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Identification of Potential Links in Biomedical Networks

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Deadline for manuscript
submissions:
closed (25 August 2022)

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



mdpi.com/si/102322

Special Issue



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Message from the Editor-in-Chief

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