Dr. Lukasz Kurgan

Department of Computer
Science, Virginia Commonwealth
University, Richmond, VA 23284,
USA

Message from the Editorial Board

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Biomolecules Editorial Office
MDPI, Grosspeteranlage 5
4052 Basel, Switzerland

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