



Molecular Epidemiology of SARS-CoV-2, 3rd Edition

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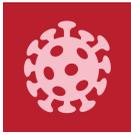
Message from the Guest Editor

Dear Colleagues,

Over the course of the SARS-CoV-2 pandemic, extensive global sustained transmission allowed for the evolution of the virus, leading to the emergence of the so-called ‘variants of concern’ (VOCs). These VOCs are characterized by several lineage-specific mutations that are suggested to impact viral transmissibility, resistance to neutralizing antibodies, and virulence. The identification of such variants has recently challenged public health authorities in terms of tracking transmission and mitigating the impact of the ongoing pandemic. This Special Issue will provide new genomic data on the SARS-CoV-2 strains circulating worldwide and present an overview of the real-time global evolution of SARS-CoV-2.

Dr. Marta Giovanetti
Guest Editor





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Editor-in-Chief

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Message from the Editor-in-Chief

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Electronic files or software regarding the full details of the calculation and experimental procedure, if unable to be published in a normal way, can be deposited as supplementary material.

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