

# Special Issue

## New Bioinformatics Tools

### Message from the Guest Editor

Modern biology is characterized by the generation of massive datasets that require a computational treatment in order to distill useful information from them. Although computers are essential today in all human activities, their importance in biological research is greater due to the irreversible data-oriented trend in this field. Consequently, from being mere helping hands to the experimental work, computational tools are now indispensable for biological research and form an intrinsic part of most workflows. Daily work in molecular biology presently depends on a large ecosystem of computational tools. This ecosystem evolves with time and adapts to different requirements imposed by the peculiarities of biological data and methods popular at a given moment. *Bioengineering* welcomes submissions for its Special Issue on “New Bioinformatics Tools”. This Special Issue aims to cover the main current trends in the development and application of computational tools and methods to biological problems.

### Guest Editor

Dr. Florencio Pazos

Spanish National Research Council, Spanish National Center for Biotechnology Madrid, Madrid, Spain

### Deadline for manuscript submissions

closed (30 June 2021)



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*Bioengineering*  
Editorial Office  
MDPI, Grosspeteranlage 5  
4052 Basel, Switzerland  
Tel: +41 61 683 77 34  
[bioengineering@mdpi.com](mailto:bioengineering@mdpi.com)

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### Editor-in-Chief

Prof. Dr. Anthony Guiseppi-Elie

Department of Biomedical Engineering, Texas A&M University, College Station, TX 77843, USA

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