

TOP 200 MEDICINES: CAN NEW ACTIONS BE DISCOVERED THROUGH COMPUTER-AIDED PREDICTION?*

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Computer-aided prediction of the biological activity spectra by the program PASS was applied to a set of 130 pharmaceuticals from the list of the Top 200 medicines. The known pharmacological effects were found in the predicted activity spectra in 93.2% of cases. Additionally, the probability of some supplementary effects was also predicted to be significant, including angiogenesis inhibition, bone formation stimulation, possible use in cognition disorders treatment, multiple sclerosis treatment, *etc.* These predictions, if confirmed experimentally, may become a cause for a new application of pharmaceuticals from the Top 200 list. Most of known side and toxic effects were also predicted by PASS. PASS predictions at earlier R & D stages may thus provide a basis for finding new “leads” among already launched drugs and may help direct more attention to those particular effects of pharmaceuticals in clinical use which become apparent only in a small part of the population and require additional precautions.

Keywords: Biological activity spectra; Top 200 medicines; Side effect; Toxicity; Computer-aided prediction; PASS

INTRODUCTION

The process of new drug research is based on the existing knowledge of the pharmacological effects that may prove useful for the treatment of certain

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disorders and the classes of molecules that are capable of producing such effects. The firm's research strategy [1] and the test standards for the drug safety assessment [2] restrict the investigation of any lead compound and drug-candidate. Besides, the information on the biological activity of each compound is always incomplete because of the concentration on only directed testing, which does not cover many possibly existing but non-tested activities. If one were able to suggest the probable activities for any compound, the directions of and priority in their testing could be established.

Computer aided structure-activity relationship (SAR) analysis is now widely used for new lead finding and optimization. Most of the SAR/QSAR/Modeling methods are only applicable to one or several types of biological activity within the same chemical series [3–5] and, therefore are unable to elucidate the general biological “potential” of a molecule under study. Recently, the possibility for simultaneous prediction of many pharmacological effects, their mechanisms of action, mutagenicity, carcinogenicity, teratogenicity and embryotoxicity has been demonstrated [6–8]. The total complex of these biological actions is called the biological activity spectrum of the compound [8]. The mean accuracy obtained with the computer program PASS [8] in leave one out cross-validation for more than 30,000 compounds with more than 500 activities from the training set was about 85% [8]. Recently, it was shown in experiments with the principal compounds from the MDDR database [9] that the approach used in PASS provided highly robust estimates for structure-activity relationships [10]. Thus, PASS ones trained can be applied to predict the activities for a new compound. In particular, PASS can be applied to already approved pharmaceuticals. Finding a new “lead” among the launched drugs is more attractive because their safety is already proven, in many cases, by previous clinical practice. On the other hand, a computer-generated “side-action” hypothesis may help focus the attention on those particular effects of pharmaceuticals in clinical use, which become apparent only in a minority of the population and require additional precautions.

The purpose of this work is to present the predictions of biological activity spectra for compounds from the list of the Top 200 pharmaceuticals [11], which are especially significant for the today's healthcare. Such computer-aided predictions may open new horizons in the study of some of the most widely used pharmaceuticals and, when compared with the experimentally established activities give an additional evaluation of the reliability for the computer program PASS.

MATERIALS AND METHODS

Data Set

The list of the Top 200 pharmaceuticals, which are the best-selling drugs in the U.S., is available from the Internet [11]. Based on their international nonproprietary and trade names, we collected the structural formulae of the substances and created the database under the ISIS/Base system (MDL Information Systems, Inc.) [9]. Information about the pharmacological main and side effects and specific toxicity of the Top 200 pharmaceuticals was collected from Refs. [11, 12]. Eighty-five percent of the pharmaceuticals had a molecular weight between 160–480 D [13] (typical drug-like compounds). The other 15% had a molecular weight between 129 and 780 D. Molecules both too small and too large were eliminated, such as: Potassium sodium chloride (K-Dur-20), Klor-Con, Potassium Chloride (ETHHEX)), Insulin (Humulin N, Humulin 70/30, Humulin R), Calcitoninum (Miacalcin Nasal), Polumyxin (Neomycin/Polymyx/HC). Mixtures of different substances or compounds with unidentified structural formulae (Premarin, Prempro) were not included in the database either. We excluded duplicates from the database which, are the same substances present in the list of Top 200 drugs under several trade names: Levothyroxine (67) – Synthroid (Knoll) and Levoxyl (Jones Pharma); Amoxicillin (10) – Amoxicillin (Teva Pharma), Trimox (Apothecon) and Amoxil (SK Beecham); Hydrocodone w/APAP (59) – Hydrocodone w/APAP (Watson, Mallincrodt, Qualitest), *etc.* The final database consisted of 130 drug substances that were used in further computer experiments.

Prediction of Biological Activity Spectra

The computer program PASS (Prediction of Activity Spectra for Substances) [6–8] predicts simultaneously several hundred biological activity types, in particular main and side pharmacological effects, mechanisms of their action, mutagenicity, carcinogenicity, teratogenicity and embryotoxicity. The prediction is based on a SAR analysis of the training set containing more than 35000 compounds, which show more than 500 kinds of biological activity. The basic elements of PASS include the representation of biological activity, the description of the chemical structures, the structure-activity relationship data and knowledge base (SAR base), and the algorithm of the activity spectra estimation.

Biological Activity

A biological activity is the result of the chemical compound's interaction with a biological entity. In a clinical study, a biological entity is represented by a human organism. In the preclinical testing, it is the experimental animals (*in vivo*) and the experimental models (*in vitro*). A biological activity depends on the peculiarities of the compound (the structure and physico-chemical properties), the biological entity (species, sex, age, *etc.*), and the mode of treatment (dose, route, *etc.*). Each biologically active compound possesses a wide spectrum of the different effects. Some of them are useful in the treatment of definite diseases but the others cause the various side and toxic effects. The total complex of activities caused by the compound in the biological entities can be called "*biological activity spectrum of a substance*".

A *biological activity spectrum* should present every activity of a compound despite of the differences in the essential conditions of their experimental determination. On the other hand, a sufficiently large set of substances can be collected only using many different sources because the information taken from a unique publication never covers all aspects of a biological action of the described substance. Thus, the "*biological activity spectrum*" may be defined as the "intrinsic" compound property depending only on its structure and physicochemical characteristics, and can be identified only qualitatively.

The biological activities are described qualitatively as the above mentioned biological activity spectra of compounds. Each compound has a list of those activities, which it possesses under some conditions.

Chemical Structure Description

In our paper published recently [14], we described the substructure descriptors called "Multilevel Neighborhoods of Atoms" (MNA). MNA descriptors are based on a structure representation that does not specify the bond types and includes hydrogen atoms according to valence and partial charge of atoms. MNA descriptors are generated as a recursively defined sequence: zero-level MNA descriptor for each atom is the mark *A* of the atom itself; any next-level MNA descriptor for each atom is the substructure notation $A(D_1D_2...D_i...)$, where D_i is the previous-level MNA descriptor for the *i*-th immediate neighbors of the atom.

This iterative process can be continued including 2nd, 3rd, *etc.*, neighborhoods of each atom. It is important to emphasize that the atom mark may include not only the atom type but also any additional information about the atom, for example, its belonging to a certain ring or

chain. A structure of a molecule is represented in PASS as a set of the 1st- and 2nd-level MNA descriptors. In 2nd-level MNA descriptors we use the mark “-” to define the belonging to a chain.

The *structure equivalence* is an important feature of the PASS approach. The structures of compounds are considered as equivalent if they have the same molecular formulae and the same set of MNA descriptors. Only unique structures are included into the SAR base. Since MNA descriptors do not represent the stereochemical peculiarities of a molecule, the compounds that differ only in stereochemistry, are formally considered as equivalent ones.

SAR Base

The prediction is carried out using the SAR base, which is created based on an analysis of the training set(s) containing the biologically active compounds. The SAR base includes the activity types and the MNA descriptor dictionaries, the substance structure and an activity description sample, and the SAR information.

To include the training set into the SAR base, the MNA descriptors are generated for each compound in this set. If a structure is not completely defined, *i.e.*, includes undetermined atoms or residues, the compound is not included in the SAR base. If an equivalent structure is found in the SAR base then new activity spectrum is combined with the existing one. The SAR data and knowledge are generated by the *training procedure* described below.

In PASS (version 1.41), as presented in this work, the SAR base contains more than 35000 substances. To create this SAR base, the training set containing drugs and well-known biologically active compounds was prepared. In the different sources the biological activities are described by different terms. Therefore the activity spectra were standardized and combined for all equivalent compounds from many sources. The MNA descriptor's dictionary contains about 36000 items. The number of different activities exceeds 800, but many of them are represented by less than 3 compounds in the training set. The total available “activity spectrum”, *i.e.*, the list of predictable types of biological activities includes more than 500 predictable biological activities. This list is modified permanently and its current version can be found on our web-site [8].

Algorithm of the Activity Spectra Estimation

The prediction algorithm was chosen from more than a hundred variants tested through the several past years. It is based on the following

SAR data:

n is the total number of compounds in the SAR base;

n_i is the number of compounds, containing the descriptor i in the structure description;

n_j is the number of compounds, containing the activity j in the activity spectrum;

n_{ij} is the number of compounds, containing both the descriptor i and the kind of activity j .

To estimate the activity spectrum the MNA descriptor set is generated for the compound under prediction. If this compound has an equivalent structure in the SAR base, that structure is excluded from the SAR base during the prediction by default. In a such case the calculations are performed by using $n-1$, n_i-1 , when the compound contains the descriptor i in the structure description, n_j-1 , when it contains the activity j in the activity spectrum, $n_{ij}-1$, when it contains both the descriptor i and the activity j . For each of the predictable activities j the initial estimation t_j is calculated as:

$$\begin{aligned} s_j &= \text{Sin}(\sum_i \text{ArcSin}(r_i * (2 * p_{ij} - 1)) / m), \\ s_{0j} &= \text{Sin}(\sum_i \text{ArcSin}(r_i * (2 * p_j - 1)) / m), \\ t_j &= (1 + (s_j - s_{0j}) / (1 - s_j * s_{0j})) / 2, \end{aligned}$$

where the summation is taken over the MNA descriptors of the compound under prediction, and m is that number, $r_i = n_i / (n_i + 0.5/m)$ is the regulating factor, $p_j = n_j / n$ is the estimation of the *a priori* probability of the activity j , and $p_{ij} = n_{ij} / n_i$ is the estimation of the conditional probability of the activity j for the MNA descriptor i .

The result of the prediction for a new compound is presented by the *activity spectrum*, which is the ranked list of the probabilities "to be active" Pa , "to be inactive" Pi , and the type of activity. The ranking is done in descending order of $Pa - Pi$; thus, more probable activities are at the top of predicted spectrum. A compound is considered active if $Pa - Pi$ exceeds the cutoff value, *e.g.*, by default $Pa - Pi < 0.0$.

The probabilities Pa and Pi are functions of the initial estimation, and defined by the equations:

$$A_j(Pa) = t_j, \quad I_j(Pi) = t_j,$$

where the functions A_j , I_j are the SAR knowledge obtained as the final result of the *training procedure*, which is described below.

Training Procedure

The SAR data n , n_i , n_j , n_{ij} are generated. For each kind of activity j , for each p of the n_j active compound, and for each q of the $n - n_j$ inactive compounds in the SAR base the leave-one-out initial estimations, *i.e.*, after exclusion this compound from the SAR base, t_{jp} and t_{jq} are calculated. The n_j estimates t_{jp} for the active compounds are sorted in ascending order; the $n - n_j$ estimates t_{jq} for the inactive compounds are sorted in descending order. The functions A_j , I_j are calculated as the conditional expectations:

$$A_j(F) = \sum_p Pr(p - 1, n_j - 1, F) t_{jp},$$

$$I_j(F) = \sum_q Pr(q - 1, n - n_j - 1, F) t_{jq},$$

where $Pr(m, n, F) = C_n^m F^m (1 - F)^{n-m}$ is the binomial distribution, $C_n^m = n! / m!(n - m)!$ is the binomial coefficient, F is in the range $[0, 1]$. It is clear that A_j and I_j are the estimations of the quantiles of the probability distributions of the initial estimation. Thus, the probabilities Pa and Pi are also the measures of belonging to subsets of "active" and "inactive" compounds, but can also be seen as the probabilities of the 1st and 2nd kinds of the prediction error, respectively. All of these possible interpretations of the probabilities Pa and Pi are equivalent and useful.

A test of PASS in blind mode by 9 scientists from 8 countries has shown that the mean accuracy of prediction is 82.6% for 108 non-congeneric compounds with 46 different types of biological activity [15]. New lead compounds with antiulcer, antitumor, hepatoprotective, antibacterial and other actions were found based on the predicted activity spectra [8].

Data Preparation

A data set of 130 drug substances, including 20 mixtures, *e.g.*, Hydrocodone w/APAP (59),¹ Amoxicillin/Clavulanate (9), *etc.* (see Tab. I), was prepared. The structure of each component of each mixture was represented separately. We exported the structural formulae of the drugs' components from ISIS/Base as SDfile [8, 9], and predicted the activity spectra for them with a cutoff condition $Pa - Pi > 0.0$. Only 2 components, Norgestimate (91), and Nitrofurantoin (87) are not found in the SAR base (they are marked in Tab. I by an asterisk *). The predictions of the activity spectra for all other substances were made with the exclusion of appropriate information from the SAR base (see above).

¹The numbers in the brackets correspond to the numbers in Table I.

TABLE I Results of activity prediction for 130 substances from the list of Top 200 drugs. The activity types are given in accordance with Ref. [11]; *Pa* is the estimation of the probability "to be active"

	<i>International Nonproprietary and Trade Names</i>	<i>Activity Types</i>	<i>Pa</i>
1	Acetaminophen/Codeine	Analgesic, general	0.960
		Analgesic-narcotic	0.957
		Analgesic-non-narcotic	0.412
		Antitussive/Expectorant	0.975
2	Albuterol; Albuterol Aerosol	Antiasthmatic	N/P
		Beta(2)-receptor agonist	0.897
		Bronchodilator	0.618
3	Alendronate; Fosamax	Calcium metabolism	0.978
4	Allopurinol	Xanthine oxidase inhibitor	0.625
5	Alprazolam	Antianxiety	0.908
		Sedative/Hypnotic	0.891
6	Amitriptyline	Antidepressant	0.970
7	Amlodipine/Benazepril; Lotrel	ACE-inhibitor	0.522
		Antianginal	0.650
		Antihypertensive	0.751
8	Amlodipine; Norvasc	Antihypertensive	0.751
		Calcium channel blocker	0.915
9	Amoxicillin/Clavulanate; Augmentin	Penicillin	0.703
10	Amoxicillin; Trimox, Amoxil	Penicillin	0.703
11	Amphetamine; Adderall	CNS, stimulant	0.838
12	Atenolol	Antianginal	0.429
		Antihypertensive	0.409
		Beta(1)-receptor blocker	0.549
13	Atorvastatin; Lipitor	Antihyperlipidemic	0.809
		HGM-CoA reductase inhibitor	0.475
14	Azithromycin; Zithromax, Zithromax Susp	Macrolide	0.865
15	Beclomethasone; Vancenase AQ DS	Antiasthmatic	N/P
		Glucocorticoid	0.785
16	Benazepril; Lotensin	ACE-inhibitor	0.522
		Antihypertensive	0.522
17	Bisoprolol/HCTZ; Ziac	Antihypertensive	0.856
		Beta(1)-receptor blocker	0.730
		Diuretic	0.974
18	Budesonide; Rhinocort	Corticosteroid	0.712
19	Bupropion; Wellbutrin SR, Zyban	Antidepressant	0.207
20	Buspirone; BuSpar	Antianxiety	0.946
21	Carisoprodol; Rela, Soma	Skeletal muscle relaxant	0.865
22	Cefprozil; Cefzi	Cephalosporin	0.633
23	Cefuroxime; Ceftin	Cephalosporin	0.711
24	Cephalexin	Cephalosporin	0.682
25	Cetirizine; Zyrtec	Antihistaminic	0.797
		H1-receptor antagonist	0.552
26	Cimetidine	Antiulcerative	0.781
		H2-receptor antagonist	0.814
27	Ciprofloxacin; Cipro	Anti-infective	0.326
		Quinolone/Derivative	0.631
28	Cisapride; Propulsid	Acid/Peptic disorder	N/P
		Gastrointestinal agent	0.685
29	Clarithromycin; Biaxin	Macrolide	0.954

TABLE I (Continued)

	<i>International Nonproprietary and Trade Names</i>	<i>Activity Types</i>	<i>Pa</i>
30	Clonazepam; Klonopin	Anticonvulsant	0.850
31	Clonidine	Antihypertensive	N/P
32	Clotrimoxazole/Betamethasone; Lotrisone	Antifungal	0.707
		Corticosteroid	0.877
33	Cyclobenzaprine	Skeletal muscle relaxant	0.502
34	Desogestrel/Ethinyl Estradiol; Desogen	Contraceptive	0.968
35	Diazepam; Valium, Valrelease	Antianxiety	0.954
		Anticonvulsant	0.884
		Skeletal muscle relaxant	0.826
36	Digoxin; Lanoxin	Antiarrhythmic	0.397
		Cardiac glycoside	0.945
37	Diltiazem; Cardizem CD	Antianginal	0.815
		Antiarrhythmic	0.490
		Antihypertensive	0.728
		Calcium channel blocker	0.927
38	Divalproex; Depakote	Anticonvulsant	0.714
39	Doxazosin; Cardura	Alpha(1)-receptor antagonist	0.734
		Antihypertensive	0.450
40	Enalapril; Vasotec	ACE-inhibitor	0.592
		Antihypertensive	0.680
41	Erythromycin; Ery-Tab	Acne product	0.538
		Macrolide	0.969
42	Estradiol	Estrogen	0.763
43	Famotidine; Pepcid	Acid/Peptic disorder	0.994
		H2-receptor antagonist	0.770
44	Felodipine; Plendil	Antihypertensive	0.769
		Calcium channel blocker	0.954
45	Fexofenadine; Allegra	Antihistaminic	0.874
		H1-receptor antagonist	0.363
46	Fluconazole; Diflucan	Antifungal	0.983
47	Fluoxetine; Prozac	Antidepressant	0.856
48	Fluticasone; Flonase	Corticosteroid – inhalation/ nasal	0.968
		Topical steroid	0.803
49	Fluvastatin; Lescol	Antihyperlipidemic	0.895
		HGM-CoA reductase inhibitor	0.732
50	Fosinopril; Monopril	ACE-inhibitor	0.944
		Antihypertensive	0.838
51	Furosemide; Furosemide Oral, Lasix	Antihypertensive	0.351
		Diuretic	0.816
52	Gabapentin; Neurontin	Anticonvulsant	0.639
53	Gemfibrozil	Antihyperlipidemic	0.572
54	Glimepiride; Amaryl	Blood glucose regulator	0.582
55	Glipizide	Blood glucose regulator	0.669
56	Glyburide; Azuglucon	Blood glucose regulator	0.677
57	Guaifenesin/Phenylpropanolamine	Antitussive/Expectorant	0.531
		Cold remedy	0.542
58	Hydrochlorothiazide; Esidrix, Dichlotride	Antihypertensive	0.856
		Diuretic	0.974
59	Hydrocodone w/APAP	Analgesic, general	0.967
		Analgesic-narcotic	0.961
		Antitussive/Expectorant	0.886

TABLE I (Continued)

	<i>International Nonproprietary and Trade Names</i>	<i>Activity Types</i>	<i>Pa</i>
60	Ibuprofen	Analgesic, general	0.558
		Analgesic-non-narcotic	0.614
		Antiarthritic	0.522
		Anti-inflammatory	0.812
61	Ipratropium; Atrovent	Antihypertensive	0.423
		Bronchodilator	N/P
62	Isosorbide Mononitrate; Imdur	Antianginal	0.986
63	Lansoprazole; Prevacid	Antiulcerative	0.978
64	Latanoprost; Xalatan	Antiglaucomic	0.683
65	Levofloxacin; Levaquin	Antibacterial	0.627
		Quinolone/Derivate	0.606
66	Levonorgestrel/Ethinyl Estradiol; Triphasil	Contraceptive	0.952
67	Levothyroxine; Synthroid, Levoxyl	Thyroid hormone	0.594
68	Lisinopril/HCTZ; Zestoretic	ACE-inhibitor	0.579
		Diuretic	0.974
69	Lisinopril; Prinivil, Zestril	ACE-inhibitor	0.575
		Antihypertensive	0.488
70	Loracarbef; Lorabid	Antibacterial, miscellaneous	0.762
		Cephalosporin	0.606
71	Loratadine/Pseudoephedrine; Claritin D 12HR, Claritin D 24HR	Antihistaminic	0.513
		Cold remedy	0.597
72	Loratadine; Claritin	Antihistaminic	0.513
73	Lorazepam; Tavor, Norlormetazepam, Temesta	Antianxiety	0.964
		Sedative/Hypnotic	0.850
74	Losartan/HCTZ; Hyzaar	Angiotensin 2 receptor antagonist	0.978
		Diuretic	0.974
75	Losartan; Cozaar	Angiotensin 2 receptor antagonist	0.992
		Antihypertensive	0.993
76	Lovastatin; Mevacor	Antihyperlipidemic	0.956
		HGM-CoA reductase inhibitor	0.936
77	Medroxyprogesterone; Cycrin, Provera	Antineoplastic	0.550
		Contraceptive	0.860
		Progestin	0.756
78	Metformin; Glucophage	Blood glucose regulator	0.807
79	Methylprednisolone	Antiarthritic	N/P
		Corticosteroid	0.519
80	Metoprolol; Toprol-XL	Antianginal	0.575
		Antihypertensive	0.534
		Beta(1)-receptor blocker	0.805
81	Mometasone; Elocon	Anti-inflammatory	0.975
		Corticosteroid	0.705
82	Mupirocin; Bactroban	Dermatologic/Miscellaneous	0.408
83	Nabumetone; Relafen	Antiarthritic	0.583
84	Naproxen; Naprosyn, Equiproxen	Analgesic, general	0.517
		Analgesic-non-narcotic	0.629
		Antiarthritic	0.409
85	Nefazodone; Serzone	Antidepressant	0.881

TABLE I (Continued)

	<i>International Nonproprietary and Trade Names</i>	<i>Activity Types</i>	<i>Pa</i>
86	Nifedipine; Procardia XL, Adalat CC	Antianginal Antihypertensive Calcium channel blocker	0.754 0.860 0.989
87	Nitrofurantoin*; Macrobid	Antiseptic, urinary tract	0.614
88	Nitroglycerin; Nitrostat	Antianginal Antihypertensive	0.995 0.857
89	Nizatidine; Axid	Antihistaminic Antiulcerative H ₂ -receptor antagonist	0.985 0.856 0.907
90	Norethindrone/Ethinyl Estradiol; Ortho-Novum	Contraceptive	0.956
91	Norgestimate*/Ethinyl Estradiol; OrthoTri-Cyclen	Contraceptive	0.896
92	Omeprazole; Prilosec	Acid/Peptic disorder	0.987
93	Oxaprozin; Daypro	Anti-inflammatory	0.811
94	Oxycodone/APAP; Roxicet	Analgesic, general Analgesic-narcotic	0.950 0.935
95	Paroxetine; Paxil	Antidepressant	0.718
96	Penicillin V Potassium; Veetids	Penicillin	0.674
97	Phenytoin; Dilantin	Anticonvulsant	0.897
98	Pravastatin; Pravachol	Antihyperlipidemic HGM-CoA reductase inhibitor	0.926 0.873
99	Prednisone	Antiarthritic Antineoplastic Corticosteroid Ocular anti-inflammatory	N/P 0.436 0.508 0.900
100	Promethazine; Phenergan	Anesthetic Antiemetic Antihistaminic Antitussive/Expectorant Sedative/Hypnotic	0.786 0.636 0.917 0.820 0.915
101	Propoxyphene N/APAP	Analgesic, general Analgesic-narcotic	0.939 0.625
102	Propranolol	Antianginal Antiarrhythmic Antimigraine Beta-receptor blocker	0.487 0.759 N/P 0.881
103	Quinapril; Accupril	ACE-inhibitor Antihypertensive	0.644 0.594
104	Ramipril; Altace	ACE-inhibitor Antihypertensive	0.802 0.619
105	Ranitidine	Antihistaminic Antiulcerative H ₂ -receptor antagonist	0.994 0.986 0.970
106	Risperidone; Risperdal	Antipsychotic/Antimanic	0.970
107	Salmeterol; Servent	Bronchodilator	0.621
108	Sertraline; Zoloft	Antidepressant Serotonin reuptake inhibitor	0.448 0.218
109	Sildenafil Citrate; Viagra	(cGMP)-specific PDE-5 inhibitor	0.839
110	Simvastatin; Zocor	Antihyperlipidemic HGM-CoA reductase inhibitor	0.982 0.975

TABLE I (Continued)

<i>International Nonproprietary and Trade Names</i>		<i>Activity Types</i>	<i>Pa</i>
111	Sumatriptan; Imitrex	5-HT ₁ receptor agonist	0.985
		Antimigraine	0.988
112	Tamoxifen	Anti-estrogen	0.953
		Antineoplastic	0.444
113	Temazepam	Sedative/Hypnotic	0.969
114	Terazosin; Hytrin	Alpha(1)-receptor antagonist	0.711
		Antihypertensive	0.461
		Diuretic	N/P
115	Tetracycline; Sumycin	Tetracycline	0.586
116	Timolol; Timoptic XE	Antiglaucomic	0.566
		Antihypertensive	0.446
		Antimigraine	N/P
		Beta-receptor blocker	0.531
117	Tobramycin/Dexamethasone; Tobradex	Aminoglycoside	0.871
118	Tramadol; Ultram	Analgesic-non-narcotic	0.877
119	Trazodone	Analgesic, general	0.545
		Antianxiety	0.720
		Antiarthritic	N/P
		Antidepressant	0.933
120	Tretinoin; Retin-A	Antineoplastic	0.638
		Dermatologic/Miscellaneous	0.954
121	Triamcinolone; Azmacort	Antiasthmatic	N/P
		Corticosteroid	0.698
		NSAID	N/P
122	Triamterene/HCTZ	Antihypertensive	0.856
		Diuretic	0.974
123	Trimeth/Sulfameth; Trimethoprim/Sulfa	Antimicrobial	0.775
124	Troglitazone; Rezulin	Antihyperglycemic	0.596
125	Valsartan; Diovan	ACE-inhibitor	N/P
		Angiotensin 2 receptor antagonist	0.370
126	Venlafaxine; Effexor	Antidepressant	0.312
127	Verapamil	Antianginal	0.700
		Antiarrhythmic	0.753
		Antiarthritic	N/P
		Antihypertensive	0.610
		Calcium channel blocker	0.559
128	Warfarin; Coumadin	Anticoagulant	0.785
129	Zafirlukast; Accolate	Antiasthmatic	N/P
		Peptide leukotriene receptor antagonist	0.457
130	Zolpidem; Ambien	Sedative/Hypnotic	0.970

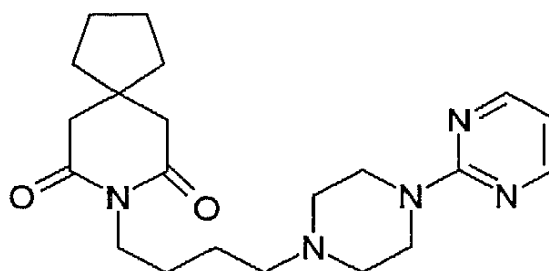
RESULTS AND DISCUSSION

Prediction of Known Activities

A comparison of the prediction results with biological activities known for each of 130 pharmaceuticals from the list of Top 200 drugs is presented in Table I. The names of activities are given in accordance with Ref. [11]. For

the majority of the compounds all known activities are predicted. There are fifteen compounds including Albuterol (2), Beclomethasone (15), Cisapride (28), *etc.*, for which some activities are not predicted. These activities are marked as "NP" in Table I. The mean accuracy of prediction, calculated for all 130 compounds and 236 activities is 93.2%.

Figure 1 presents, as an example, the predicted biological activity spectrum for Buspiron (20). All information about this compound and its



PASS 1.41 - Prediction of Activity Spectra for Substances
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Chemical Structure File: buspiron.mol
> <ACTIVITY_PREDICTION>
  34 Substructure descriptors; 0 new.
Exclude Original PASS Structure 25891 with activities:
    5 Hydroxytryptamine 1 agonist
    5 Hydroxytryptamine 1A agonist
    5 Hydroxytryptamine agonist
    Antidepressant
    Antiemetic
    Antipsychotic
    Anxiolytic
    Chemoprotective
    Hypothermic
    Psychosexual dysfunction treatment
    Psychotropic
  81 Possible activities at Pa > Pi.
  Pa   Pi   for Activity:
0.981 0.003 5 Hydroxytryptamine 1A agonist
0.969 0.006 Psychotropic
0.946 0.006 Anxiolytic
0.933 0.005 5 Hydroxytryptamine agonist
0.925 0.004 5 Hydroxytryptamine 1 agonist
0.851 0.006 Antipsychotic
0.836 0.007 Psychosexual dysfunction treatment
0.801 0.008 Antidepressant
0.709 0.007 Alpha adrenoreceptor antagonist
0.623 0.010 Neurotrophic factor
0.597 0.008 Dopamine antagonist
0.579 0.007 Dopamine D2 antagonist
0.589 0.037 Rhinitis treatment
0.571 0.020 Cognition disorders treatment
0.573 0.038 Cytokine modulator
0.566 0.044 Hypothermic
0.501 0.019 Sodium channel blocker

```

FIGURE 1 The predicted biological activity spectrum for Buspiron. Only activities with $Pa > 0.5$ are shown.

activities (5 Hydroxytryptamine 1 agonist, 5 Hydroxytryptamine 1A agonist, 5 Hydroxytryptamine agonist, Antidepressant, Antiemetic, Antipsychotic, Anxiolytic, Chemoprotective, Hypothermic, Psychosexual dysfunction treatment, Psychotropic) was excluded from the training set during the prediction. Its main activity "Tranquilizer (minor)" known from Ref. [11] corresponds to the activity "Anxiolytic" (position 3 in the prediction list, $Pa = 0.946$). New prospective activity (Neurotrophic factor) is predicted for Buspirone (20) with $Pa = 0.623$.

Prediction of New Activities in Some Drugs from Top 200 List

Figure 2 illustrates the percentage of the known activity distribution *versus* the values of the calculated probability to be active. As can be seen from Figure 2, about 89% of the predicted activities that coincided with the known activities had Pa values exceeding 50% while 63% of these activities had Pa values of more than 70%. However, in many cases additional (previously unknown) activities were predicted with rather high probability, which sometimes exceeded the probability values calculated for the known ones.

For instance, Buspirone (20) (see Fig. 1) has the highest values of probability Pa in prediction for the known activities, but there are many exceptions, the examples of which are shown in Figure 3. New prospective effects are likely to be found in some pharmaceuticals from the Top 200 list, in particular: Antiparkinsonian for Sertraline (108); Cognition disorders treatment for Ramipril (104); Multiple sclerosis treatment for Amlodipine (8) and Carisoprodol (21); Bone formation stimulant for Oxaparin (93);

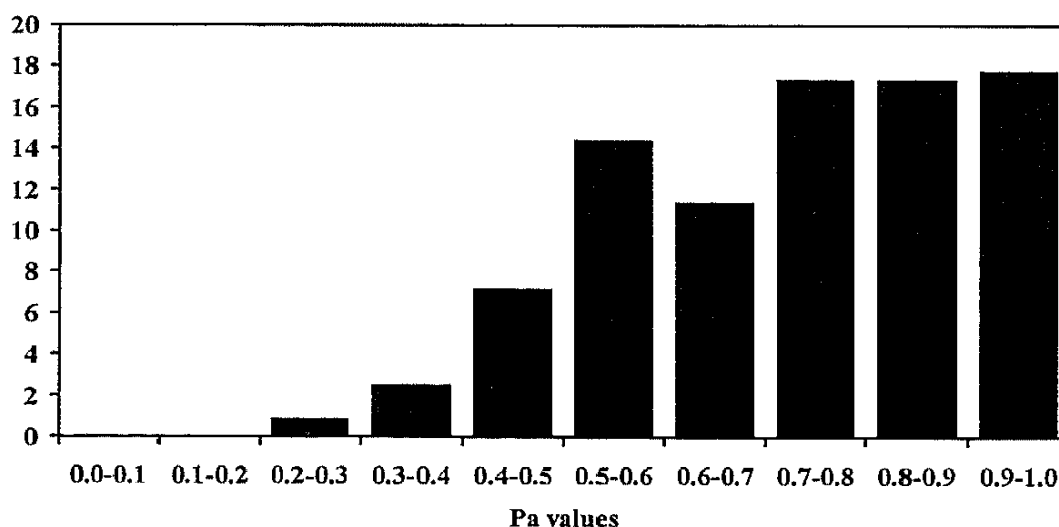


FIGURE 2 Distribution of the known activities *versus* the calculated Pa values.

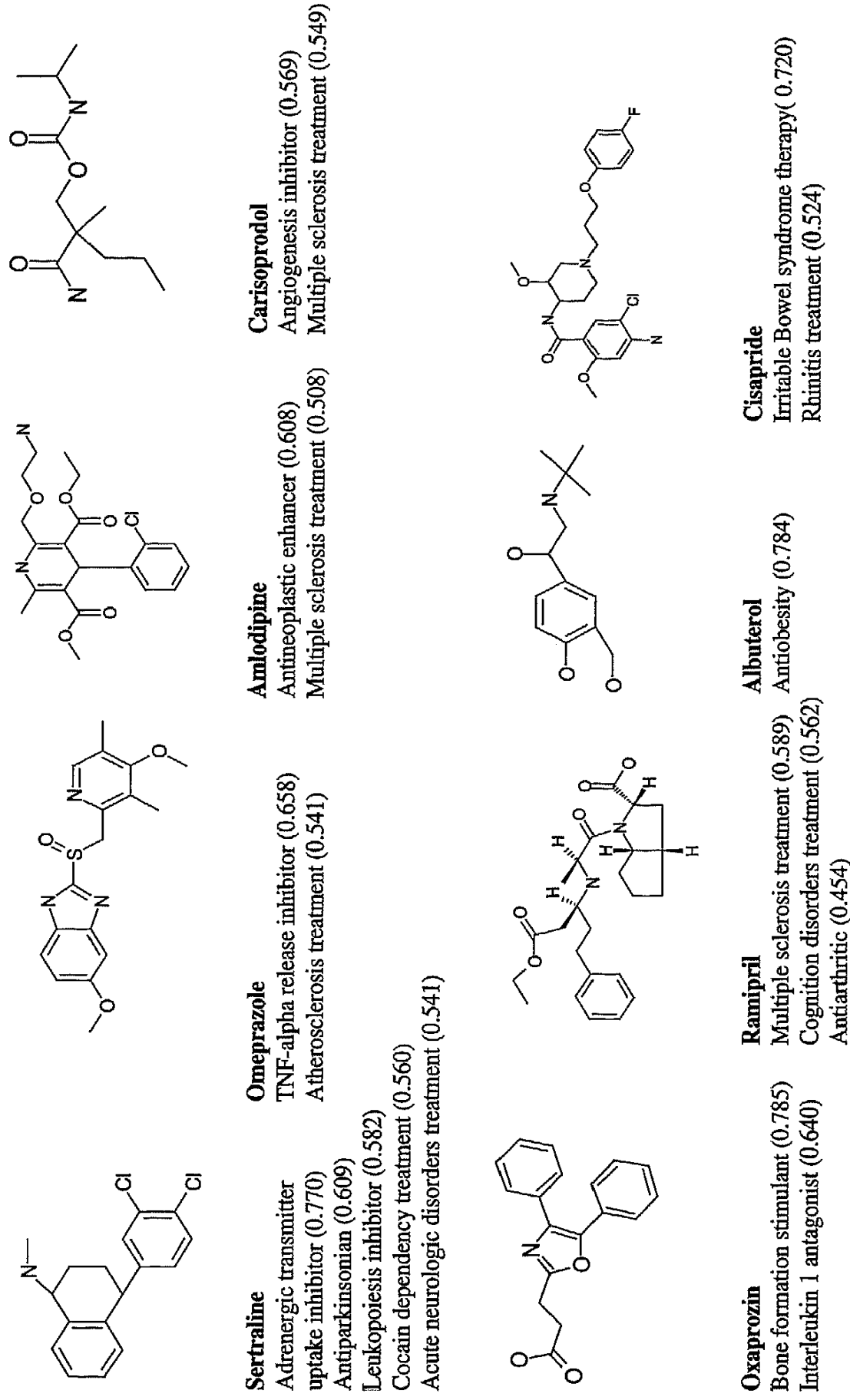


FIGURE 3 Examples of biological activities predicted de novo for some pharmaceuticals from the Top 200 list, which may become a reason for a new application. Pa values are given in brackets.

Angiogenesis inhibitor for Carisoprodol (21); *etc.* These new predicted activities have to be tested experimentally, because they may become a reason for a new application of some approved drugs.

Prediction of Side Effects in Some Drugs from Top 200 List

On the other hand, predicted biological activity spectra may include some side and toxic actions for pharmaceuticals from the Top 200 list. These data are summarized in Table II. Among the toxic effects predicted by PASS are teratogenicity and embryotoxicity, carcinogenicity; and among the side effects are abort induction, ulcerogenicity, and cardiotoxicity.

Eighty three percent of the predicted side/toxic effects coincided with the data known from literature.

Just one overprediction of teratogenicity and embryotoxicity was obtained for Prednisone (99); and three overpredictions of cardiotoxic effects were obtained for Cyclobenzaprine (33), Tramadol (118) and Amitriptyline (6). Taking into account that the equivalent structures and their biological activities were removed from the training set during the prediction, one

TABLE II Prediction of side and toxic actions for some drugs from Top 200 list

<i>Action</i>	<i>Pharmaceuticals</i>	<i>Prediction</i>	<i>Experiment</i>
Teratogen	Sertraline	+	+
	Furosemide	+	+
Embryotoxic	Prednisone	+	+
	Norgestimate/Ethyl Estradiol	+	—
	Beclomethasone	+	+
	Clotrimoxazole/Betamethasone	+	+
	Medroxyprogesterone	+	+
	Desogestrel/Ethyl Estradiol	+	+
	Triamcinolone	+	+
	Temazepam	+	+
	Estradiol	+	+
	Methylprednisolone	+	+
	Tobramycin/Dexamethasone	+	+
	Mometasone	+	+
	Budesonide	+	+
	Tetracycline	+	+
	Lorazepam	+	+
Carcinogenic	Nitrofurantoin	+	+
Abort	Propoxyphene N/APAP	+	+
Inducer	Tamoxifen	+	+
Ulcerogenic	Nabumetone	+	+
Cardiotoxic	Cyclobenzaprine	+	—
	Tramadol	+	—
	Amitriptyline	+	—

might conclude that the average prediction accuracy is reasonable to recommend PASS use for identification of such effects at the early stages of R & D.

CONCLUSIONS

- (1) The known biological actions of pharmaceuticals from the Top 200 list were predicted with the computer program PASS in 93% of the cases. 83% of predicted side and toxic effects were known for the pharmaceuticals from the Top 200 list previously. Despite the exclusion of each equivalent compound from the training set during the prediction, the mean prediction accuracy is even better than in LOO cross-validation.
- (2) High accuracy of PASS predictions provides the evidence that this program can be effectively used in search for new lead compounds with desirable activities. In particular, some new prospective biological actions were predicted for the pharmaceuticals from the Top 200 list, including possible also as: angiogenesis inhibitor, bone formation stimulant, cognition disorders treatment, multiple sclerosis treatment, *etc.*

References

- [1] Walker, S. (1994). The future of the Japanese pharmaceutical industry. – How can R & D be really international. *Pharma Japan*, **1424**, 7–10.
- [2] Maggon, K. K. and Mechkovski, A. (1992). Total quality management. *Drug News and Perspect*, **5**, 261–570.
- [3] Wermuth, C. (1996). *The practice of Medicinal Chemistry*. Academic Press.
- [4] Van de Waterbeemd, H. (1996). *Structure-property Correlations in Drug Research*. Academic Press.
- [5] Kubinyi, H., Folkers, G. and Martin, Y. C. (1997). *3D QSAR in Drug Design. Vol. II and Vol. III*. Kluwer, The Netherlands.
- [6] Filimonov, D. A. and Poroikov, V. V. (1996). PASS: Computerized prediction of biological activity spectra for chemical substances. In: *Bioactive Compound Design: Possibilities for Industrial Use*. BIOS Scientific Publishers, Oxford, pp. 47–56.
- [7] Poroikov, V. V., Filimonov, D. A., Stepanchikova, A. V. *et al.* (1996). Optimization of synthesis and pharmacological testing of new compounds based on computerized prediction of their biological activity spectra. *Chim.-Pharm. J. (Rus)*, **30**(9), 20–23.
- [8] [<http://www.ibmh.msk.su/PASS>]
- [9] [<http://www.mdli.com>]
- [10] Poroikov, V. V., Filimonov, D. A., Borodina, Yu. V., Lagunin, A. A. and Kos, A. (2000). Robustness of biological activity spectra predicting by computer program PASS for noncongeneric sets of chemical compounds. *J. Chem. Inf. Comput. Sci.*, **40**(6), 1349–1355.
- [11] [<http://www.rxlist.com>]

- [12] Drug Information for Healthcare Professionals. US Pharmacopeia, 2000.
- [13] Ghose, A. K., Viswanadhan, V. N. and Wendolovski, J. J. (1999). A knowledge-based approach in designing combinatorial or medicinal chemistry libraries for drug discovery. 1. A qualitative and quantitative characterization of known drug databases. *J. Comb. Chem.*, 1, 55–68.
- [14] Filimonov, D. A., Poroikov, V. V., Borodina, Yu. V. and Gloriovova, T. A. (1999). Chemical similarity assessment through multilevel neighborhoods of atoms: Definition and comparison with the other descriptors. *J. Chem. Inf. Comput. Sci.*, 39, 666–670.
- [15] [http://www.vei.co.uk/chemweb/library/lecture17/slideroom_babaev/transcript.html]