



Computer-Aided Discovery of Hidden Pharmacotherapeutic Potential in Phytoconstituents from Traditional Medicine

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Computer-Aided Discovery of Hidden Pharmacotherapeutic Potential in Phytoconstituents from Traditional Indian Medicine (TIM)

Supported by RFBR-DST grant No. 11-04-92713-IND_a/RUSP-1176

Meeting with Prof. Rajesh K. Goel at CRDRI (Feb. 2010)















Some Information About Natural Compounds

- ✓ At the end of XX century about 80 percent of the world population to some extend are using natural compounds as medicines¹.
- ✓ Pharmaceuticals of vegetable or microbial origin make up more than 30% of global sales².
- ✓ About 70% of NCEs were obtained on the basis of natural products in 1981-2006³.

¹ World Health Organization

² Sneader W. Drug Discovery: A History. Wiley: Chichester, 2005, 88-150.

³ Newman D.J., Cragg G.M. J. Nat. Prod., 2007, 70:461-477.

Advantages of Natural Compounds

✓ Greater chemical diversity in comparison with the compounds obtained using only synthetic methods.

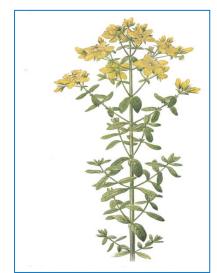
✓ Better ADME/T characteristics.







Гинкго двулопастной Ginkgo biloba



Зверобой продырявленный Hypericum perforatum



Черника обыкновеннаяVaccinium myrtillus



Эхинацея пурпурная Echinacea purpurea

Traditional Indian Medicine (Ayurveda)

- ✓ A lot of empirical knowledge about pharmacotherapeutic properties of natural products is accumulated in Traditional Indian Medicine (TIM) Ayurveda, which is known earlier than 1000 years BC.
- ✓ Some Ayurvedic plants are included into the list of national Indian priorities.







Problems with Study of Phytocomponents

- ✓ Utilization of extracts in a folk medicine.
- ✓ The difficulty of separation of each component in pure form.
- ✓ Changes in composition of components depending
 on the location and time of collection.
- ✓ Pleiotropic (multitargeted) action.

Analysis of Ayurvedic (TIM) Medicines Using In Silico Approaches

The empirical knowledge contained in Ayurveda can be currently analyzed using modern computer-aided methods.

Such studies could give information about the basic mechanisms of TIM actions, providing the basis for rational design of new medicinal plant combinations, and identification of new lead compounds for future pharmaceuticals.

Some Publications on Computational Studies of TIM



Evidence-Based Complementary and Alternative Medicine Volume 2013, Article ID 627375, 25 pages http://dx.doi.org/10.1155/2013/62737

Volume 29. Issue Number 4. (2012)

©Adenine Press (2012)

Review Article

New Perspectives on How to Discover Drugs from Herbal Medicines: CAM's Outstanding Contribution to Modern Therapeutics

Si-Yuan Pan, 1 Shu-Feng Zhou, 2 Si-Hua Gao, 3 Zhi-Ling Yu, 4 Shuo-Feng Zhang, Min-Ke Tang, Jian-Ning Sun, Dik-Lung Ma, 5 Yi-Fan Han,6 Wang-Fun Fong,4 and Kam-Ming Ko

School of Chinese Medicine, Beijing University of Chinese Medicine, Beijing 100102, China

Evidence-Based Complementary and Alternative Medicine

- ² College of Pharmacy, University of South Florida, Tampa, FL 33612, USA
- ³ School of basic medicine, Beijing University of Chinese Medicine, Beijing 100102, China School of Chinese Medicine, Hong Kong Baptist University, Hong Kong

Journal of Biomolecular Structure & ng Kong Polytechnic University, Hong Kong Dynamics, ISSN 0739-1102

Technology, Hong Kong -pan@163.com

host proteins during tuberculosis infection

Molecular Biology Laboratory, Department of Biochemistry and Molecular Biology and Department of Genetic Engineering and

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Volume 8(21)

Docking studies and network analyses reveal

capacity of compounds from Kandelia rheedii to

strengthen cellular immunity by interacting with

Aubhishek Zaman

open access

lesh; Aubhishek Zaman - Email: aubhishek@gmail.com; Phone

Volume 2013, Article ID 781216, 12 pages http://dx.doi.org/10.1155/2013/781216 Research Article

Hindawi Publishing Corporation

In Vitro and In Vivo Evaluation of Polyherbal Formulation against Russell's Viper and Cobra Venom and Screening of **Bioactive Components by Docking Studies**

und and used in Indian subcontinent, is a well-known herbal cure to omponents of the plant extract responsible for mediating this action tions of three compounds (emodin, fusaric acid and skyrin) from the NK), estrogen receptor (ERBB), dopamine β-hydroxylase (DBH) and ets are known to be responsible for strengthening cellular immunity se three compounds with the respective protein targets have been designing molecular medicines against tuberculosis

Computational Evidence to Inhibition of Human Acetyl Cholinesterase by Withanolide A for Alzheimer Treatment

http://www.jbsdonline.com

Alzheimer's disease (AD) a neurodegenerative disorder is the most common cause of

G. Sakthivel, 1,2 Amitabha Dey, 2 Kh. Non N. Surjit Singh,2 and Lokesh Deb2

Department of Nanotechnology, Noorul Islam Centre fo ² Pharmacology Laboratory, Medicinal Plants & Horticulti Department of Biotechnology, Government of India, Tak

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Bioorganic & Medicinal Chemistry 19 (2011) 6779-679 Contents lists available at SciVerse ScienceDirect

Bioorganic & Medicinal Chemistry

journal homepage: www.elsevier.com/locate/bmc



Hypothesis

Drug Design, Development and Therapy

Dovepress

CHEMICAL INFORMATION



Development of QSAR model immunomodulatory activity of coumarinolignoids

> This article was published in the following Dove I Drug Design, Development and Therapy

Number of times this article has been viewed

Dharmendra K Yadav Abha Meena Ankit Srivastava D Chanda Feroz Khan

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Open Access Full Text Article

paradigm.

SK Chattopadhyay Metabolic and Structural Biology Department, Central Institute of Medicinal and Aromatic Plants. Council of Scientific and Industrial Research, PO-CIMAP, India

Abstract: Immunomodulation is the p intrusion of molecules inside the body. drugs are promoted in traditional Indian coumarinolignoids isolated from the se hepatoprotective action and have recentl activity affecting both cell-mediated an modulatory compound from derivative relationship (QSAR) and molecular doc accord with the in vivo experimental d activity was predicted through OSAR m sion method with leave-one-out approx was 99% ($R^2 = 0.99$) and predictive acc that dipole moment, steric energy, amirefractivity correlates well with biolog energy, and molar refractivity has neg binding affinity to immunomodulatory

Keywords: coumarinolignoids, immu

Chemogenomics Approaches to Rationalizing the Mode-of-Action of Traditional Chinese and Ayurvedic Medicines

Fazlin Mohd Fauzi, II. Alexios Koutsoukas, Robert Lowe, Kalpana Joshi, Tai-Ping Fan, Robert C. Glen, and Andreas Bender and

Unilever Centre for Molecular Science Informatics, Department of Chemistry, University of Cambridge, Lensfield Road, Cambridge CB2 1EW, United Kingdom

[†]Universiti Teknologi MARA (U/TM) Malaysia, 40 450 Shah Alam, Selangor, Malaysia

[‡]Blizard Institute of Cell and Molecular Science, Barts and The London School of Medicine and Dentistry, The Blizard Building, 4 Newark Street, London E1 2AT, United Kingdom

Symbiosis School of Biomedical Sciences, Symbiosis International University, Pune, India

Department of Pharmacology, University of Cambridge, Tennis Court Road, Cambridge CB2 1PD, United Kingdom

3 Supporting Information

AND MODELING

ABSTRACT: Traditional Chinese medicine (TCM) and Ayurveda have been used in humans for thousands of years. While the link to a particular indication has been established in man, the mode-of-action (MOA) of the formulations often remains unknown. In this study, we aim to understand the MOA of formulations used in traditional medicine using an in silico target prediction algorithm, which aims to predict protein targets (and hence MOAs), given the chemical structure of a compound. Following this approach we were able to establish several links between suggested MOAs and rimental evidence. In particular, compounds from the 'tonifying and replenishing medicinal' class from TCM exhibit a hypoglycemic effect which can be related to activity of the ingredients against the Sodium-Glucose Transporters (SGLT) 1 and 2 as well as Protein Tyrosine Phosphatase (PTP). Similar results were obtained for Ayurvedic anticancer drugs. Here, both primary anticancer targets (those directly involved in cancer andsances usings (1975), routh plantaly anticolates trapple (1920) much with plantal machine participation and participation and a strender 3-dipharenductuse I and 2 over predicted as well as targets which act synergistically with the primary target, such as the eillast pump Peglycoprotein
(Peg). In addition, we were all to the clusticales some targets which may point us to novel MOAs as well as explain side effects.



Most notably, GPBAR1, which was predicted as a target for both 'tonifying and replenishing medicinal' and anticancer classes, suggests an influence of the compounds on metabolism. Understanding the MOA of these compounds is beneficial as it provides a resource for NMEs with possibly higher efficacy in the clinic than those identified by single-target biochemical assays.

Pharmacophore-based discovery of FXR-agonists. Part II: Identification of bioactive triterpenes from Ganoderma lucidum

Ulrike Grienke a, Judit Mihály-Bison b, Daniela Schuster c, Taras Afonyushkin b, Markus Binder b, Shu-hong Guan d, Chun-ru Cheng d, Gerhard Wolber e, Hermann Stuppner a, De-an Guo d, Valery N. Bochkov b, Judith M. Rollinger a,*

*Institute of Pharmacy/Pharmacognosy and Center for Molecular Biosciences Innsbruck, University of Innsbruck, Innrain 52c, 6020 Innsbruck, Austria Center of Biomolecular Medicine and Pharmacology, Department of Vascular Biology and Thrombosis Research, Medical University of Vienna, Schwarzspanierstraße 17,

⁶Computer-Aided Molecular Design Group, Institute of Pharmacy/Pharmaceutical Chemistry and Center for Molecular Biosciences Innsbruck, University of Innsbruck

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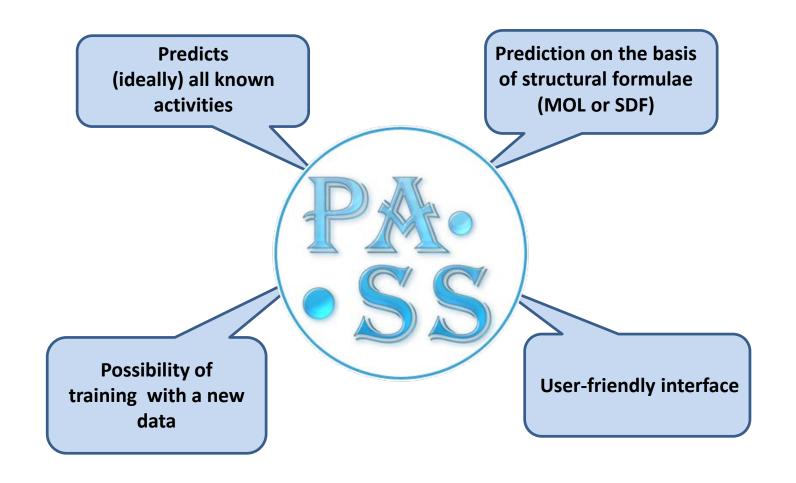
Keywords: Farnesoid X receptor Ganoderma lucidum Ganoderic acids Molecular modeling Virtual screening Natural products

ABSTRACT

The farnesoid X receptor (FXR) belonging to the metabolic subfamily of nuclear receptors is a ligandinduced transcriptional activator. Its central function is the physiological maintenance of bile acid homeostasis including the regulation of glucose and lipid metabolism. Accessible structural information about its ligand-binding domain renders FXR an attractive target for in silico approaches. Integrated to natural product research these computational tools assist to find novel bioactive compounds showing beneficial effects in prevention and treatment of, for example, the metabolic syndrome, dyslipidemia, atherosclerosis, and type 2 diabetes. Virtual screening experiments of our in-house Chinese Herbal Medicine database with structure-based pharmacophore models, previously generated and validated revealed mainly lanostane-type triterpenes of the TCM fungus Ganoderma lucidum Karst, as putative FXR ligands. To verify the prediction of the in silico approach, two Ganoderma fruit body extracts and compounds isolated thereof were pharmacologically investigated. Pronounced FXR-inducing effects were observed for the extracts at a concentration of 100 µg/mL. Intriguingly, five lanostanes out of 25 secondary metabolites from G. lucidum, that is, ergosterol peroxide (2), lucidumol A (11), ganoderic acid TR (12) ganodermanontriol (13), and ganoderiol F (14), dose-dependently induced FXR in the low micromola range in a reporter gene assay. To rationalize the binding interactions, additional pharmacophore profiling and molecular docking studies were performed, which allowed establishing a first structure-activity relationship of the investigated tritemenes

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Requirements for a computer program evaluated biological activity profiles (spectra)

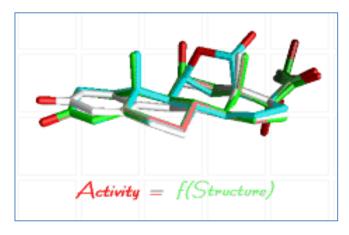


Biological activity spectra of organic compound

Biological activity is one of the most important characteristics of organic compound, which provides the basis for its use in therapeutic purposes. Biological activity reflects the result of interaction between the substance and biological object, and depends on substance structure and properties, biological object (species, sex, age), and mode of action (administration route, dose). Biological activity spectrum of an organic compound is the set of different kinds of biological activity that reflect the results of the compound's interaction with various biological entities. It represents the "intrinsic" property of a substance depending only on its structure. This is a qualitative characteristic property of a substance that depends only on its molecular structure.

Poroikov V. et al. *Automatic Documentation and Mathematical Linguistics*, 1993, 27: 40-43. Filimonov D. *Chemistry of Heterocyclic Compounds*, 2014, No. 3, 483-499.

Structure-activity relationships: (Q)SAR

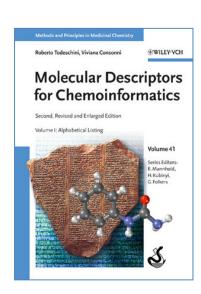


Molecular descriptors

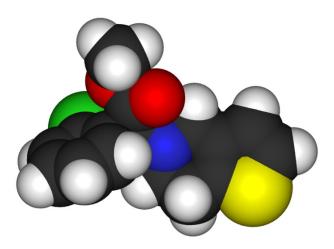
Sub-structural (-COO, -NH2, -OH, C6H5, и др.); physical-chemical (molecular weight, melting point, IR frequencies, chemical shifts in NMR, etc.); molecular connectivity, Wiener indices, Balaban indices, hydrophobicity constant, pKa, van der Waals volume, Log P, water solubility, etc. (several thousand).

Mathematical methods

Multiple linear regression (MLR); non-linear regression; partial least squares (PLS); regression on principal components (PCR); artificial neural networks (ANN); similarity matrices; genetic algorithms; support vector machine (SVM); cluster analysis (CA); discriminant analysis; etc.



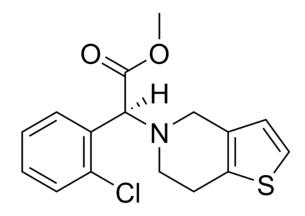
Chemical structure representation



The spatial configuration of the free uncharged molecules in the ground state in a vacuum is a necessary and sufficient description of its structure.

The use of this molecular structure description requires substantial computational resources for molecular modeling and/or quantum-chemical calculations.

However, the basis of all calculations is the traditional structural formula.



Thus, the structural formula uniquely determines all properties of the organic molecule.

Influence of the environment?

- Structural formula determines, at least, potential "intrinsic" properties of the molecule.

Neighborhoods of atoms descriptors

The most biological activities of organic compounds are the result of molecular recognition, which in turn depends on the correspondence between particular atoms of the ligand and the target.

MOLECULAR BIOLOGY QUANTUM CHEMISTRY QUANTUM FIELDS THEORY

$$\mathbf{M} = \mathbf{V} + \mathbf{V}\mathbf{g}\mathbf{M} = \mathbf{V} + \mathbf{V}\mathbf{g}\mathbf{V} + \mathbf{V}\mathbf{g}\mathbf{V}\mathbf{g}\mathbf{V} + \mathbf{V}\mathbf{g}\mathbf{V}\mathbf{g}\mathbf{V}\mathbf{g} + \dots$$
$$\mathbf{M}_{i} = \mathbf{V}_{i} + \mathbf{V}_{i}\mathbf{g}\mathbf{M} = \mathbf{V}_{i} + \mathbf{V}_{i}\mathbf{g}(\mathbf{M}_{1} + \mathbf{M}_{2} + \dots + \mathbf{M}_{m})$$

D.A. Filimonov

Descriptors are based on the concept of atoms' of molecule taking into account the influence of the neighborhoods:

MNA - Multilevel Neighborhoods of Atoms

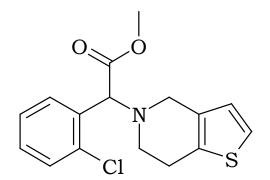
QNA - **Quantitative Neighborhoods** of **Atoms**

LMNA - Labeled Multilevel Neighborhoods of Atoms

Filimonov D.A., Poroikov V.V. *In: Chemoinformatics Approaches to Virtual Screening*. Eds. Alexandre Varnek and Alexander Tropsha. Cambridge (UK): RSC Publishing, 2008, 182-216. Filimonov D.A. et al. *SAR and QSAR Environ. Res.*, 2009, 20: 679-709. Rudik A.V. et al. *J. Chem. Inform. Model.*, 2014, 54: 498–507.

Substance representation: Clopidogrel

Structural formula



Activity Spectrum

Abdominal pain

Acute neurologic disorders treatment

Agranulocytosis

Allergic reaction

Anaphylaxis

Anemia

Angioedema

Angiogenesis inhibitor

Antianginal

Antiarthritic

Anticoagulant

Antineoplastic

Antipsoriatic

Antithrombotic

MNA Descriptors (1st and 2nd levels)

 $\begin{array}{lll} \text{HC} & \text{C(C(CCC)C(CC-H-H)S(CC))} \\ \text{CHHHO} & \text{C(C(CCC)C(CS-H)-H(C))} \\ \text{CHHCC} & \text{C(C(CCC)N(CC-C)-H(C)-H(C))} \\ \text{CHHCN} & \text{C(C(CCS)C(CC-H)C(CN-H-H))} \\ \text{CHCC} & \text{C(C(CCS)C(CN-H-H)-H(C)-H(C))} \\ \text{CHCCN} & \text{C(C(CC-H-H)N(CC-C)-H(C)-H(C))} \end{array}$

CHCS C(C(CC-H)C(CC-H)-H(C))
CCCC C(C(CC-H)C(CC-C)-H(C))
CCCS C(C(CC-H)C(CC-C)-CI(C))
CCCCI C(C(CC-H)C(CC-CI)-H(C))

CCOO C(C(CC-H)C(CC-CI)-C(CN-H-C))

NCCC C(C(CC-H)S(CC)-H(C))

OC N(C(CN-H-H)C(CN-H-H)-C(CN-H-C))

OCC S(C(CCS)C(CS-H))

SCC -H(C(CC-H)) CIC -H(C(CC-H-H))

-H(C(CN-H-H)) -H(C(CS-H))

-H(-C(CN-H-C))

-H(-C(-H-H-H-O))

-C(C(CC-C)N(CC-C)-H(-C)-C(-C-O-O))

-C(-H(-C)-H(-C)-H(-C)-O(-C-C)) -C(-C(CN-H-C)-O(-C)-O(-C-C)) -O(-C(-H-H-H-O)-C(-C-O-O))

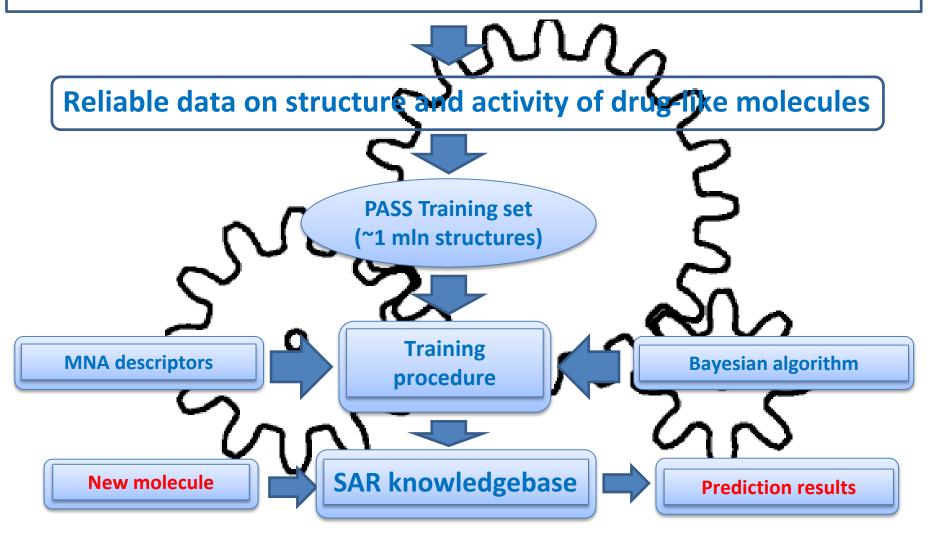
-O(-C(-C-O-O))

-CI(C(CC-CI))

112 known activities in PASS SAR Base

PASS: Prediction of Activity Spectra for Substances

Full text publications, databases, presentations at conferences etc.



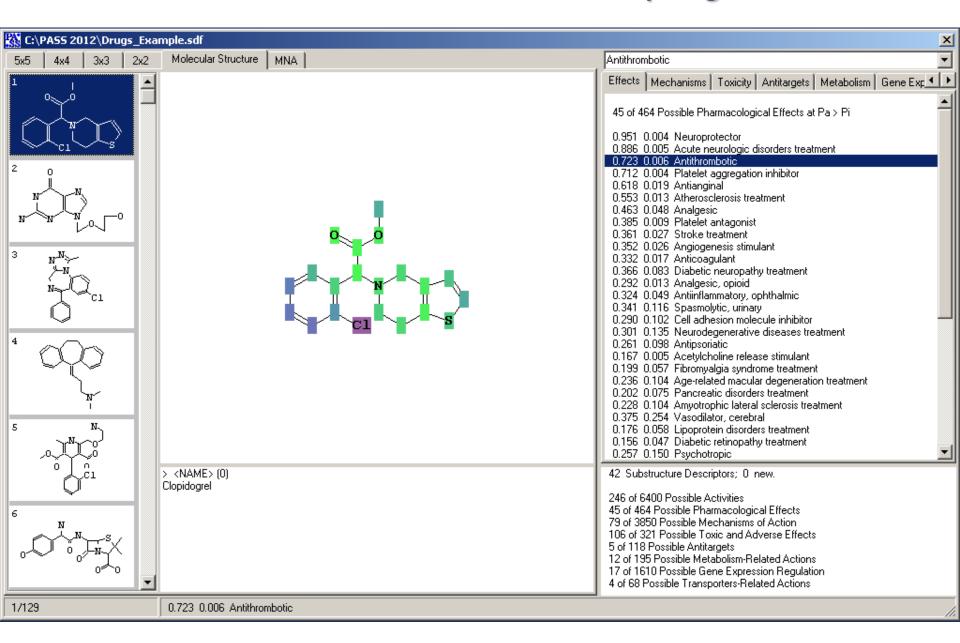
Filimonov D.A. et al. Chem. Heterocycl. Comps., 2014, 50: 444-457.

PASS 2014 Characteristics

Training Set	959,801 drugs, drug-candidates, pharmacological and toxic substances comprise the training set
Biological Activity	7,158 biological activities can be predicted (Active vs. Inactive)
Chemical Structure	Multilevel Neighborhoods of Atoms (MNA) descriptors [1, 2]
Mathematical Algorithm	Bayesian approach was selected by comparison of many different methods [2]
Validation	Average accuracy of prediction in LOO CV for the whole training set is ~95% [2]; robustness was shown using principal compounds from MDDR database [3]

- 1. Filimonov D.A. et al. J. Chem. Inform. Computer Sci., 1999, 39: 666-670.
- 2. Filimonov D.A., Poroikov V.V. Chemoinformatics Approaches to Virtual Screening, 2008, 182-216.
- 3. Poroikov V.V. et al. J. Chem. Inform. Computer Sci., 2000, 40: 1349-1355.

Results of PASS Prediction for Clopidogrel



Results of PASS Prediction for Clopidogrel

Abdominal pain

Acute neurologic disorders

treatment

Agranulocytosis

Allergic reaction

Anaphylaxis

Anemia

Angioedema

Angiogenesis inhibitor

Antianginal Antiarthritic

Anticoagulant

Antineoplastic

Antipsoriatic

Antithrombotic

Anxiety

Arthralgia

Atherosclerosis treatment

Back pain

Behavioral disturbance

Blindness

Bronchoconstrictor

Cardiotoxic

Cataract

CCL4 expression enhancer

CCL5 expression enhancer

Chest pain

Colic Colitis

Conjunctivitis

Consciousness alteration

Constipation

Cough

CYP2 substrate

CYP2C substrate

CYP2C19 inhibitor CYP2C19 substrate

CYP2C9 inhibitor

CYP3A substrate

CYP3A4 substrate

Cytochrome P450 inhibitor

Dermatitis

Dermatologic

Dizziness

Drug eruption Dyspepsia

Emetic

Eosinophilia

Erythema multiforme

Exanthema

Flatulence

GP IIb/IIIa receptor antagonist

Hallucinogen

Headache

Heart failure

Hematotoxic

Hemorrhage

Henoch-Schonlein purpura

Hepatic failure

Hepatitis

Hepatotoxic

Hypertensive

Hyperthermic

Hypotension

Infection Insomnia

Lassitude

Leukopenia

Lichen planus

Lichenoid eruption

Malaise

Menstruation disturbance

Myalgia Nausea

Necrosis

Nephrotoxic Neuroprotector

Neutropenia .

Ocular toxicity

Pain

Pancreatitis

Pancytopenia

Platelet aggregation inhibitor

Platelet antagonist

Pruritus

Pulmonary embolism

Purinergic P2 antagonist

Purinergic P2T antagonist

Purinergic P2Y antagonist

Purinergic P2Y12 antagonist

Purinergic receptor antagonist

Purpura

Renal colic

Reproductive dysfunction

Rhinitis

Sensory disturbance

Serum sickness

Shock

Sinusitis

Sleep disturbance

Stomatitis

Syncope

THBS1 expression enhancer

Thrombocytopenia

Toxic

Toxic epidermal necrolysis

Toxic, gastrointestinal

TP53 expression enhancer

Urticaria

Vasculitis

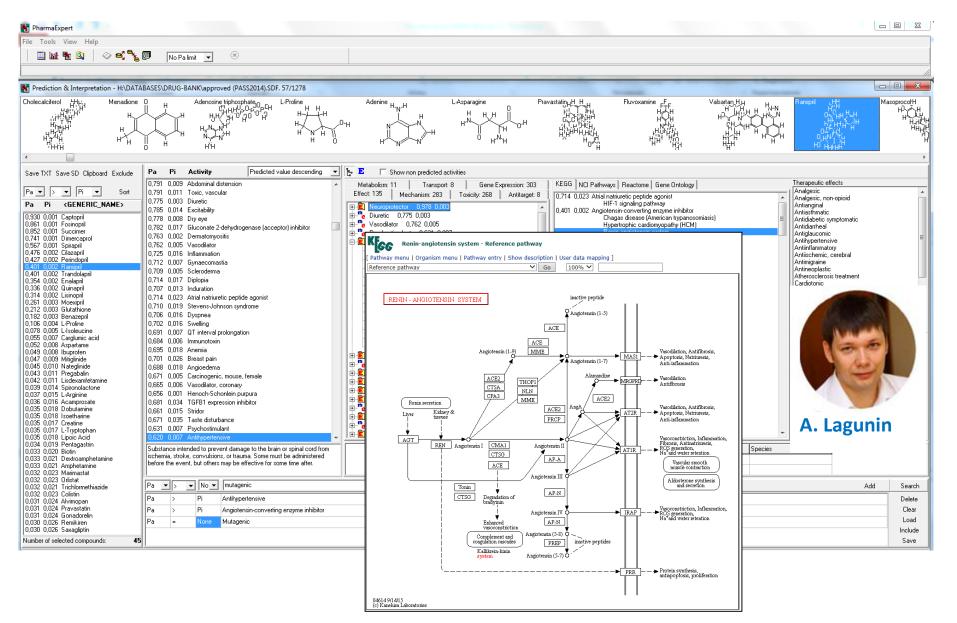
Vertigo

Vision disturbance

Blue – predictions coincided with the experiment.

Black – unpredictable activities. Red – unpredicted activities.

PharmaExpert: Interpretation of the prediction results



Web-services based on our methods





D. Druzhilovsky



A. Rudik



A. Zakharov

www.way2drug.com/Projects.php

Some examples of practical applications of biological activity spectra prediction



Athina Geronikaki, AUT, Greece

Search for multitarget drugs

J. Med. Chem. 2008, 51: 1601.





Rajesh Goel, PU, India



Drug reporpusing

Pharm. Chem. J. 2011, 45: 605.



Sergey Kryzhanovsky, Inst. of Pharmacol., Russia

Estimating drug-drug interactions for phytoconstituents of medicinal plants

Nat. Prod. Rep., 2014, 31: 1585.

Examples of search for new compounds based on predictions

J. Chem. Inf. Comput. Sci. 2003, 43, 228-236

PASS Biological Activity Spectrum Predictions in the Enhanced Open NCI Database Browser

Vladimir V. Poroikov,[‡] Dmitrii A. Filimonov,[‡] Wolf-Dietrich Ihlenfeldt,[#] Tatyana A. Gloriozova,[‡] Alexey A. Lagunin,[‡] Yulia V. Borodina,[‡] Alla V. Stepanchikova,[‡] and Marc C. Nicklaus*,[†]

Laboratory of Structure-Function Based Drug Design, V.N. Orekhovich Institute of Biomedical Chemistry of the Russian Academy of Medical Sciences, 10 Pogodinskaya Street, Moscow 119121, Russia, Computer Chemistry Center and Institute for Organic Chemistry, University of Erlangen-Nürnberg, Nägelsbachstrasse

2870

J. Med. Chem. 2004, 47, 2870-2876

Design of New Cognition Enhancers: From Computer Prediction to Synthesis and Biological Evaluation

Athina A. Geronikaki,*,† John C. Dearden,‡ Dmitrii Filimonov,§ Irina Galaeva," Taissia L. Garibova," Tatiana Gloriozova,§ Valentina Krajneva," Alexey Lagunin,§ Fliur Z. Macaev,↓ Guenadij Molodavkin," Vladimir V. Poroikov,§ Serghei I. Pogrebnoi,↓ Felix Shepeli,↓ Tatiana A. Voronina," Maria Tsitlakidou,† and Liudmila Vlad↓

School of Pharmacy, Department of Pharmaceutical Chemistry, Aristotle University of Thessaloniki, Thessaloniki, Greece,

3326 J. Med. Chem. 2003, 46, 3326-3332

Computer-Aided Selection of Potential Antihypertensive Compounds with Dual Mechanism of Action

Alexey A. Lagunin,* Oleg A. Gomazkov, Dmitrii A. Filimonov, Tatyana A. Gureeva, Elvira A. Dilakyan, Elena V. Kugaevskaya, Yulia E. Elisseeva, Nina I. Solovyeva, and Vladimir V. Poroikov

Institute of Biomedical Chemistry of Russian Academy of Medical Sciences, Pogodinskaya Street, 10, Moscow 119121, Russia

J. Med. Chem. 2008, 51, 1601-1609

Computer-Aided Discovery of Anti-Inflammatory Thiazolidinones with Dual Cyclooxygenase/Lipoxygenase Inhibition

Athina A. Geronikaki, [†] Alexey A. Lagunin, ** Dimitra I. Hadjipavlou-Litina, [†] Phaedra T. Eleftheriou, [†] Dmitrii A. Filimonov, [‡] Vladimir V. Poroikov, [‡] Intekhab Alam, [§] and Anil K. Saxena [§]

European Journal of Medicinal Chemistry 47 (2012) 111-124

Contents lists available at SciVerse ScienceDirect

European Journal of Medicinal Chemistry

journal homepage: http://www.elsevier.com/locate/ejmech

Original article

Fragment-based design, docking, synthesis, biological evaluation and structure—activity relationships of 2-benzo/benzisothiazolimino-5-aryliden-4



Available online at www.sciencedirect.com

SCIENCE DIRECT®

Bioorganic & Medicinal Chemistry

Bioorganic & Medicinal Chemistry 12 (2004) 6559-6568

Design, synthesis, computational and biological evaluation of new anxiolytics

Athina Geronikaki, ^{a,*} Eugeni Babaev, ^b John Dearden, ^c Wim Dehaen, ^d Dmitrii Filimonov, ^e Irina Galaeva, ^f Valentina Krajneva, ^f Alexey Lagunin, ^e Fliur Macaev ^g Guenadiy Molodaykin ^f Vladimir Poroikov ^e Serghei Pogrebnoi ^g

Chemistry of Heterocyclic Compounds, Vol. 42, No. 5, 2006

SYNTHESIS AND ANTI-INFLAMMATORY ACTIVITY OF ETHYNYLTHIAZOLES

A. Geronikaki¹, S. Vasilevsky², D. Hadjipavlou-Litina¹, A. Lagunin, and B. V. Poroikov³

A series of acetylene derivatives of thiazole using the Sonogashira cross-coupling method was synthesized and evaluated in vivo for their anti-inflammatory activity. Four compounds exhibited good anti-inflammatory activity and two inhibited soybean lipoxygenase.



Available online at www.sciencedirect.com



European Journal of Medicinal Chemistry 44 (2009) 473-481



http://www.elsevier.com/locate/eimecl

Original article

Evaluation of the local anaesthetic activity of 3-aminobenzo[d]isothiazole derivatives using the rat sciatic nerve model

Athina Geronikaki ^{a,*}, Paola Vicini ^b, Nikos Dabarakis ^c, Alexey Lagunin ^d, Vladimir Poroikov ^d, John Dearden ^e, Hassan Modarresi ^e, Mark Hewitt ^e, George Theophilidis ^f

Current Pharmaceutical Design, 2010, 16, 1703-1717

1703

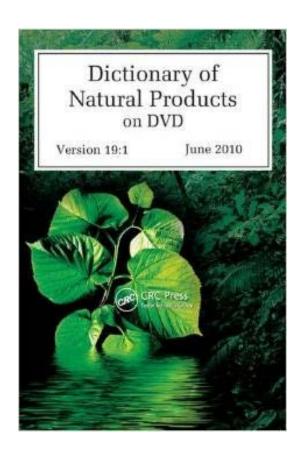
Multi-Targeted Natural Products Evaluation Based on Biological Activity Prediction with PASS

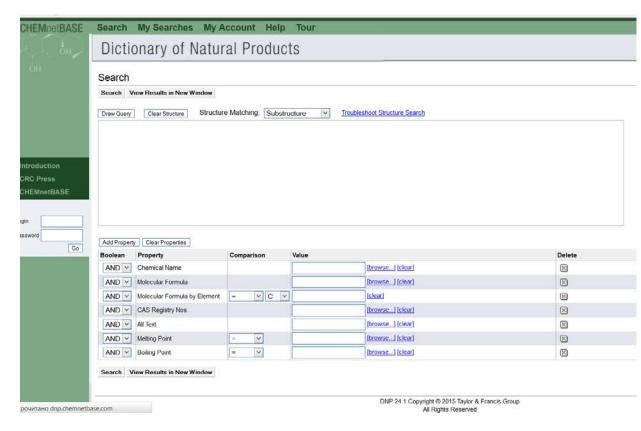
Alexey Lagunin, Dmitry Filimonov and Vladimir Poroikov*

Institute of Biomedical Chemistry of Rus. Acad. Med. Sci., 10, Pogodinskaya Str., Moscow, 119121, Russia

Abstract: Natural products found a wide use in folk medicine. Presently, when routine development of new drugs faced a considerable challenge, they become an inspiration and valuable source in drugs discovery. Rather complex and diverse chemical structures of natural compounds provide a basis for modulation of different biological targets. Natural compounds exhibit a multitargeted action that may lead to additive/synergistic or antagonistic effects. Rational design of more safe and potent pharmaceuticals requires an estimation of probable multiple actions of natural products. Our software PASS can perform such estimation. It predicts with reasonable accuracy over 3500 pharmacotherapeutic effects, mechanisms of action, interaction with the metabolic system, and specific toxicity for drug-like molecules on the basis of their structural formulae. We analyzed PASS predictions utilizing PharmaExpert, which performs selection of compounds with multiple mechanisms of action, analysis of activity-activity relationships and drug-drug interactions. The naner describes an

Sources of information about natural products: Dictionary of Natural Products

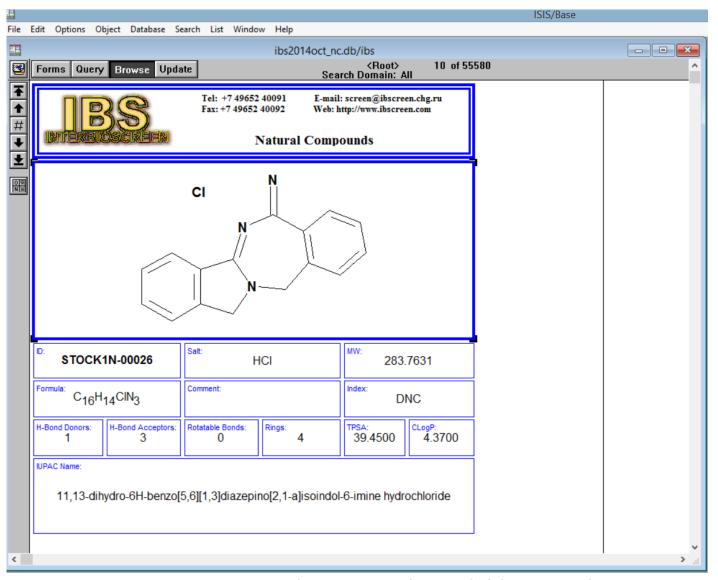




Chemical, physical, and biological data on more than 259,859 compounds contained in over 68,000 entries.

http://dnp.chemnetbase.com/

Sources of information about natural products: InterBioScreen Natural Compounds database



Over 55 000 natural compounds available in stock.

Sources of information about natural products: our Ayurvedic Medicines database



Natural products are used in folk medicine since many thousands year. They represent a significant, though often underappreciated resource for the development of new medicines.

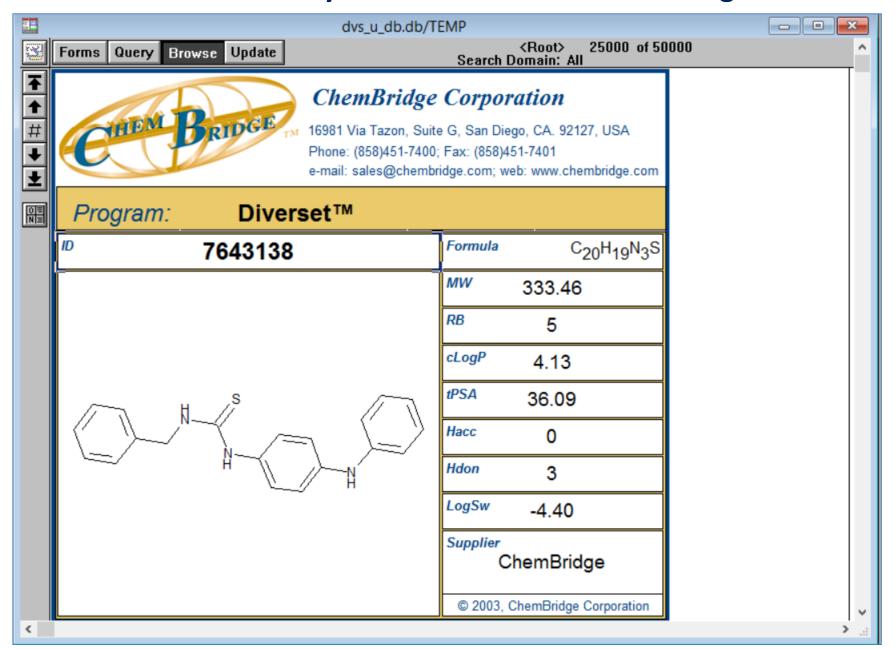
Content:

- (1) 50 medicinal plants;
- (2) structural formulae of 1906 phytochemicals;
- (3) biological activity of 288 phytochemicals.

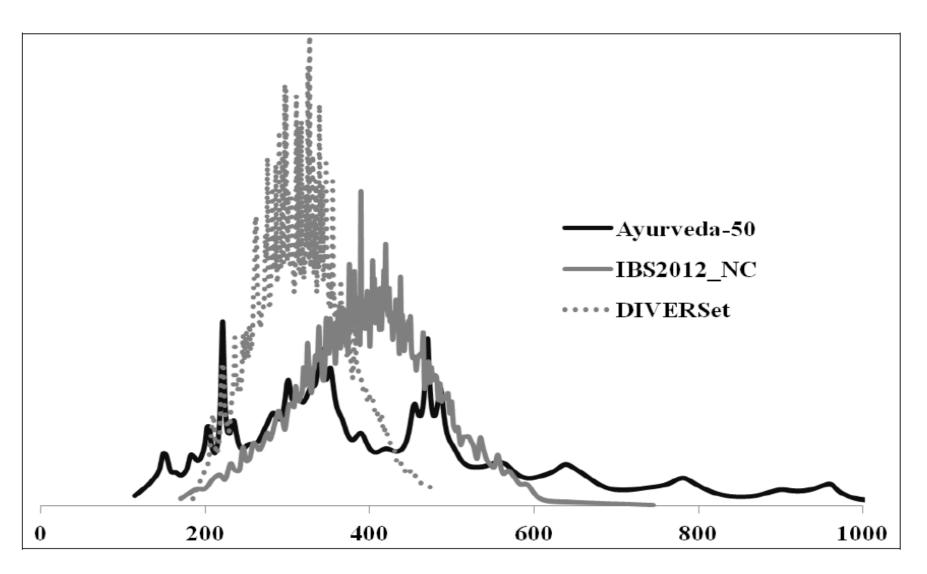
Criteria:

- (1) Ayurvedic /traditional medicinal use;
- (2) adequately explored for phytochemical analysis;
- (3) unexplored for pleiotropic pharmacological studies.

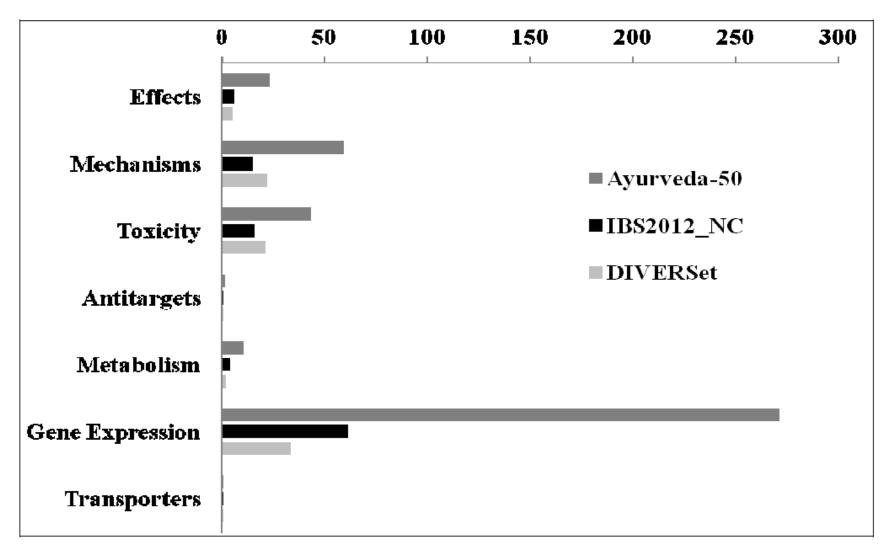
Reference set of synthetic molecules: ChemBridge DVS



Comparison of three chemical libraries: distribution by molecular weights

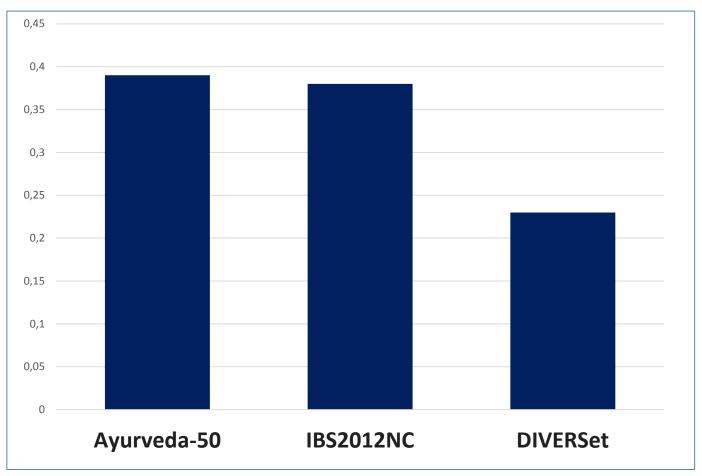


Comparison of three chemical libraries: average number of biological activities*



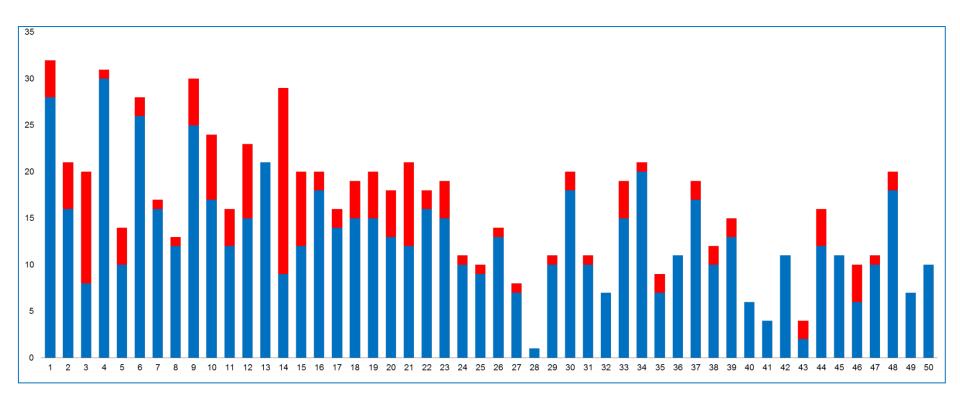
^{*}Predicted by PASS for one structure.

Comparison of three chemical libraries: pharmacological potential (PP)*



*PP = Number of predicted effects / Number of predicted mechanisms

Comparison of predicted and known activities for phytoconsituents from fifty TIM medicinal plants

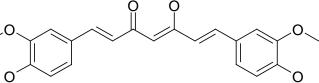


Blue - predicted; red - not predicted

Druzhilovskiy D., Rudik A., Ivanov S., Lagunin A., Filimonov D., Dinesh Gawande, Rajesh Kumar Goel, Vladimir Poroikov. (2012). Computer-aided analysis of hidden potential in traditional Indian medicine Ayurveda. Abstr. 19th EuroQSAR Symposium, Vienna, Austria, August 26-30, p. 128.

Exploited and Hidden Pharmacological Potential of Curcumin





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Review

Multi-targeted therapy by curcumin: how spicy is it?

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- ² Cytokine Research Laboratory, Department of Experimental Therapeutics, The University of Texas M. D. Anderson Cancer Center, Houston, TX, USA

Although traditional medicines have been used for thousands of years, for most such medicines neither the active component nor their molecular targets have been very well identified. Curcumin, a yellow component of turmeric or curry powder, however, is an exception. Although inhibitors of cycloxygenase-2, HER2, tumor necrosis factor, EGFR, Ber-abl, proteosome, and vascular endothelial cell growth factor have been approved for human use by the United States Food and Dny Administration (FDA), curcumin as a single agent can down-regulate all these targets. Curcumin can also activate apoptosis, down-regulate cell survival gene products, and up-regulate p53, p21, and p27. Although curcumin is poorly absorbed after ingestion, multiple studies have suggested that even low levels of physiologically achievable concentrations of curcumin may be sufficient for its chemopreventive and chemotherapeutic activity. Thus, curcumin regulates multiple targets (multitargeted therapy), which is needed for treatment of most diseases, and it is inexpensive and has been found to be safe in human clinical trials. The present article reviews the key molecular mechanisms of curcumin action and compares this to some of the single-targeted therapis; variety available for human acneer.

Keywords: Cancer / Curcumin / Cyclooxygenases / Multi-targeted therapy / Tumor necrosis factor Received: September 7, 2007; revised: Ocober 12, 2007; accepted: October 21, 2007



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Current Drug Targets, 2011, 12, 332-347

The Targets of Curcumin

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Abstract: Curcumin (diferuloylmethane), an orange-yellow component of turmeric or curry powder, is a polyphenol natural product isolated from the rhizome of the plant Curcuma longa. For centuries, curcumin has been used in some medicinal preparation or used as a food-coloring agent. In recent years, extensive in vitro and in vivo studies suggested curcumin has anticancer, antiviral, antiarthritic, anti-amyloid, antioxidant, and anti-inflammatory properties. The underlying mechanisms of these effects are diverse and appear to involve the regulation of various molecular targets, including transcription factors (such as nuclear factor-wB), growth factors (such as vascular endothelial cell growth factor), inflammatory cytokines (such as nuclear factor-wB), growth factors (such as vascular endothelial cell growth factor), inflammatory cytokines (such as nuclear factor-wB), growth factors (such as vascular endothelial cell growth factor), inflammatory cytokines (such as nuclear factor-wB), growth factors (such as vascular endothelial cell growth factor), inflammatory cytokines (such as cyclooxygenase 2 and 5 lipoxygenase). Thus, due to its efficacy and regulation of multiple targets, as well as its safety for human use curcumin has received considerable interest as a potential therapeutic agent for the prevention and or various malignant diseases, arthritis, allergies, Alzheimer's disease, and other inflammatory illnesses. This review summarizes various in vitro and in vivo pharmacological aspects of curcumin as well as the underlying action mechanisms. The recently identified undecular targets and signaling pathways modulated by curcumin are also discussed here.

Keywords: Curcumin, molecular targets, transcription factors, growth factors, inflammatory cytokines, protein kinases, enzymes.

PASS Prediction

0,933	0,003	Feruloyi esterase inhibitor	Data (17. 61 at., 2000
0,931	0,002	Ulcer, gastric Retinal oxidase inhibitor	
0,896	0,002	Beta-carotene 15,15'-monocxygenase inhibitor	
0,900	0,009	Beta-carotene 15,15'-monocrygenase inhibitor Aspulvinose dimethylallyttransferase inhibitor CVP3 inhibitor	Patel R., et al., 2007
0,893	0,005	Sneezing NADH dehydrogenase (ubiquinone) inhibitor Membrane integrity agonist Monophenol monooxygenase inhibitor	
0,897	0,012	NADH dehydrogenase (ubiquinone) inhibitor Membrane integrity agonist	
0,864	0,003	Monophenol monooxygenase inhibitor Non mutagenic, Salmonelia	Panich U. et al., 2010
0,851	0,003	Ulcer, poptic	
0,860	0,012	Mices, pegitic Antacid Historie acetyltransferase inhibitor	Israel M. et al. 2011
			Goel A. et al., 2008 Alhosin M. et a., 2011
0.922	0.000	Antieczematic	Alhosin M. et a., 2011 Yoon J-H. et al., 2005
0,834	0,005	Linoleate did synthase inhibitor CYP1A1 inhibitor	10017-11. 61 81., 2007
0,828	0,002	CYP1A1 inhibitor	
0,831	0,010	Reductant Gluconate 2-dehydrogenase (acceptor) inhibitor CVP3-k inducer	
0,822	0,003	CYP1A inducer Chlorde cone reductase inhibitor	
0,840	0,023	CPF A misuser Chlords cone reductase inhibitor Ubiquinol-cytochrome-c reductase inhibitor	
0,819	0,003	Choleretic GST P1-1 substrate	Deters M. et al., 2003
0,814	0,002	Vanillyl-alcohol oxidase inhibitor	
0,813	0,002	GSTP1:1 substrate VanillyI-alcohol oxidase inhibitor GSTP substrate HIF1A expression inhibitor	Choi H. et al., 2006
0,806	0,002	CYP1A1 inducer	
0,796	0,004	Ulceration Antimutagenic	Biswas J. et al, 2010
0,821	0,032	Subarachnoid hemorrhage GST A substrate	
0,823	0,041	Or or automatic Perphyria Steroid N-acetylglucosaminyltransferase inhibitor	
0,784	0,003	Steroid N-acetylglucosaminyltransferase inhibitor Antihypercholesterolemic	Deters M. et al., 2003
0.774	0.003	Leukotriene Cantagonist	Yoon J-H. et al., 2005
0,771	0,005	UDP-glucuronosytransferase substrate Apoptosis agonist	Goel A. et al., 2008
			Hatcher H. et al., 2008
0.74	0.008	Hepstellussendering Caspass Stimulant Caminative	Karmakar S. et al., 2007
0,742	0,004	Carminative	
0,741	0,004	Carminative MAP Kinase stimulant Mucomembranous protector	Suh HW et al., 2007
0,744	0,020	CVP2 substrate Fibrinolytic Aldehyde oxidase inhibitor	Ameye L.G. et al., 2005
0,728	0,018	Aldehyde oxidase inhibitor	
0,711	0,003	4-Coumarate-CoA ligase inhibitor CYP2D6 substrate	
		Membrane permeability inhibitor Sweating	Morin D. et al., 2001
0,733	0,034	Vasoprotector	
0,708	0,007	Chemopreventive	Goel A. et al., 2008
0,733	0,002	Glaucoma HIV-1 integrase inhibitor Chorea	Burke T. et al., 1995
0,721	0,023	Unine esscoioration Toxic, vascular Hematotoxic	
0,734	0,040	Hematotoxic	
0,700	0,010	Irritation Laryngospasm	
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In silico mining several medicinal plants with desirable pleiotropic (anticonvulsant, antidepressant, and nootropic) effects

No	Plant	Structure	Name	PASS predictions
1	Achyranthes aspera	N	3-Pyrrolidine- carboxylic acid; N- Me	0.547 0.098 Nootropic0.593 0.024 Anticonvulsant0.281 0.069 Antidepressant
2	Aerva lanata		6H-Indolo(3,2,1-de)(1,5)naphthyrid in-6-one, 10-methoxy-	0.687 0.041 Nootropic 0.285 0.138 Anticonvulsant 0.205 0.111 Antidepressant
3	Berberis vulgaris		Lambertine	0.386 0.231 Nootropic 0.326 0.109 Anticonvulsant 0.469 0.026 Antidepressant
4	Glycyrrhiza glabra		Cyclotetradecane	0.544 0.100 Nootropic 0.406 0.067 Anticonvulsant 0.218 0.102 Antidepressant

Predicted biological activities for major phytocomponents of Passiflora Incarnata: Harmine

0.674 0.045 Nootropic

0.698 0.004 5 Hydroxytryptamine 3A antagonist

0.557 0.081 Nerve growth factor agonist

0.469 0.031 Cyclic AMP phosphodiesterase inhibitor

0.499 0.091 Calcium channel (voltage-sensitive) activator

0.437 0.072 Cyclic AMP agonist

0.367 0.030 Calcium channel activator

0.116 0.336 Anticonvulsant

0.698 0.004 5 Hydroxytryptamine 3A antagonist

0.629 0.005 Cyclic AMP antagonist

0.438 0.038 Calmodulin antagonist

0.374 0.010 Benzodiazepine agonist partial

0.253 0.083 Antidepressant

0.745 0.023 5 Hydroxytryptamine release stimulant

0.698 0.004 5 Hydroxytryptamine 3A antagonist

0.557 0.081 Nerve growth factor agonist

0.414 0.046 5 Hydroxytryptamine 7 antagonist

0.351 0.039 Imidazoline receptor agonist

Predicted biological activities for major phytocomponents of Passiflora Incarnata: Harmaline

0.704 0.036 Nootropic

0.478 0.052 5 Hydroxytryptamine 3A antagonist

0.443 0.039 Cyclic AMP phosphodiesterase inhibitor

0.269 0.150 Anticonvulsant

0.478 0.052 5 Hydroxytryptamine 3A antagonist

0.440 0.037 Calmodulin antagonist

0.424 0.126 Cyclic AMP antagonist

0.394 0.062 5 Hydroxytryptamine 7 antagonist

0.344 0.048 Antidepressant

0.745 0.023 5 Hydroxytryptamine release stimulant

0.478 0.052 5 Hydroxytryptamine 3A antagonist

0.388 0.005 MAO inhibitor

0.409 0.027 Imidazoline receptor agonist

0.394 0.062 5 Hydroxytryptamine 7 antagonist

Predicted biological activities for major phytocomponents of Passiflora Incarnata: Harmalol

0.672 0.046 Nootropic

0.662 0.005 5 Hydroxytryptamine 3A antagonist

0.568 0.044 Calcium channel (voltage-sensitive) activator

0.409 0.050 Cyclic AMP phosphodiesterase inhibitor

0.330 0.034 Adrenaline release stimulant

0.245 0.170 Anticonvulsant

0.662 0.005 5 Hydroxytryptamine 3 antagonist

0.409 0.050 Cyclic AMP phosphodiesterase inhibitor

0.431 0.116 Cyclic AMP antagonist

0.425 0.048 Calmodulin antagonist

0.209 0.108 Antidepressant

0.844 0.011 5 Hydroxytryptamine release stimulant

0.662 0.005 5 Hydroxytryptamine 3 antagonist

0.387 0.005 MAO inhibitor

0.400 0.056 5 Hydroxytryptamine 7 antagonist

Revealing Medicinal Plants That Are Useful for the Comprehensive Management of Epilepsy and Associated Comorbidities through *In Silico* Mining of Their Phytochemical Diversity

Authors

Rajesh Kumar Goel¹, Dinesh Gawande¹, Alexey Lagunin², Puneet Randhawa¹, Awanish Mishra¹, Vladimir Poroikov²

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Orekhovich Institute of Biomedical Chemistry, Moscow, Russia

Key words

- Passifibra incarnata
- Passifloraceae
- epilepsy
- PASS
- PharmaExpert
- depression
- memory deficit

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Abstract

In silico techniques in drug discovery may rationalise and speed up the identification of lead molecules from nature. Drug discovery from medicinal plants has mostly been confined to indications in accordance with their ethnical use only. However, the availability of multiple phytoconstituents in medicinal plants suggests that these may be much more useful beyond their traditional uses and in the management of chronic diseases, along with their comorbidities. In this study, the computer programmes PASS and PharmaExpert were used to reveal the medicinal plants useful in the comprehensive management of epilepsy and associated psychiatric disorders based on the pleiotropic effects predicted for their phytoconstituents. In silico analysis revealed that seven of 50 medicinal plants from traditional Indian medicine possessed the desired pleiotropic effect, i.e., anticonvulsant, antidepressant, and nootropic activities, The majority of phytoconstituents from Passiflora incarnata were concurrently predicted to have the desired pleiotropic effects. Therefore, P. incarnata was pharmacologically validated using the pentylenetetrazole kindling mouse model, Behavioural and neurochemical evaluations confirmed the ameliorative role of P. incarnata in epilepsy and the associated depression and memory deficit. The pharmacological findings from this study propose that PASS and PharmaExpert may serve as good tools for the optimisation of the selection of plants based on their phytoconstituents

for the treatment of different ailments, even bevond their traditional use.

Abbreviations

AChE: acetylcholinesterase AUC: area under the curve

CPCSEA: Committee for the Purpose of Control and Supervision of Experiments on

Animals

NP: Dictionary of Natural Products

EPM: elevated plus maze FST: forced swim test

LOO CV: leave-one-out cross-validation MNA: multilevel neighbourhoods of atoms

MOA: mechanisms of action NO: nitric oxide

Pa: probability "to be active"

PASS: prediction activity spectra of

Pi: probability "to be inactive"
PIHE: Passiflora incarnata hydroethanolic

extrac

PTZ: pentylenetetrazole
PHT: phenytoin
SFZ: shock-free zone

TIM: traditional Indian medicine

TST: tail suspension test

Supporting Information available online at http://www.thieme-connect/products

Introduction

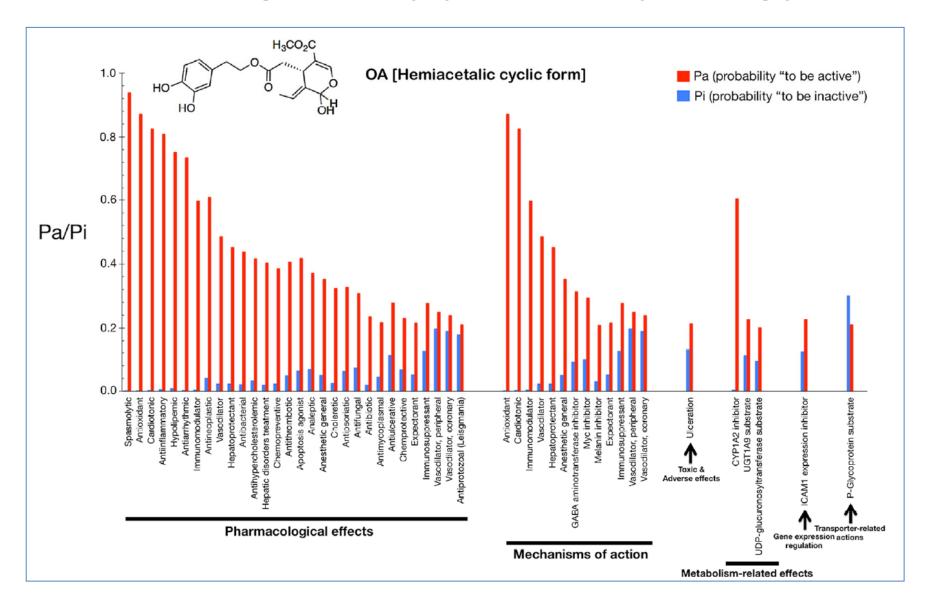
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Medicinal plants have been used for the treatment of various ailments in different systems of traditional medicine and have also served as a source for many drugs in modern medicine [1,2]. These plants have been explored either based on bioactivity-guided fractionation to identify bioactive principles for traditional activities of interest or based on the random exploration of the phytoconstituents without assigning any specific pharmacological activity [3, 4]. In any case, these explorations have led to enrichment of the phytochemical information about these medicinal

Computer-aided discovery of biological activity spectra for anti-aging and anti-cancer olive oil oleoproteins

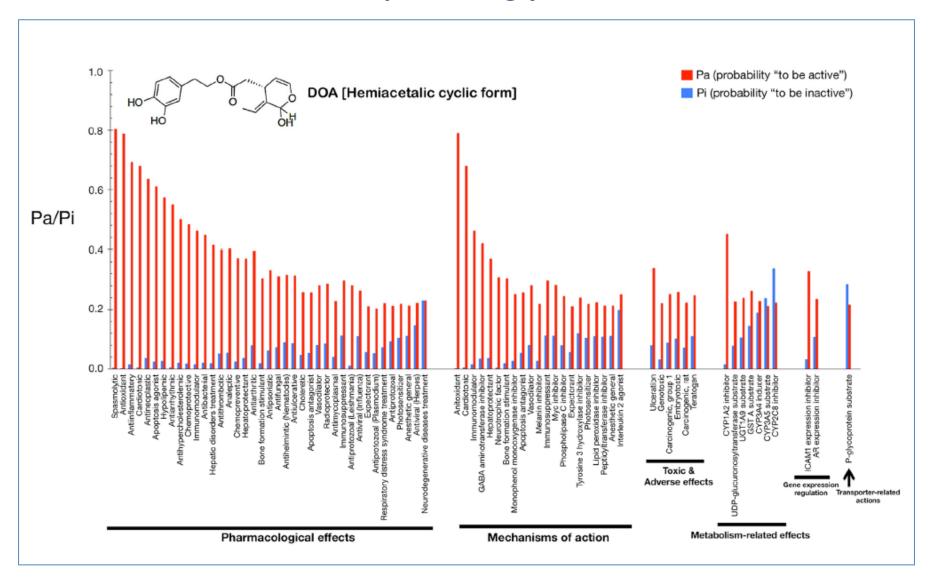
Corominas-Faja B. et al. *Aging*, 2014, 6: 731-741.

Predicted biological activity spectra for oleoprotein aglycone



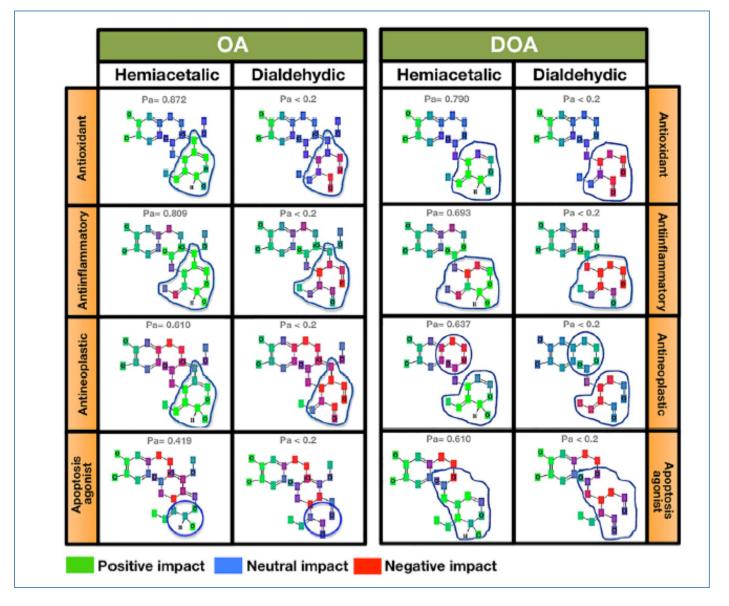
Corominas-Faja B. et al. *Aging*, 2014, 6: 731-741.

Predicted biological activity spectra for decarboxymetyl oleoprotein aglycone



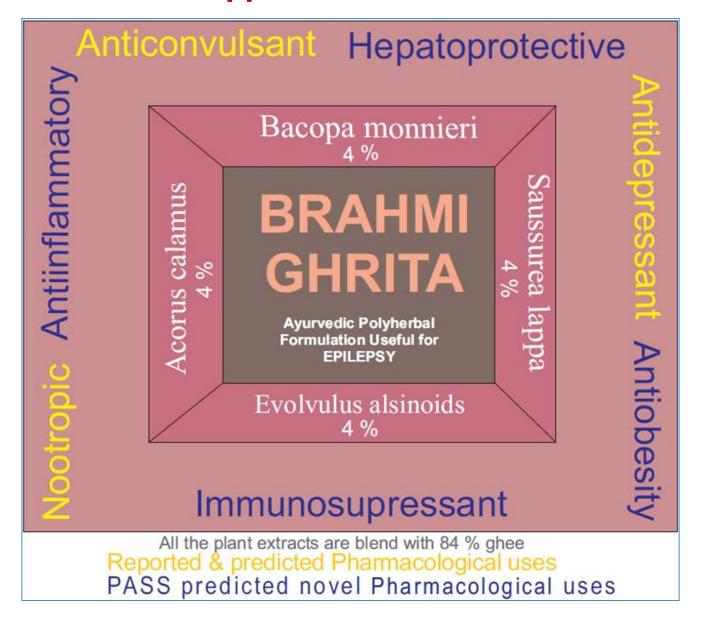
Corominas-Faja B. et al. *Aging*, 2014, 6: 731-741.

Contribution of particular atoms to anti-aging/anticancer activities of OA and DOA



Corominas-Faja B. et al. *Aging*, 2014, 6: 731-741.

Brahmi Ghrita: Bacopa monnieri, Acorus calamus, Saussurea lappa and Evolvulus alsinoids



More information could be found in our joint publications

Med Chem Res (2011) 20:1509-1514 DOI 10.1007/s00044-010-9398-x

RESEARCH

ORIGINAL RESEARCH

PASS-assisted exploration of new therapeutic potential of natural products

Rajesh Kumar Goel · Damanpreet Singh · Alexev Lagunin · Vladimir Poroikov

Received: 15 March 2010/Accepted: 22 July 2010/Published online: 6 August 2010 © Springer Science+Business Media, LLC 2010

Abstract The use of drug substances derived from plants, fungi, bacteria, and marine organisms are "Mother Nature Gift" for diseases of mankind. Many of these are discovered serendipitously and have a long tradition in medicine. Till date, the use of natural products, their semisynthetic and synthetic derivatives have been mostly confined to their ethnic use. But it has been well known that each Natural products (NPs) are used in folk medicine since substance has a wide spectrum of biological activities as evident from some new uses of many old drugs. PASS ADME/T (absorption, distribution, metabolism, and (Prediction of Activity Spectra for Substances) has been excretion/toxicity) characteristics and high chemical employed as a strong potential tool to predict the biological diversity. Presently, NPs are considered as a valuable activity spectrum of synthetic substances for the discovery of new drugs. But the potential of PASS to predict the biological activity spectra of natural products is still

Keywords Ayurveda · Biological activity spectrum Herbal drugs · Natural products · PASS

many thousands year, due to their biological origin, better source of lead structures for new pharmaceutical agents. Over 70% of New Chemical Entities (NCEs) introduced into medical practice from 1981-2006 were obtained on

Chemo- and Bioinformatics resources and in silico approaches for drug discovery from

Plants used in Traditional Indian Medicine: A Critical Review.

Lagunin A.A.1, Goel R.K.2, Gawande D.Y.2, Pahwa P.2, Gloriozova T.A.1, Dmitriev A.V.1, Ivanov S.A.¹, Rudik A.V.¹, Konova V.I.¹, Pogodin P.V.^{1,3}, Druzhilovsky D.S.¹, Poroikov V.V.^{1,3}

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³ Russian National Research Medical University, department of Biochemistry of N Biological Faculty, 117997, Ostrovitianov str. 1, Moscow, Russia.

Abstract

In silico studies are widely recognized as a useful stage of new drug discov appropriate databases. Here we review the significance of chemo- and bioinforr and in silico approaches for new drug discovery form medicinal plants used in Tr. Medicine including reverse pharmacology, QSAR, structure- and ligand-based methods. ADME/T assessment and network analysis. The review contains a prac-

combination of shows, and biginformatics mathods in the study of thoronaut

European Journal of Pharmacology 704 (2013) 33-40



Contents lists available at SciVerse ScienceDirect

European Journal of Pharmacology

journal homepage: www.elsevier.com/locate/ejphar



Behavioural pharmacology

Ameliorative effect of Curcumin on seizure severity, depression like behavior. learning and memory deficit in post-pentylenetetrazole-kindled mice

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ARTICLE INFO

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Keywords: Curcumin PTZ kindling Epileptic comorbidities Depression Learning and memory deficit Компьютерная оценка скрытого потенциала фитокомпонентов лекарственных растений из традиционной индийской медицины Аюрведа

Лагунин А.А.¹, Дружиловский Д.С.¹, Рудик А.В.¹, Филимонов Д.А.¹, Gawande D.², Suresh K2, Goel R.2, Поройков B.B.1

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Реферат

С целью изучения скрытого потенциала традиционной индийской медицины Аюрведа создан вебресурс по фитокомпонентам 50 лекарственных растений из ("http://ayurveda.pharmaexpert.ru). В реляционную централизованную специализированную базу данных введена информация о 50

REVIEW

NPR



Cite this: Nat Prod Rep., 2014, 31,

Chemo- and bioinformatics resources for in silico drug discovery from medicinal plants beyond their traditional use: a critical review†

Alexey A. Lagunin,*ac Rajesh K. Goel,*b Dinesh Y. Gawande,b Priynka Pahwa,b Tatyana A. Gloriozova, Alexander V. Dmitriev, Sergey M. Ivanov, Anastassia V. Rudik, a Varvara I. Konova, a Pavel V. Pogodin, ac Dmitry S. Druzhilovsky and Vladimir V. Poroikov*ac

ийской медицине и входящих в их ости 288 фитокомпонентов. гической активности 946 вли биологической активности и ыборку компью терной программы ппированной обучающей выборки и о одному и кросс-валидация сь, что значения средней ошибки вчений, полученных при 5%, соответственно), что иализированной версии программы вы PASS получен прогноз спектров карственных растений ТИМ. С вализ результатов прогноза для ; для ряда растений проведено экстрактов из лекарственных акотерапевтические эффекты

Covering: up to 2014

Way2Drug web platform



We have proposed the local correspondence concept, which is on the fact that hased biological activities of compounds the result are molecular recognition, which in turn depends on the correspondence between the particular atoms of the ligand and the target.

Using this concept, developed a consistent system of atom-centered neighborhoods of atoms descriptors including MNA, QNA, and LMNA. and have implemented them several SAR/QSAR/QSPR modeling approaches.

Components of Way2Drug platform (I)



Predicts about 4000 biological activity types of organic compounds by their structural formulas, including pharmacological effects, mechanisms of action, toxicity and side effects, interaction with metabolic enzymes, effects on gene expression, etc.



GUSAR online presents: consensus prediction, applicability domain assessment, internal and external models validation and clearly interpretations of obtaining results (acute rodents toxicity, antitargets, etc.).



Gene expression profiles are used to solve various problems in pharmaceutical research, such as the repositioning of drugs, overcoming resistance, estimating toxicity and drug-drug interactions.

Components of Way2Ddug platform (II)



Web-service for *in silico* prediction of cytotoxicity to the tumor and non-tumor cell-lines based on structural formula of chemical compound.



Prediction of interaction with 18 cytochrome P450 and UGT isoforms: CYP1A2, CYP2C9, CYP2C19, CYP2D6, CYP3A4, UGT1A10, UGT1A1, UGT2B7, UGT1A7, UGT2B15, UGT1A8, UGT1A4, UGT2B17, UGT2B10, UGT1A3, UGT1A9, UGT1A6, UGT2B4.



Prediction of sites of metabolism for drug-like compounds for (five major human) cytochrome P450s: CYP1A2, CYP2C9, CYP2C19, CYP2D6 and CYP3A4. Also in the training set were included the sites of glucoronidation, catalyzed by UGT.

Way2Drug services are available online



Over 300 papers published citing our web-services (>50% with the experimental confirmation; the other 50% just with the prediction results without experiments)





European Journal of Medicinal Chemistry 43 (2008) 1015-1024

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Bioorganic Medicinal Chemistry Letters

UDC 547.67

V.I. Zvarych, R.Ya. Musyanovych, V.G Chervetsov

Department of Technology of Biologically Active Substance

Lviv Polytechnic National Universi

Pharmacy and Biotechnolog

O.Z. Komarovska-Porokhnyavets, M.V. Stasevych, V.P. Noviko

Original article

Synthesis, properties, and perspectives of gem-diphosphono substituted-thiazoles

Vol. 47, No. 1

620, Russian Journal of Bioorganic Chemistry, 2013, Vol. 39, No. 2, pp. 202-210. © Pleiades Publishing, Ltd., 2013.

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A.F. Ismagilova, K.Yu. Suponitsky, D.V. Kazakov, F.E. Safarov, G.A. Tolstikov, 2013, published in Bioorganicheskaya Khimiya, 20

Synthesis, Structure, and Pharmacological Ac

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Quinazolines revisited: search for novel anxiolytic nd CADAquaia aganta

Bioorganic & Medicinal Chemistry Letters 15 (2005) 2145-2148

0066-4804/03/\$08.00+0 DOI: 10.1128/AAC.47.1.174-180.2003 Copyright © 2003, American Society for Microbiology. All Rights Reserved.

In Vitro Activities of 7-Substituted 9-Chloro and 9-Amino-2-Methoxyacridines and Their Bis- and Tetra-Acridine Complexes against Leishmania infantum

Carole Di Giorgio, 1* Florence Delmas, 1 Nathalie Filloux, 2 Maxime Robin, 2 Lactitia Seferian, 2 Nadine Azas, Monique Gasquet, Muriel Costa, Pierre Timon-David, and Jean-Pierre Galv²

Laboratoire de Parasitologie, Hygiène et Zoologie, Faculté de Pharmacie, Marseille Cedex 05,1 and Laboratoire de Parasitologie, Hygiène et Zoologie, Faculté de Pharmacie, Marseille Cedex 05,1 and Laboratoire de Parasitologie, Hygiène et Zoologie, Faculté de Pharmacie, Marseille Cedex 05,1 and Laboratoire de Parasitologie, Hygiène et Zoologie, Faculté de Pharmacie, Marseille Cedex 05,1 and Laboratoire de Pharmacie, Marseille Cedex 05,1 and Cede Valorisation de la Chimie Fine, Université d'Aix-Marseille III, Site de Saint Jérome, Marseilles,² Fra

Bioorganic & Medicinal Chemistry 20 (2012) 2930-2939

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Bioorganic & Medicinal Chemistry



SYNTHESIS OF NEW DERIVATIVES OF 2-ACYLISOTHIOCYANATE OF 1-NITRO-9,10-ANTHRAQUINONE WITH ANTIMICROBIAL ACTIVITY

УДК 378.147:547

Комбинаторная химия в высшей школе: десятилетний опыт научных, учебных и организационных проектов

journal homepage: www.elsevier.com/locate/bmc

Identification of novel isocytosine derivatives as xanthine oxidase inhibitors from a set of virtual screening hits

European Journal of Medicinal Chemistry 45 (2010) 2606-2612

Contents lists available at ScienceDirect

European Journal of Medicinal Chemist

Antimicrobial Agents and Chemotherapy, Jan. 2003, p. 174-180

Somw publications with the experimental confirmation of prediction results for natural products

Nº	Natural product	Activity	Experimental confirmation
1	Spirosolenol from roots of <i>Solanum</i> anguivi	Antiinflammatory	in vitro
2	Phytocomponents of Vitex negundo	Antioxidant, antineoplastic	in vitro
3	Phytocomponents of <i>Ficus religiosa L. (Moraceae)</i>	Anticonvulsant GABA, Aminotransferase inhibitor	in vitro
4	Quercetin	Antiinflammatory, antibacterial	in vitro
5	Polyketides from marine-derived fungus <i>Ascochyta salicorniae</i>	Protein phosphatase inhibitor	in vitro

There is dozen publications where the authors used our web-services for prediction of the biological activity spectrum of natural products with the experimental confirmation of the prediction results.

More info about the computational resources:



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REVIEW



Cite this: Nat. Prod. Rep., 2014, 31, 1585

Chemo- and bioinformatics resources for in silico drug discovery from medicinal plants beyond their traditional use: a critical review?

Alexey A. Lagunin, *ac Rajesh K. Goel, *b Dinesh Y. Gawande, b Priynka Pahwa, b Tatyana A. Gloriozova, a Alexander V. Dmitriev, a Sergey M. Ivanov, a Anastassia V. Rudik, a Varvara I. Konova, a Pavel V. Pogodin, ac Dmitry S. Druzhilovsky and Vladimir V. Poroikov*ac

REVIEWS

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Reviews • KEYNOTE REVIEW

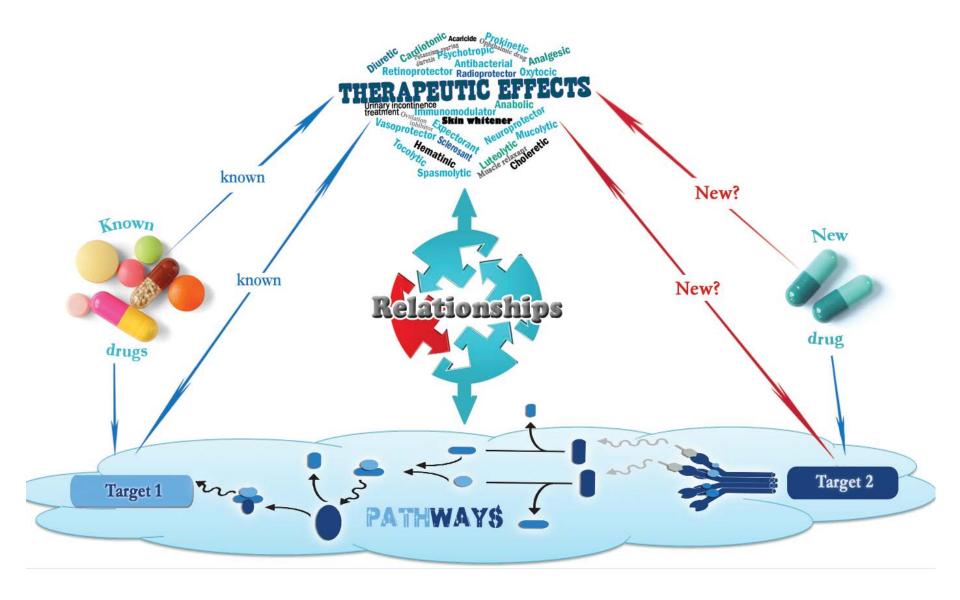
Teaser In silico approaches reveal mechanisms of adverse drug reactions and predict them at the earliest stages of drug development.



In silico assessment of adverse drug reactions and associated mechanisms

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General computer-aided approach to estimating the hidden pharmacotherapeutic potential of medicinal plants



Comparison of predictions with known effects of *Aloe vera* phytocomponents

		Prediction by PASS and PharmExpert with the cutoff Pa>0.5					
No.	Known effects	Effects	MOA	KEGG	NCI pathways	Reactome	Any approach
1	Antibacterial	+	+	+	+	+	+
2	Antifungal	+	+	+	+	+	+
3	Anti-inflammatory	+	+	+	+	+	+
4	Antimutagenic	+	-	-	-	-	+
5	Antioxidant	+	+	-	-	+	+
6	Antiprotozoal (Leishmania)	+	-	-	-	-	+
7	Antiulcerative	+	+	+	-	+	+
8	Cardioprotectant	+	-	+	+	+	+
9	Cytostatic	+	-	-	-	+	+
10	Cytotoxic	+	-	+	+	+	+
11	Hepatoprotectant	+	+	+	+	+	+
12	Hypoglycemic	-	+	+	+	+	+
13	Hypolipemic	-	+	+	-	+	+
14	Immunostimulant	-	-	+	+	+	+
	Neurodegenerative diseases						
15	treatment	-	-	+	+	+	+
16	Wound-healing agent	-	+	+	+	+	+
	True Positives (TP)	11	9	12	10	14	16
	True Negatives (TN)	67	63	54	67	49	28
	False positives (FP)	39	43	52	39	57	78
	False negatives (FN)	5	7	4	6	2	0
	Sensitivity, TP/(TP+FN)	0.69	0.56	0.75	0.63	0.88	1.00
	Specificity, TN/(TN+FP)	0.63	0.59	0.51	0.63	0.46	0.26
	Precision, TP/(TP+FP)	0.22	0.17	0.19	0.20	0.20	0.17

Summary

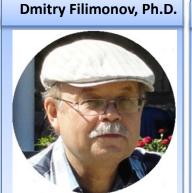
- ✓ Natural products is a valuable source for creating novel medicines because they are particularly designed by "Mother Nature" for interaction with biological systems, and due to their great chemical and pharmacological diversity.
- ✓ The hidden pharmacological potential of medicinal plants, their phytoconstituents and other natural products may be discovered using computer-aided analysis by PASS and PharmaExpert.
- ✓ Way2Drug containing many computational predictive resources may become a platform for different collaborative projects in the field of drug discovery.

Acknowledgements to the key persons and to the financial support of our long-term efforts

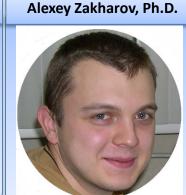


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РОССИЙСКИЙ НАУЧНЫЙ ФОНД





And to many other colleagues who participate(d) in our projects





Sixth Framework Programme 2002 - 2006

RESEARCH & INNOVATION



















Thank you for your kind attention!



We are open for collaboration.

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