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Pharmacogenomics and Hypertension: Problems and Prospects

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Deadline for manuscript submissions:

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Message from the Guest Editors

Arterial hypertension has a strong underlying hereditary and environmental basis. Treatment has remained empiric, control rates are suboptimal, and there is minimal translation of genetics from the bench to the clinic. Although rare monogenic forms of hypertension with a distinct phenotype were initially well documented and provided specific treatments, the overall genetic risk for hypertension was considered to be polygenetic in origin. The initial GWAS studies could only explain 2% of the overall risk, which led to a degree of nihilism in relation to the pharmacogenetics of hypertension.

The aim of this Special Issue is to review and publish cutting-edge research on new developments in the understanding of the genetics of arterial hypertension, the translation of genetic research to the clinic, and future directions for pharmacogenomics. We are soliciting original articles related to understanding the genetic architecture of hypertension, genetic risk scoring, Mendelian randomization, epigenetics, the pharmacokinetics and pharmacodynamics of antihypertensive drugs, translational research, and state-of-the-art manuscripts on the status of pharmacogenomics and future directions.







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Message from the Editor-in-Chief

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