Special Issue

Recent Advances in Chromatin Structure and Dynamics

Message from the Guest Editor

We now understand that the genome is hierarchically organized into loops, topologically associated domains, compartments, and chromosome territories.

Additionally, epigenetic regulation through post-translational modifications of histone tails, DNA methylation, histone variants, and protein binding provides a diverse array of mechanisms for modulating chromatin structure and dynamics, which in turn influences gene expression. In this Special Issue, we invite researchers to submit original articles and reviews that focus on chromatin structure and dynamics, as well as its regulation by epigenetic modifications in the context of cellular function and human diseases.

Guest Editor

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Message from the Editor-in-Chief

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Editor-in-Chief

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