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Identification of Potential Links in Biomedical Networks

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Message from the Guest Editors

Dear Colleagues,

Predicting the potential associations in the biomedical bipartite network can provide useful insights for the diagnosis and treatment of complex diseases and the discovery of new drug targets.

This Special Issue is about solving potential link (association) prediction in biomedical networks. Areas to be covered in this Special Issue may include but are not limited to:

Gene–disease network
Drug–disease network
Drug–target network
Drug–side effect network
Drug–drug network
Protein–protein interaction network
circRNA–disease network
microRNA–disease network
lncRNA–disease network
lncRNA–protein interaction network

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Guest Editors



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Special Issue



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Message from the Editor-in-Chief

Genes are central to our understanding of biology, and modern advances such as genomics and genome editing have maintained genetics as a vibrant, diverse and fastmoving field. There is a need for good quality, open access journals in this area, and the *Genes* team aims to provide expert manuscript handling, serious peer review, and rapid publication across the whole discipline of genetics. Starting in 2010, the journal is now well established and recognised.

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