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# Artificial Intelligence in Bioinformatics: Current Status and Future Prospects

Guest Editors:

#### Dr. Chao Yang

Department of Chemistry, New York University, New York, NY 10003, USA

#### Dr. Peng Zhou

School of Life Science and Technology, University of Electronic Science and Technology of China, Chengdu, China

Deadline for manuscript submissions:

closed (20 July 2023)

## **Message from the Guest Editors**

In recent years, artificial intelligence (AI) has received a great deal of attention in diverse research fields, and bioinformatics is no exception. With the rapid progress of computational power and exponential increase of data, Al has been applied in many branches of bioinformatics, such biological/clinical data analysis and modeling, molecular structure prediction and structure-function analysis. The two major subsets of Al—machine learning (ML) and deep learning (DL)—have created a great deal of excitement in the research community. These methods can aid the interpretation or prediction of complex systems related to biology, chemistry, and pharmaceutical sciences, among others. Currently, the applications of ML/DL modeling in bioinformatics research are still in the preliminary stage; it would be valuable to understand how Al boosts performance in many aspects of bioinformatics research











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#### **Editor-in-Chief**

# **Prof. Dr. Giulio Nicola Cerullo**Dipartimento di Fisica, Politecnico di Milano, Piazza L. da Vinci 32, 20133 Milano, Italy

### **Message from the Editor-in-Chief**

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