SARS-CoV-2 Bioinformatics

Guest Editor:

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Message from the Guest Editor

Dear Colleagues,

   A Special Issue such as the present one offers the opportunity to portray the main contributions achieved at present in a single forum. Published in a common Special Issue, they can be a practical reference source for the topic and catalyze the cross-polymerization of the different computational approaches. High-quality scientific papers applying computational simulation methods to characterize any biological aspect of SARS-CoV-2, its infectivity, transmission, pathogenesis, diagnosis, treatments, and prevention are welcome. In particular, we would like to receive contributions that offer breakthrough contributions to the field. Works presenting new computational methods and tools are also welcome. Pure computational papers across different scales can be accepted providing they discuss the relevant experimental and/or clinical literature.

Special Issue