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Advance in Transcriptional Regulation by Conventional Metabolic Enzymes

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Message from the Guest Editor

Dear Colleagues,

Recently, many conventional metabolic enzymes are localized in nucleus, where they play novel nonconventional functions in nucleus including particularly transcriptional regulation through epigenetic modifications.

Pyruvate dehydrogenase complex (PDC) in nucleus can provide acetyl-coA, which is used for histone acetylation to regulate gene expression; Pyruvate kinase M2 (PKM2) in nucleus plays a conventional function to provide pyruvate, which is used for acetyl-CoA production to modify histone protein and a non-conventional function of protein kinase to phosphorylate histone protein and STAT3: Phosphofructokinase 1 (PFK1) binds to the transcriptional coactivator TEADS to stabilizes YAP/TAZ. PFK3B binds to Cdk1, cyclin D3 and Cdc25C. PFK1 product fructose-2,6bisphosphate is used for the phosphorylation of p27 by Cdk1, undergoing p27 degradation.

In this Special Issue, we would like to publish new papers related to "Advance in Transcriptional Regulation by Conventional Metabolic Enzymes".

- metabolic enzymes
- nuclear translocation
- transcriptional regulation
- epigenetic modification
- acetylation of DNA and histone
- methylation of NA and histone





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Message from the Editor-in-Chief

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