

Accepted Manuscript

Title: *In silico* assessment of adverse drug reactions and associated mechanisms

Author: Sergey M. Ivanov Alexey A. Lagunin Vladimir V. Poroikov



PII: S1359-6446(15)00289-5
DOI: <http://dx.doi.org/doi:10.1016/j.drudis.2015.07.018>
Reference: DRUDIS 1663

To appear in:

Received date: 13-4-2015
Revised date: 15-7-2015
Accepted date: 31-7-2015

Please cite this article as: Ivanov, S.M., Lagunin, A.A., Poroikov, V.V., *In silico* assessment of adverse drug reactions and associated mechanisms, *Drug Discovery Today* (2015), <http://dx.doi.org/10.1016/j.drudis.2015.07.018>

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

Highlights:

- Early assessment of adverse drug reactions (ADRs) requires knowledge of their mechanisms
- Chemical and biological drug features can be used for assessment of mechanisms
- Predicted protein targets, genes, pathways, chemical fragments are used in ADR analysis
- Network and pathway analysis reveal true associations of features with ADR mechanisms
- Predictive models for adverse reactions can be created based on revealed features

Download English Version:

<https://daneshyari.com/en/article/10885683>

Download Persian Version:

<https://daneshyari.com/article/10885683>

[Daneshyari.com](https://daneshyari.com)