



MOL2NET, International Conference Series on Multidisciplinary Sciences

Note on: A Computational Toxicology Approach to Screen the Hepatotoxic Ingredients in Traditional Chinese Medicines

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Fragment

MOL2NET Conference highlights fragments of abstracts published in special issues if journals associated to the conference. This is a fragment of the abstract of the original article that belongs to: A Computational Toxicology Approach to Screen the Hepatotoxic Ingredients in Traditional Chinese Medicines: Polygonum multiflorum Thunb as a Case Study by Shuaibing He, Xuelian Zhang, Shan Lu, Ting Zhu, Guibo Sun and Xiaobo Sun *Biomolecules* 2019, 9(10), 577; <https://doi.org/10.3390/biom9100577> - 07 Oct 2019.

In recent years, liver injury induced by Traditional Chinese Medicines (TCMs) has gained increasing attention worldwide. Assessing the hepatotoxicity of compounds in TCMs is essential and inevitable for both doctors and regulatory agencies. However, there has been no effective method to screen the [...] Read more. (This article belongs to the Special Issue Big Data Analysis in Biomolecular Research, Bioinformatics, and Systems Biology with Complex Networks and Multi-Label Machine Learning Models)

References

Reference (Read Full Paper Free):

Biomolecules 2019, 9(10), 577; <https://doi.org/10.3390/biom9100577>

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