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Phenotypic Consequences of Human Genetic Diversity

Guest Editor:

Message from the Guest Editor

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Deadline for manuscript submissions: closed (30 April 2024)

Dear Colleagues,

Over the past 15 years, studies on human genetics have unveiled remarkable discoveries in genomic medicine while elucidating the diverse genomic structure across various populations. I aspire to contribute to the expanding repository of documented genomic variants that impact human phenotypes at both DNA and RNA levels.

This Special Issue will serve as an open access platform to address the aforementioned concerns by presenting original research articles and reviews focused on exploring the phenotypic consequences of human genetic diversity. I welcome studies of diverse designs (e.g., GWAS, gene expression, SNP studies, variant discoveries, population genetic studies) conducted in this field.

Dr. Mahmut Cerkez Ergoren *Guest Editor*









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Message from the Editor-in-Chief

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