

# Igor V Tetko

## List of Publications by Year in descending order

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191  
papers

14,103  
citations

31976

53  
h-index

20961

115  
g-index

202  
all docs

202  
docs citations

202  
times ranked

14300  
citing authors

#	ARTICLE	IF	CITATIONS
1	Virtual Computational Chemistry Laboratory – Design and Description. <i>Journal of Computer-Aided Molecular Design</i> , 2005, 19, 453-463.	2.9	1,250
2	The FunCat, a functional annotation scheme for systematic classification of proteins from whole genomes. <i>Nucleic Acids Research</i> , 2004, 32, 5539-5545.	14.5	988
3	The <i>Fusarium graminearum</i> Genome Reveals a Link Between Localized Polymorphism and Pathogen Specialization. <i>Science</i> , 2007, 317, 1400-1402.	12.6	837
4	Neural network studies. 1. Comparison of overfitting and overtraining. <i>Journal of Chemical Information and Computer Sciences</i> , 1995, 35, 826-833.	2.8	570
5	Application of Associative Neural Networks for Prediction of Lipophilicity in ALOGPS 2.1 Program. <i>Journal of Chemical Information and Computer Sciences</i> , 2002, 42, 1136-1145.	2.8	523
6	Calculation of Molecular Lipophilicity: State-of-the-Art and Comparison of LogP Methods on more than 96,000 Compounds. <i>Journal of Pharmaceutical Sciences</i> , 2009, 98, 861-893.	3.3	517
7	Online chemical modeling environment (OCHEM): web platform for data storage, model development and publishing of chemical information. <i>Journal of Computer-Aided Molecular Design</i> , 2011, 25, 533-554.	2.9	453
8	QSAR without borders. <i>Chemical Society Reviews</i> , 2020, 49, 3525-3564.	38.1	427
9	Prediction of n-Octanol/Water Partition Coefficients from PHYSPROP Database Using Artificial Neural Networks and E-State Indices. <i>Journal of Chemical Information and Computer Sciences</i> , 2001, 41, 1407-1421.	2.8	360
10	Critical Assessment of QSAR Models of Environmental Toxicity against <i>Tetrahymena pyriformis</i> : Focusing on Applicability Domain and Overfitting by Variable Selection. <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 1733-1746.	5.4	350
11	Gene selection from microarray data for cancer classification – a machine learning approach. <i>Computational Biology and Chemistry</i> , 2005, 29, 37-46.	2.3	336
12	Estimation of Aqueous Solubility of Chemical Compounds Using E-State Indices. <i>Journal of Chemical Information and Computer Sciences</i> , 2001, 41, 1488-1493.	2.8	319
13	CERAPP: Collaborative Estrogen Receptor Activity Prediction Project. <i>Environmental Health Perspectives</i> , 2016, 124, 1023-1033.	6.0	264
14	Combinatorial QSAR Modeling of Chemical Toxicants Tested against <i>Tetrahymena pyriformis</i> . <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 766-784.	5.4	258
15	Can we estimate the accuracy of ADME-Tox predictions?. <i>Drug Discovery Today</i> , 2006, 11, 700-707.	6.4	242
16	Applicability Domains for Classification Problems: Benchmarking of Distance to Models for Ames Mutagenicity Set. <i>Journal of Chemical Information and Modeling</i> , 2010, 50, 2094-2111.	5.4	202
17	ToxAlerts: A Web Server of Structural Alerts for Toxic Chemicals and Compounds with Potential Adverse Reactions. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 2310-2316.	5.4	197
18	Computing chemistry on the web. <i>Drug Discovery Today</i> , 2005, 10, 1497-1500.	6.4	193

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19	A renaissance of neural networks in drug discovery. <i>Expert Opinion on Drug Discovery</i> , 2016, 11, 785-795.	5.0	182
20	ISIDA - Platform for Virtual Screening Based on Fragment and Pharmacophoric Descriptors. <i>Current Computer-Aided Drug Design</i> , 2008, 4, 191-198.	1.2	173
21	Application of ALOGPS to predict 1-octanol/water distribution coefficients, logP, and logD, of AstraZeneca in-house database. <i>Journal of Pharmaceutical Sciences</i> , 2004, 93, 3103-3110.	3.3	171
22	State-of-the-art augmented NLP transformer models for direct and single-step retrosynthesis. <i>Nature Communications</i> , 2020, 11, 5575.	12.8	171
23	In Silico Approaches to Prediction of Aqueous and DMSO Solubility of Drug-Like Compounds: Trends, Problems and Solutions. <i>Current Medicinal Chemistry</i> , 2006, 13, 223-241.	2.4	159
24	Parvalbumin deficiency affects network properties resulting in increased susceptibility to epileptic seizures. <i>Molecular and Cellular Neurosciences</i> , 2004, 25, 650-663.	2.2	149
25	Neural Network Studies. 2. Variable Selection. <i>Journal of Chemical Information and Computer Sciences</i> , 1996, 36, 794-803.	2.8	146
26	Neural Network Studies. 4. Introduction to Associative Neural Networks. <i>Journal of Chemical Information and Computer Sciences</i> , 2002, 42, 717-728.	2.8	146
27	Transformer-CNN: Swiss knife for QSAR modeling and interpretation. <i>Journal of Cheminformatics</i> , 2020, 12, 17.	6.1	134
28	Exhaustive QSPR Studies of a Large Diverse Set of Ionic Liquids: How Accurately Can We Predict Melting Points?. <i>Journal of Chemical Information and Modeling</i> , 2007, 47, 1111-1122.	5.4	129
29	Application of ALOGPS 2.1 to Predict log <sub>D</sub> Distribution Coefficient for Pfizer Proprietary Compounds. <i>Journal of Medicinal Chemistry</i> , 2004, 47, 5601-5604.	6.4	123
30	CoMPARA: Collaborative Modeling Project for Androgen Receptor Activity. <i>Environmental Health Perspectives</i> , 2020, 128, 27002.	6.0	120
31	Spatiotemporal activity patterns of rat cortical neurons predict responses in a conditioned task. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999, 96, 1106-1111.	7.1	112
32	Neural Network Modeling for Estimation of Partition Coefficient Based on Atom-Type Electrotopological State Indices. <i>Journal of Chemical Information and Computer Sciences</i> , 2000, 40, 947-955.	2.8	111
33	Applications of neural networks in structure-activity relationships of a small number of molecules. <i>Journal of Medicinal Chemistry</i> , 1993, 36, 811-814.	6.4	88
34	BIGCHEM: Challenges and Opportunities for Big Data Analysis in Chemistry. <i>Molecular Informatics</i> , 2016, 35, 615-621.	2.5	85
35	Applicability domain for <i>in silico</i> models to achieve accuracy of experimental measurements. <i>Journal of Chemometrics</i> , 2010, 24, 202-208.	1.3	80
36	Identification of Small-Molecule Frequent Hitters from AlphaScreen High-Throughput Screens. <i>Journal of Biomolecular Screening</i> , 2014, 19, 715-726.	2.6	77

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37	Internet Software for the Calculation of the Lipophilicity and Aqueous Solubility of Chemical Compounds. <i>Journal of Chemical Information and Computer Sciences</i> , 2001, 41, 246-252.	2.8	74
38	Associative Neural Network. <i>Neural Processing Letters</i> , 2002, 16, 187-199.	3.2	72
39	Inductive Transfer of Knowledge: Application of Multi-Task Learning and Feature Net Approaches to Model Tissue-Air Partition Coefficients. <i>Journal of Chemical Information and Modeling</i> , 2009, 49, 133-144.	5.4	71
40	How Accurately Can We Predict the Melting Points of Drug-like Compounds?. <i>Journal of Chemical Information and Modeling</i> , 2014, 54, 3320-3329.	5.4	70
41	Neural Network Studies. 3. Variable Selection in the Cascade-Correlation Learning Architecture. <i>Journal of Chemical Information and Computer Sciences</i> , 1998, 38, 651-659.	2.8	69
42	MitoP2: An Integrative Tool for the Analysis of the Mitochondrial Proteome. <i>Molecular Biotechnology</i> , 2008, 40, 306-315.	2.4	69
43	Benchmarking of Linear and Nonlinear Approaches for Quantitative Structure-Property Relationship Studies of Metal Complexation with Ionophores. <i>Journal of Chemical Information and Modeling</i> , 2006, 46, 808-819.	5.4	68
44	Optimization models for cancer classification: extracting gene interaction information from microarray expression data. <i>Bioinformatics</i> , 2004, 20, 644-652.	4.1	65
45	A Transformer Model for Retrosynthesis. <i>Lecture Notes in Computer Science</i> , 2019, , 817-830.	1.3	65
46	Calculation of lipophilicity for Pt(II) complexes: Experimental comparison of several methods. <i>Journal of Inorganic Biochemistry</i> , 2008, 102, 1424-1437.	3.5	64
47	Comparative Study of Multitask Toxicity Modeling on a Broad Chemical Space. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 1062-1072.	5.4	63
48	CATMoS: Collaborative Acute Toxicity Modeling Suite. <i>Environmental Health Perspectives</i> , 2021, 129, 47013.	6.0	63
49	A review of recent advances towards the development of QSAR models for toxicity assessment of ionic liquids. <i>Journal of Hazardous Materials</i> , 2020, 384, 121429.	12.4	61
50	The development of models to predict melting and pyrolysis point data associated with several hundred thousand compounds mined from PATENTS. <i>Journal of Cheminformatics</i> , 2016, 8, 2.	6.1	60
51	Efficient Partition of Learning Data Sets for Neural Network Training. <i>Neural Networks</i> , 1997, 10, 1361-1374.	5.9	59
52	Prediction of Acute Aquatic Toxicity toward <i>Daphnia Magna</i> by using the GA-k-NN Method. <i>ATLA Alternatives To Laboratory Animals</i> , 2014, 42, 31-41.	1.0	59
53	Prediction of partition coefficient based on atom-type electrotopological state indices. <i>Journal of Pharmaceutical Sciences</i> , 1999, 88, 229-233.	3.3	57
54	A Survey of Multi-task Learning Methods in Chemoinformatics. <i>Molecular Informatics</i> , 2019, 38, e1800108.	2.5	57

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55	Consensus Modeling for HTS Assays Using In silico Descriptors Calculates the Best Balanced Accuracy in Tox21 Challenge. <i>Frontiers in Environmental Science</i> , 2016, 4, .	3.3	55
56	A pattern grouping algorithm for analysis of spatiotemporal patterns in neuronal spike trains. 1. Detection of repeated patterns. <i>Journal of Neuroscience Methods</i> , 2001, 105, 1-14.	2.5	54
57	Corticofugal modulation of functional connectivity within the auditory thalamus of rat, guinea pig and cat revealed by cooling deactivation. <i>Journal of Neuroscience Methods</i> , 1999, 86, 161-178.	2.5	53
58	HIV-1 Reverse Transcriptase Inhibitor Design Using Artificial Neural Networks. <i>Journal of Medicinal Chemistry</i> , 1994, 37, 2520-2526.	6.4	51
59	Predicting the pKa of Small Molecules. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2011, 14, 307-327.	1.1	50
60	Volume Learning Algorithm Artificial Neural Networks for 3D QSAR Studies. <i>Journal of Medicinal Chemistry</i> , 2001, 44, 2411-2420.	6.4	49
61	Development of Dimethyl Sulfoxide Solubility Models Using 163â€™000 Molecules: Using a Domain Applicability Metric to Select More Reliable Predictions. <i>Journal of Chemical Information and Modeling</i> , 2013, 53, 1990-2000.	5.4	48
62	Different tonic regulation of neuronal activity in the rat dorsal raphe and medial prefrontal cortex via 5-HT1A receptors. <i>Neuroscience Letters</i> , 2001, 304, 129-132.	2.1	47
63	Modeling the Biodegradability of Chemical Compounds Using the Online CHEmical Modeling Environment (OCHEM). <i>Molecular Informatics</i> , 2014, 33, 73-85.	2.5	47
64	An unsupervised automatic method for sorting neuronal spike waveforms in awake and freely moving animals. <i>Methods</i> , 2003, 30, 178-187.	3.8	46
65	A comparison of different QSAR approaches to modeling CYP450 1A2 inhibition. <i>Journal of Chemical Information and Modeling</i> , 2011, 51, 1271-1280.	5.4	46
66	Extended Functional Groups (EFG): An Efficient Set for Chemical Characterization and Structure-Activity Relationship Studies of Chemical Compounds. <i>Molecules</i> , 2016, 21, 1.	3.8	46
67	Differences in locomotor behavior revealed in mice deficient for the calcium-binding proteins parvalbumin, calbindin D-28k or both. <i>Behavioural Brain Research</i> , 2007, 178, 250-261.	2.2	45
68	Prediction of logP for Pt(II) and Pt(IV) complexes: Comparison of statistical and quantum-chemistry based approaches. <i>Journal of Inorganic Biochemistry</i> , 2016, 156, 1-13.	3.5	45
69	Associative Neural Network. <i>Methods in Molecular Biology</i> , 2008, 458, 180-197.	0.9	44
70	Large, chemically diverse dataset of logP measurements for benchmarking studies. <i>European Journal of Pharmaceutical Sciences</i> , 2013, 48, 21-29.	4.0	42
71	Prediction-driven matched molecular pairs to interpret QSARs and aid the molecular optimization process. <i>Journal of Cheminformatics</i> , 2014, 6, 48.	6.1	42
72	Modelling the toxicity of a large set of metal and metal oxide nanoparticles using the OCHEM platform. <i>Food and Chemical Toxicology</i> , 2018, 112, 507-517.	3.6	42

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73	Large-scale Evaluation of log <i>P</i> Predictors: Local Corrections May Compensate Insufficient Accuracy and Need of Experimentally Testing Every Other Compound. <i>Chemistry and Biodiversity</i> , 2009, 6, 1837-1844.	2.1	40
74	Modeling of non-additive mixture properties using the Online CHEMical database and Modeling environment (OCHEM). <i>Journal of Cheminformatics</i> , 2013, 5, 4.	6.1	39
75	A pattern grouping algorithm for analysis of spatiotemporal patterns in neuronal spike trains. 2. Application to simultaneous single unit recordings. <i>Journal of Neuroscience Methods</i> , 2001, 105, 15-24.	2.5	37
76	Super paramagnetic clustering of protein sequences. <i>BMC Bioinformatics</i> , 2005, 6, 82.	2.6	37
77	Non-linear cortico-cortical interactions modulated by cholinergic afferences from the rat basal forebrain. <i>BioSystems</i> , 2000, 58, 219-228.	2.0	36
78	Chemical space exploration guided by deep neural networks. <i>RSC Advances</i> , 2019, 9, 5151-5157.	3.6	33
79	CADASTER QSPR Models for Predictions of Melting and Boiling Points of Perfluorinated Chemicals. <i>Molecular Informatics</i> , 2011, 30, 189-204.	2.5	32
80	Public (Q)SAR Services, Integrated Modeling Environments, and Model Repositories on the Web: State of the Art and Perspectives for Future Development. <i>Molecular Informatics</i> , 2017, 36, 1600082.	2.5	32
81	Trade-off Predictivity and Explainability for Machine-Learning Powered Predictive Toxicology: An in-Depth Investigation with Tox21 Data Sets. <i>Chemical Research in Toxicology</i> , 2021, 34, 541-549.	3.3	31
82	Augmentation Is What You Need!. <i>Lecture Notes in Computer Science</i> , 2019, , 831-835.	1.3	31
83	Spatiotemporal Expression Control Correlates with Intragenic Scaffold Matrix Attachment Regions (S/MARs) in <i>Arabidopsis thaliana</i> . <i>PLoS Computational Biology</i> , 2006, 2, e21.	3.2	29
84	Accurate In Silico log <i>P</i> Predictions: One Can't Embrace the Unembraceable. <i>QSAR and Combinatorial Science</i> , 2009, 28, 845-849.	1.4	29
85	The WWW as a Tool to Obtain Molecular Parameters. <i>Mini-Reviews in Medicinal Chemistry</i> , 2003, 3, 809-820.	2.4	28
86	Application of a Pruning Algorithm To Optimize Artificial Neural Networks for Pharmaceutical Fingerprinting. <i>Journal of Chemical Information and Computer Sciences</i> , 1998, 38, 660-668.	2.8	27
87	Identifying potential endocrine disruptors among industrial chemicals and their metabolites - development and evaluation of in silico tools. <i>Chemosphere</i> , 2015, 139, 372-378.	8.2	27
88	Polynomial Neural Network for Linear and Non-linear Model Selection in Quantitative-Structure Activity Relationship Studies on the Internet. <i>SAR and QSAR in Environmental Research</i> , 2000, 11, 263-280.	2.2	26
89	Electronic-Topological Investigation of the Structure - Acetylcholinesterase Inhibitor Activity Relationship in the Series of N-Benzylpiperidine Derivatives. <i>QSAR and Combinatorial Science</i> , 2001, 20, 31-45.	1.2	26
90	QSAR-derived affinity fingerprints (part 1): fingerprint construction and modeling performance for similarity searching, bioactivity classification and scaffold hopping. <i>Journal of Cheminformatics</i> , 2020, 12, 39.	6.1	26

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91	The perspectives of computational chemistry modeling. <i>Journal of Computer-Aided Molecular Design</i> , 2012, 26, 135-136.	2.9	25
92	ToxCast EPA <i>in Vitro</i> to <i>in Vivo</i> Challenge: Insight into the Rank-I Model. <i>Chemical Research in Toxicology</i> , 2016, 29, 768-775.	3.3	25
93	Fast combinatorial methods to estimate the probability of complex temporal patterns of spikes. <i>Biological Cybernetics</i> , 1997, 76, 397-408.	1.3	24
94	Dynamical cell assemblies in the rat auditory cortex in a reaction-time task. <i>BioSystems</i> , 1998, 48, 269-277.	2.0	23
95	Support vector machines for separation of mixed plant-pathogen EST collections based on codon usage. <i>Bioinformatics</i> , 2005, 21, 1383-1388.	4.1	23
96	Estimation of Acid Dissociation Constants Using Graph Kernels. <i>Molecular Informatics</i> , 2010, 29, 731-740.	2.5	23
97	PLS-Optimal: A Stepwise D-Optimal Design Based on Latent Variables. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 975-983.	5.4	23
98	GEN: highly efficient SMILES explorer using autodidactic generative examination networks. <i>Journal of Cheminformatics</i> , 2020, 12, 22.	6.1	23
99	Does "Big Data" exist in medicinal chemistry, and if so, how can it be harnessed?. <i>Future Medicinal Chemistry</i> , 2016, 8, 1801-1806.	2.3	22
100	Title is missing!. <i>Neural Processing Letters</i> , 1997, 6, 43-50.	3.2	21
101	Lattice Model for QSAR Studies. <i>Journal of Molecular Modeling</i> , 2000, 6, 517-526.	1.8	21
102	Surrogate data "a secure way to share corporate data. <i>Journal of Computer-Aided Molecular Design</i> , 2005, 19, 749-764.	2.9	21
103	Luciferase Advisor: High-Accuracy Model To Flag False Positive Hits in Luciferase HTS Assays. <i>Journal of Chemical Information and Modeling</i> , 2018, 58, 933-942.	5.4	19
104	Evaluation of CADASTER QSAR Models for the Aquatic Toxicity of (Benzo)triazoles and Prioritisation by Consensus Prediction. <i>ATLA Alternatives To Laboratory Animals</i> , 2013, 41, 49-64.	1.0	18
105	Matched Molecular Pair Analysis on Large Melting Point Datasets: A Big Data Perspective. <i>ChemMedChem</i> , 2018, 13, 599-606.	3.2	18
106	From Big Data to Artificial Intelligence: chemoinformatics meets new challenges. <i>Journal of Cheminformatics</i> , 2020, 12, 74.	6.1	17
107	Pharmaceutical Fingerprinting in Phase Space. 1. Construction of Phase Fingerprints. <i>Analytical Chemistry</i> , 1999, 71, 2423-2430.	6.5	16
108	Separation of sequences from host-pathogen interface using triplet nucleotide frequencies. <i>Fungal Genetics and Biology</i> , 2007, 44, 231-241.	2.1	16

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109	Identification of Small-Molecule Frequent Hitters of Glutathione S-Transferaseâ€“Glutathione Interaction. <i>Journal of Biomolecular Screening</i> , 2016, 21, 596-607.	2.6	16
110	The Mouse Functional Genome Database (MfunGD): functional annotation of proteins in the light of their cellular context. <i>Nucleic Acids Research</i> , 2006, 34, D568-D571.	14.5	15
111	A systematic approach to infer biological relevance and biases of gene network structures. <i>Nucleic Acids Research</i> , 2006, 34, e6-e6.	14.5	15
112	Cross-frequency coupling in mesiotemporal EEG recordings of epileptic patients. <i>Journal of Physiology (Paris)</i> , 2010, 104, 197-202.	2.1	15
113	QSAR models and scaffold-based analysis of non-nucleoside HIV RT inhibitors. <i>Chemometrics and Intelligent Laboratory Systems</i> , 2015, 148, 134-144.	3.5	15
114	Deep neural network model for highly accurate prediction of BODIPYs absorption. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2022, 267, 120577.	3.9	15
115	Deterministic Behaviour of Short Time Series. <i>Meccanica</i> , 1999, 34, 145-152.	2.0	14
116	Self-organising Modelling as a Part of Simulation Process. <i>Systems Analysis Modelling Simulation</i> , 2003, 43, 1331-1339.	0.1	14
117	MIPS bacterial genomes functional annotation benchmark dataset. <i>Bioinformatics</i> , 2005, 21, 2520-2521.	4.1	13
118	Rational design of isonicotinic acid hydrazide derivatives with antitubercular activity: Machine learning, molecular docking, synthesis and biological testing. <i>Chemical Biology and Drug Design</i> , 2018, 92, 1272-1278.	3.2	13
119	Structureâ€“Activity Relationship in Pyrazolo[4,3-c]pyridines, First Inhibitors of PEX14â€“PEX5 Proteinâ€“Protein Interaction with Trypanocidal Activity. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 847-879.	6.4	13
120	Beyond the â€˜bestâ€™ match: machine learning annotation of protein sequences by integration of different sources of information. <i>Bioinformatics</i> , 2008, 24, 621-628.	4.1	12
121	Computational Toxicology. <i>Chemical Research in Toxicology</i> , 2020, 33, 687-688.	3.3	12
122	<title>Spatiotemporal activity patterns detected from single cell measurements from behaving animals</title>. , 1999, 3728, 20.		10
123	Calculation of molecular lipophilicity: state of the art and comparison of methods on more than 96000 compounds. <i>Chemistry Central Journal</i> , 2009, 3, .	2.6	10
124	Prioritisation of Polybrominated Diphenyl Ethers (PBDEs) by Using the QSPR-THESAURUS Web Tool. <i>ATLA Alternatives To Laboratory Animals</i> , 2013, 41, 127-135.	1.0	10
125	The QSPR-THESAURUS: The Online Platform of the CADASTER Project. <i>ATLA Alternatives To Laboratory Animals</i> , 2014, 42, 13-24.	1.0	10
126	Structure-Activity Relationship Modeling and Experimental Validation of the Imidazolium and Pyridinium Based Ionic Liquids as Potential Antibacterials of MDR <i>Acinetobacter baumannii</i> and <i>Staphylococcus aureus</i> . <i>International Journal of Molecular Sciences</i> , 2021, 22, 563.	4.1	10



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127	Introduction to Special Issue: Computational Toxicology. <i>Chemical Research in Toxicology</i> , 2021, 34, 171-175.	3.3	10
128	Pharmaceutical Fingerprinting in Phase Space. 2. Pattern Recognition. <i>Analytical Chemistry</i> , 1999, 71, 2431-2439.	6.5	9
129	Volume learning algorithm significantly improved PLS model for predicting the estrogenic activity of xenoestrogens. <i>Journal of Molecular Graphics and Modelling</i> , 2007, 26, 591-594.	2.4	9
130	From Descriptors to Predicted Properties: Experimental Design by Using Applicability Domain Estimation. <i>ATLA Alternatives To Laboratory Animals</i> , 2013, 41, 33-47.	1.0	9
131	Nonlinear Dimensionality Reduction for Visualizing Toxicity Data: Distance-Based Versus Topology-Based Approaches. <i>ChemMedChem</i> , 2014, 9, 1047-1059.	3.2	9
132	An evaluation of experimental design in QSAR modelling utilizing the $k$ -medoid clustering. <i>Journal of Chemometrics</i> , 2012, 26, 509-517.	1.3	8
133	More Is Not Always Better: Local Models Provide Accurate Predictions of Spectral Properties of Porphyrins. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1201.	4.1	8
134	Theoretical and Experimental Studies of Phosphonium Ionic Liquids as Potential Antibacterials of MDR <i>Acinetobacter baumannii</i> . <i>Antibiotics</i> , 2022, 11, 491.	3.7	8
135	Focused Library Generator: case of Mdmx inhibitors. <i>Journal of Computer-Aided Molecular Design</i> , 2020, 34, 769-782.	2.9	7
136	Water envelope has a critical impact on the design of protein-protein interaction inhibitors. <i>Chemical Communications</i> , 2020, 56, 4360-4363.	4.1	7
137	Calculation of lipophilicity for Pt(II) complexes: experimental comparison of several methods. <i>Chemistry Central Journal</i> , 2008, 2, .	2.6	6
138	ROBUSTNESS IN EXPERIMENTAL DESIGN: A STUDY ON THE RELIABILITY OF SELECTION APPROACHES. <i>Computational and Structural Biotechnology Journal</i> , 2013, 7, e201305002.	4.1	6
139	Joint Virtual Special Issue on Computational Toxicology. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 1069-1071.	5.4	6
140	Simple heuristic methods for input parameters' estimation in neural networks. , 1994, , .		5
141	Evaluation of potential HIV-1 reverse transcriptase inhibitors by artificial neural networks. , 0, , .		5
142	Computer assisted neurophysiological analysis of cell assemblies activity. <i>Neurocomputing</i> , 2001, 38-40, 1025-1030.	5.9	5
143	Dopamine modulation of activity of cat sensorimotor cortex neurons during conditioned reflexes. <i>Neuroscience Letters</i> , 2002, 330, 171-174.	2.1	5
144	The effects of activation of glutamate ionotropic connections of neurons in the sensorimotor cortex in a conditioned reflex. <i>Neuroscience and Behavioral Physiology</i> , 2003, 33, 479-488.	0.4	5

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145	In silico and in vitro studies of a number PILs as new antibacterials against MDR clinical isolate <i>Acinetobacter baumannii</i> . <i>Chemical Biology and Drug Design</i> , 2020, 95, 624-630.	3.2	5
146	Using Online Tool (iPrior) for Modeling ToxCast&#8482; Assays Towards Prioritization of Animal Toxicity Testing. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2015, 18, 420-438.	1.1	5
147	Title is missing!. <i>Neural Processing Letters</i> , 1997, 6, 51-59.	3.2	4
148	A web portal for classification of expression data using maximal margin linear programming. <i>Bioinformatics</i> , 2004, 20, 3284-3285.	4.1	4
149	Dopamine modulation of glutamate metabotropic receptors in conditioned reaction of sensory motor cortex neurons of the cat. <i>Neuroscience Letters</i> , 2004, 356, 127-130.	2.1	4
150	Eclair--a web service for unravelling species origin of sequences sampled from mixed host interfaces. <i>Nucleic Acids Research</i> , 2005, 33, W724-W727.	14.5	4
151	The Prediction of Physicochemical Properties. , 0, , 240-275.		4
152	Online chemical modeling environment (OCHEM): web platform for data storage, model development and publishing of chemical information. <i>Journal of Cheminformatics</i> , 2011, 3, .	6.1	4
153	Anti-MRSA drug discovery by ligand-based virtual screening and biological evaluation. <i>Bioorganic Chemistry</i> , 2021, 114, 105042.	4.1	4
154	Variable Selection in the Cascade-Correlation Learning Architecture. , 2000, , 472-473.		4
155	Early ADME/T Predictions: Toy or Tool?. , 2008, , 240-267.		4
156	Pattern grouping algorithm and de-convolution filtering of non-stationary correlated Poisson processes. <i>Neurocomputing</i> , 2001, 38-40, 1709-1714.	5.9	3
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