



Epigenetics of Diabetes and Related Complications

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Deadline for manuscript
submissions:

closed (30 November 2019)

Message from the Guest Editors

A growing body of evidence suggests that dynamic alterations in the epigenome (i.e., DNA methylation, histone marks, and non-coding RNAs) play a causative role in the activation of the molecular events responsible for diabetes progression and associated complications. Unveiling the epigenetic landscape in diabetic subjects and people at risk of developing diabetes is beneficial to provide tools for personalized epigenetic-based therapies.

The aim of this Special Issue is to provide a collection of original and review articles aimed at advancing the current knowledge of the epigenetics of diabetes and its implications in classic diabetes complications and co-morbidities.





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Message from the Editor-in-Chief

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