

# Robert Preissner

## List of Publications by Year in descending order

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Version: 2024-02-01

135  
papers

9,654  
citations

47006

47  
h-index

42399

92  
g-index

141  
all docs

141  
docs citations

141  
times ranked

13491  
citing authors

#	ARTICLE	IF	CITATIONS
1	ProTox-II: a webservice for the prediction of toxicity of chemicals. <i>Nucleic Acids Research</i> , 2018, 46, W257-W263.	14.5	1,328
2	The target landscape of clinical kinase drugs. <i>Science</i> , 2017, 358, .	12.6	609
3	SuperTarget and Matador: resources for exploring drug-target relationships. <i>Nucleic Acids Research</i> , 2007, 36, D919-D922.	14.5	518
4	SDM—a server for predicting effects of mutations on protein stability and malfunction. <i>Nucleic Acids Research</i> , 2011, 39, W215-W222.	14.5	453
5	ProTox: a web server for the in silico prediction of rodent oral toxicity. <i>Nucleic Acids Research</i> , 2014, 42, W53-W58.	14.5	400
6	mVOC: a database of microbial volatiles. <i>Nucleic Acids Research</i> , 2014, 42, D744-D748.	14.5	337
7	SuperPred: update on drug classification and target prediction. <i>Nucleic Acids Research</i> , 2014, 42, W26-W31.	14.5	288
8	mVOC 2.0: a database of microbial volatiles. <i>Nucleic Acids Research</i> , 2018, 46, D1261-D1265.	14.5	288
9	Super Natural II—a database of natural products. <i>Nucleic Acids Research</i> , 2015, 43, D935-D939.	14.5	279
10	SuperCYP: a comprehensive database on Cytochrome P450 enzymes including a tool for analysis of CYP-drug interactions. <i>Nucleic Acids Research</i> , 2010, 38, D237-D243.	14.5	215
11	WITHDRAWN—a resource for withdrawn and discontinued drugs. <i>Nucleic Acids Research</i> , 2016, 44, D1080-D1086.	14.5	210
12	PROMISCUOUS: a database for network-based drug-repositioning. <i>Nucleic Acids Research</i> , 2011, 39, D1060-D1066.	14.5	203
13	Polymorphic Cytochrome P450 Enzymes (CYPs) and Their Role in Personalized Therapy. <i>PLoS ONE</i> , 2013, 8, e82562.	2.5	198
14	SuperTarget goes quantitative: update on drug-target interactions. <i>Nucleic Acids Research</i> , 2012, 40, D1113-D1117.	14.5	174
15	SuperPred: drug classification and target prediction. <i>Nucleic Acids Research</i> , 2008, 36, W55-W59.	14.5	144
16	<i>Legionella pneumophila</i> Induces IFN $\beta$ in Lung Epithelial Cells via IPS-1 and IRF3, Which Also Control Bacterial Replication. <i>Journal of Biological Chemistry</i> , 2006, 281, 36173-36179.	3.4	118
17	SuperScent—a database of flavors and scents. <i>Nucleic Acids Research</i> , 2009, 37, D291-D294.	14.5	106
18	Evidence for treatment with estradiol for women with SARS-CoV-2 infection. <i>BMC Medicine</i> , 2020, 18, 369.	5.5	106

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19	Prediction Is a Balancing Act: Importance of Sampling Methods to Balance Sensitivity and Specificity of Predictive Models Based on Imbalanced Chemical Data Sets. <i>Frontiers in Chemistry</i> , 2018, 6, 362.	3.6	97
20	A Comprehensive View on Proteasomal Sequences: Implications for the Evolution of the Proteasome. <i>Journal of Molecular Biology</i> , 2003, 326, 1437-1448.	4.2	96
21	TRPC6 G757D Loss-of-Function Mutation Associates with FSGS. <i>Journal of the American Society of Nephrology: JASN</i> , 2016, 27, 2771-2783.	6.1	94
22	SuperLooper--a prediction server for the modeling of loops in globular and membrane proteins. <i>Nucleic Acids Research</i> , 2009, 37, W571-W574.	14.5	85
23	SuperNatural: a searchable database of available natural compounds. <i>Nucleic Acids Research</i> , 2006, 34, D678-D683.	14.5	84
24	SuperSweet--a resource on natural and artificial sweetening agents. <i>Nucleic Acids Research</i> , 2011, 39, D377-D382.	14.5	84
25	Chemical Proteomics Reveals Ferrochelatase as a Common Off-target of Kinase Inhibitors. <i>ACS Chemical Biology</i> , 2016, 11, 1245-1254.	3.4	82
26	Evidence-Based Considerations Exploring Relations between SARS-CoV-2 Pandemic and Air Pollution: Involvement of PM2.5-Mediated Up-Regulation of the Viral Receptor ACE-2. <i>International Journal of Environmental Research and Public Health</i> , 2020, 17, 5573.	2.6	82
27	SuperDRUG2: a one stop resource for approved/ marketed drugs. <i>Nucleic Acids Research</i> , 2018, 46, D1137-D1143.	14.5	81
28	SuperDrug: a conformational drug database. <i>Bioinformatics</i> , 2005, 21, 1751-1753.	4.1	80
29	IL-13 is a driver of COVID-19 severity. <i>JCI Insight</i> , 2021, 6, .	5.0	80
30	Voronoi: analyzing packing in protein structures. <i>Nucleic Acids Research</i> , 2009, 37, D393-D395.	14.5	78
31	CancerResource: a comprehensive database of cancer-relevant proteins and compound interactions supported by experimental knowledge. <i>Nucleic Acids Research</i> , 2011, 39, D960-D967.	14.5	70
32	Statin and rottlerin small-molecule inhibitors restrict colon cancer progression and metastasis via MACC1. <i>PLoS Biology</i> , 2017, 15, e2000784.	5.6	70
33	SuperToxic: a comprehensive database of toxic compounds. <i>Nucleic Acids Research</i> , 2009, 37, D295-D299.	14.5	69
34	Structural features of transmembrane helices. <i>FEBS Letters</i> , 2004, 559, 145-151.	2.8	63
35	Characterization of GD2 Peptide Mimotope DNA Vaccines Effective against Spontaneous Neuroblastoma Metastases. <i>Cancer Research</i> , 2006, 66, 10567-10575.	0.9	63
36	Survivin minigene DNA vaccination is effective against neuroblastoma. <i>International Journal of Cancer</i> , 2009, 125, 104-114.	5.1	63

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37	Natural products: sources and databases. <i>Natural Product Reports</i> , 2006, 23, 347-356.	10.3	62
38	Analysis of sex and gender-specific research reveals a common increase in publications and marked differences between disciplines. <i>BMC Medicine</i> , 2010, 8, 70.	5.5	60
39	Computational methods for prediction of in vitro effects of new chemical structures. <i>Journal of Cheminformatics</i> , 2016, 8, 51.	6.1	58
40	The Transformer database: biotransformation of xenobiotics. <i>Nucleic Acids Research</i> , 2014, 42, D1113-D1117.	14.5	57
41	The RTP Site Shared by the HIV-1 Tat Protein and the 11S Regulator Subunit $\hat{\pm}$ is Crucial for their Effects on Proteasome Function Including Antigen Processing. <i>Journal of Molecular Biology</i> , 2002, 323, 771-782.	4.2	56
42	SynSysNet: integration of experimental data on synaptic protein-protein interactions with drug-target relations. <i>Nucleic Acids Research</i> , 2012, 41, D834-D840.	14.5	54
43	Hydrogen-Bonding and Packing Features of Membrane Proteins: Functional Implications. <i>Biophysical Journal</i> , 2008, 94, 1945-1953.	0.5	52
44	Novel curcumin- and emodin-related compounds identified by in silico 2D/3D conformer screening induce apoptosis in tumor cells. <i>BMC Cancer</i> , 2005, 5, 97.	2.6	51
45	Photoactivation of an Inhibitor of the 12/15-lipoxygenase Pathway. <i>ChemBioChem</i> , 2006, 7, 1089-1095.	2.6	50
46	Docking without docking: ISEARCH prediction of interactions using known interfaces. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007, 69, 839-844.	2.6	50
47	BitterSweetForest: A Random Forest Based Binary Classifier to Predict Bitterness and Sweetness of Chemical Compounds. <i>Frontiers in Chemistry</i> , 2018, 6, 93.	3.6	50
48	Comparison of 2D Similarity and 3D Superposition. Application to Searching a Conformational Drug Database. <i>Journal of Chemical Information and Computer Sciences</i> , 2004, 44, 1816-1822.	2.8	47
49	SuperCYPsPred a web server for the prediction of cytochrome activity. <i>Nucleic Acids Research</i> , 2020, 48, W580-W585.	14.5	47
50	The Exponential Phase of the Covid-19 Pandemic in Central Italy: An Integrated Care Pathway. <i>International Journal of Environmental Research and Public Health</i> , 2020, 17, 3792.	2.6	46
51	SuperPred 3.0: drug classification and target prediction a machine learning approach. <i>Nucleic Acids Research</i> , 2022, 50, W726-W731.	14.5	46
52	Prediction of prolyl residues in cis-conformation in protein structures on the basis of the amino acid sequence. <i>FEBS Letters</i> , 1990, 277, 159-163.	2.8	45
53	Conservation of cis prolyl bonds in proteins during evolution. <i>Proteins: Structure, Function and Bioinformatics</i> , 2004, 58, 589-595.	2.6	43
54	Molecular similarity-based predictions of the Tox21 screening outcome. <i>Frontiers in Environmental Science</i> , 2015, 3, .	3.3	43

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55	Columba: an integrated database of proteins, structures, and annotations. BMC Bioinformatics, 2005, 6, 81.	2.6	40
56	Suppressive Effects of Clerodendrum volubile P Beauv. [Labiatae] Methanolic Extract and Its Fractions on Type 2 Diabetes and Its Complications. Frontiers in Pharmacology, 2018, 9, 8.	3.5	40
57	Molecular Packing and Packing Defects in Helical Membrane Proteins. Biophysical Journal, 2005, 88, 1970-1977.	0.5	39
58	The Catch-22 of Predicting hERG Blockade Using Publicly Accessible Bioactivity Data. Journal of Chemical Information and Modeling, 2018, 58, 1224-1233.	5.4	39
59	SuperLigands - a database of ligand structures derived from the Protein Data Bank. BMC Bioinformatics, 2005, 6, 122.	2.6	36
60	Anti-diabetic effect of the ethyl acetate fraction of Clerodendrum volubile: protocatechuic acid suppresses phagocytic oxidative burst and modulates inflammatory cytokines. Biomedicine and Pharmacotherapy, 2017, 86, 307-315.	5.6	36
61	Inhomogeneous molecular density: reference packing densities and distribution of cavities within proteins. Bioinformatics, 2003, 19, 2112-2121.	4.1	35
62	A rationally designed tyrosine hydroxylase DNA vaccine induces specific antineuroblastoma immunity. Molecular Cancer Therapeutics, 2008, 7, 2241-2251.	4.1	35
63	Sequence information within proteasomal prosequences mediates efficient integration of $\hat{1}^2$ -subunits into the 20 s proteasome complex. Journal of Molecular Biology, 1999, 288, 117-128.	4.2	34
64	CancerResourceâ€”updated database of cancer-relevant proteins, mutations and interacting drugs. Nucleic Acids Research, 2016, 44, D932-D937.	14.5	33
65	A New Algorithm for Integrated Analysis of miRNA-mRNA Interactions Based on Individual Classification Reveals Insights into Bladder Cancer. PLoS ONE, 2013, 8, e64543.	2.5	33
66	Superimpose: a 3D structural superposition server. Nucleic Acids Research, 2008, 36, W47-W54.	14.5	31
67	Representation of target-bound drugs by computed conformers: implications for conformational libraries. BMC Bioinformatics, 2006, 7, 293.	2.6	30
68	SuperSite: dictionary of metabolite and drug binding sites in proteins. Nucleic Acids Research, 2009, 37, D195-D200.	14.5	30
69	Analysis and prediction of helixâ€”helix interactions in membrane channels and transporters. Proteins: Structure, Function and Bioinformatics, 2006, 64, 253-262.	2.6	28
70	SuperMimic--fitting peptide mimetics into protein structures. BMC Bioinformatics, 2006, 7, 11.	2.6	28
71	Oestrogen-mediated upregulation of the Mas receptor contributes to sex differences in acute lung injury and lung vascular barrier regulation. European Respiratory Journal, 2021, 57, 2000921.	6.7	28
72	SL2: an interactive webtool for modeling of missing segments in proteins. Nucleic Acids Research, 2016, 44, W390-W394.	14.5	26

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73	Enzymatic, expression and structural divergences among carboxyl O-methyltransferases after gene duplication and speciation in <i>Nicotiana</i> . <i>Plant Molecular Biology</i> , 2010, 72, 311-330.	3.9	25
74	Computational Prediction of Potential Inhibitors of the Main Protease of SARS-CoV-2. <i>Frontiers in Chemistry</i> , 2020, 8, 590263.	3.6	24
75	Reynoutria Rhizomes as a Natural Source of SARS-CoV-2 Mpro Inhibitors—Molecular Docking and In Vitro Study. <i>Pharmaceuticals</i> , 2021, 14, 742.	3.8	24
76	Volatile Organic Compounds in the Breath of Oral Squamous Cell Carcinoma Patients: A Pilot Study. <i>Otolaryngology - Head and Neck Surgery</i> , 2017, 157, 981-987.	1.9	23
77	Computational prediction of immune cell cytotoxicity. <i>Food and Chemical Toxicology</i> , 2017, 107, 150-166.	3.6	23
78	Identification of signature volatiles to discriminate <i>Candida albicans</i> , <i>glabrata</i> , <i>krusei</i> and <i>tropicalis</i> using gas chromatography and mass spectrometry. <i>Mycoses</i> , 2016, 59, 117-126.	4.0	22
79	Adjuvant antifungal therapy using tissue tolerable plasma on oral mucosa and removable dentures in oral candidiasis patients: a randomised double-blind split-mouth pilot study. <i>Mycoses</i> , 2016, 59, 467-475.	4.0	21
80	PROMISCUOUS 2.0: a resource for drug-repositioning. <i>Nucleic Acids Research</i> , 2021, 49, D1373-D1380.	14.5	21
81	VirtualTaste: a web server for the prediction of organoleptic properties of chemical compounds. <i>Nucleic Acids Research</i> , 2021, 49, W679-W684.	14.5	21
82	SuperHapten: a comprehensive database for small immunogenic compounds. <i>Nucleic Acids Research</i> , 2007, 35, D906-D910.	14.5	19
83	FragmentStore—a comprehensive database of fragments linking metabolites, toxic molecules and drugs. <i>Nucleic Acids Research</i> , 2011, 39, D1049-D1054.	14.5	19
84	GenderMedDB: an interactive database of sex and gender-specific medical literature. <i>Biology of Sex Differences</i> , 2014, 5, 7.	4.1	19
85	Computational and experimental validation of antioxidant properties of synthesized bioactive ferulic acid derivatives. <i>International Journal of Food Properties</i> , 2018, 21, 86-98.	3.0	19
86	Allosteric Modulation of the CB1 Cannabinoid Receptor by Cannabidiol—A Molecular Modeling Study of the N-Terminal Domain and the Allosteric-Orthosteric Coupling. <i>Molecules</i> , 2021, 26, 2456.	3.8	19
87	Sixty-One Volatiles Have Phylogenetic Signals Across Bacterial Domain and Fungal Kingdom. <i>Frontiers in Microbiology</i> , 2020, 11, 557253.	3.5	17
88	Oxidative testicular injury: effect of l-leucine on redox, cholinergic and purinergic dysfunctions, and dysregulated metabolic pathways. <i>Amino Acids</i> , 2021, 53, 359-380.	2.7	17
89	Real-world evidence for improved outcomes with histamine antagonists and aspirin in 22,560 COVID-19 patients. <i>Signal Transduction and Targeted Therapy</i> , 2021, 6, 267.	17.1	17
90	SUPERFICIAL—surface mapping of proteins via structure-based peptide library design. <i>BMC Bioinformatics</i> , 2005, 6, 223.	2.6	15

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91	Theoretical Modeling Techniques and Their Impact on Tumor Immunology. <i>Clinical and Developmental Immunology</i> , 2010, 2010, 1-11.	3.3	15
92	Outcome of revascularization therapy in traumatized immature incisors. <i>BMC Oral Health</i> , 2020, 20, 207.	2.3	15
93	Onset of Oral Lichenoid Lesions and Oral Lichen Planus Following COVID-19 Vaccination: A Retrospective Analysis of about 300,000 Vaccinated Patients. <i>Vaccines</i> , 2022, 10, 480.	4.4	15
94	Accelerating screening of 3D protein data with a graph theoretical approach. <i>Bioinformatics</i> , 2003, 19, 2442-2447.	4.1	14
95	Sensitive to Infection but Strong in Defense—Female Sex and the Power of Oestradiol in the COVID-19 Pandemic. <i>Frontiers in Global Women S Health</i> , 2021, 2, 651752.	2.3	14
96	Inverse sequence similarity of proteins does not imply structural similarity. <i>FEBS Letters</i> , 2003, 545, 105-109.	2.8	13
97	Exploring Activity Profiles of PAINS and Their Structural Context in Target–Ligand Complexes. <i>Journal of Chemical Information and Modeling</i> , 2018, 58, 1847-1857.	5.4	13
98	Norditerpenoids with Selective Anti-Cholinesterase Activity from the Roots of <i>Perovskia atriplicifolia</i> Benth.. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4475.	4.1	13
99	Vacclmm: simulating peptide vaccination in cancer therapy. <i>BMC Bioinformatics</i> , 2013, 14, 127.	2.6	12
100	Development of Immune-Specific Interaction Potentials and Their Application in the Multi-Agent-System Vacclmm. <i>PLoS ONE</i> , 2011, 6, e23257.	2.5	12
101	DNA Vaccination: Using the Patient's Immune System to Overcome Cancer. <i>Clinical and Developmental Immunology</i> , 2010, 2010, 1-14.	3.3	11
102	Acyclovir Has Low but Detectable Influence on HLA-B*57:01 Specificity without Inducing Hypersensitivity. <i>PLoS ONE</i> , 2015, 10, e0124878.	2.5	11
103	Drugs as habitable planets in the space of dark chemical matter. <i>Drug Discovery Today</i> , 2018, 23, 481-486.	6.4	11
104	Cytotoxic Effect of Vanicosides A and B from <i>Reynoutria sachalinensis</i> against Melanotic and Amelanotic Melanoma Cell Lines and in silico Evaluation for Inhibition of BRAFV600E and MEK1. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4611.	4.1	11
105	Cellular Fingerprints: A Novel Approach Using Large-Scale Cancer Cell Line Data for the Identification of Potential Anticancer Agents. <i>Chemical Biology and Drug Design</i> , 2009, 74, 439-448.	3.2	10
106	Inhibiting the inhibitors: Retro-inverso Smac peptides. <i>Peptides</i> , 2009, 30, 2374-2379.	2.4	10
107	Inhibition of DNA–Topoisomerase I by Acylated Triterpene Saponins from <i>Pittosporum angustifolium</i> Lodd.. <i>Natural Products and Bioprospecting</i> , 2016, 6, 141-147.	4.3	10
108	Personalized Cancer Therapy Considering Cytochrome P450 Variability. <i>Advances in Pharmacology</i> , 2015, 74, 113-130.	2.0	9

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109	2D and 3D similarity landscape analysis identifies PARP as a novel off-target for the drug Vatalanib. <i>BMC Bioinformatics</i> , 2015, 16, 308.	2.6	9
110	A Comparative Analysis of COVID-19 Vaccines Based on over 580,000 Cases from the Vaccination Adverse Event Reporting System. <i>Vaccines</i> , 2022, 10, 408.	4.4	9
111	Cobweb: a Java applet for network exploration and visualisation. <i>Bioinformatics</i> , 2011, 27, 1725-1726.	4.1	8
112	L-leucine stimulation of glucose uptake and utilization involves modulation of glucose $\alpha$ lipid metabolic switch and improved bioenergetic homeostasis in isolated rat psoas muscle ex vivo. <i>Amino Acids</i> , 2021, 53, 1135-1151.	2.7	8
113	A minigene DNA vaccine encoding peptide epitopes derived from Galectin-1 has protective antitumoral effects in a model of neuroblastoma. <i>Cancer Letters</i> , 2021, 509, 105-114.	7.2	8
114	Pain-Prescription Differences - An Analysis of 500,000 Discharge Summaries. <i>Current Drug Research Reviews</i> , 2019, 11, 58-66.	1.4	8
115	Prediction of oral squamous cell carcinoma based on machine learning of breath samples: a prospective controlled study. <i>BMC Oral Health</i> , 2021, 21, 500.	2.3	8
116	Comparison of five-year survival rates among patients with oral squamous cell carcinoma with and without association with syphilis: a retrospective case-control study. <i>BMC Cancer</i> , 2022, 22, 454.	2.6	8
117	JAIL: a structure-based interface library for macromolecules. <i>Nucleic Acids Research</i> , 2009, 37, D338-D341.	14.5	7
118	Binding sites in membrane proteins $\alpha$ Diversity, druggability and prospects. <i>European Journal of Cell Biology</i> , 2012, 91, 326-339.	3.6	7
119	Voroia 4-ever. <i>Nucleic Acids Research</i> , 2021, 49, W685-W690.	14.5	7
120	Non-Psychotropic Cannabinoids as Inhibitors of TET1 Protein. <i>Bioorganic Chemistry</i> , 2022, 124, 105793.	4.1	7
121	Residues of a proposed gate region in type I ATP-binding cassette import systems are crucial for function as revealed by mutational analysis. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2013, 1828, 2164-2172.	2.6	6
122	CardioScape mapping the cardiovascular funding landscape in Europe. <i>European Heart Journal</i> , 2018, 39, 2423-2430.	2.2	6
123	Evaluation of Drug-Drug Interactions in EGFR-Mutated Non-Small-Cell Lung Cancer Patients during Treatment with Tyrosine-Kinase Inhibitors. <i>Journal of Personalized Medicine</i> , 2021, 11, 424.	2.5	6
124	The druggability of bitter taste receptors for the treatment of neurodegenerative disorders. <i>Biochemical Pharmacology</i> , 2022, 197, 114915.	4.4	6
125	A Birds-Eye (Re)View of Acid-Suppression Drugs, COVID-19, and the Highly Variable Literature. <i>Frontiers in Pharmacology</i> , 2021, 12, 700703.	3.5	5
126	In silico screening of drug databases for TSE inhibitors. <i>BioSystems</i> , 2005, 80, 117-122.	2.0	4



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127	Mapping discontinuous proteinâ€binding sites via structureâ€based peptide libraries: combining <i>in silico</i> and <i>in vitro</i> approaches. <i>Journal of Molecular Recognition</i> , 2013, 26, 23-31.	2.1	4
128	Volatile organic compounds in the breath of oral candidiasis patients: a pilot study. <i>Clinical Oral Investigations</i> , 2018, 22, 721-731.	3.0	4
129	Individualized Drugsâ€™ Selection by Evaluation of Drug Properties, Pharmacogenomics and Clinical Parameters: Performance of a Bioinformatic Tool Compared to a Clinically Established Counselling Process. <i>Pharmacogenomics and Personalized Medicine</i> , 2021, Volume 14, 955-962.	0.7	3
130	Man vs. machine: comparison of pharmacogenetic expert counselling with a clinical medication support system in a study with 200 genotyped patients. <i>European Journal of Clinical Pharmacology</i> , 2022, 78, 579-587.	1.9	2
131	Computer-assisted identification of small-molecule Bcl-2 modulators. <i>Computational Biology and Chemistry</i> , 2009, 33, 451-456.	2.3	1
132	Molecular basis for the sensitivity of TRP channels to polyunsaturated fatty acids. <i>Naunyn-Schmiedeberg's Archives of Pharmacology</i> , 2018, 391, 833-846.	3.0	1
133	Comparison of Computerized Prescription Support Systems in COVID-19 Patients: INTERCheck and Drug-PIN. <i>SN Comprehensive Clinical Medicine</i> , 2021, 4, 3.	0.6	1
134	DBEndo: a web-based endodontic case management tool. <i>BMC Research Notes</i> , 2015, 8, 685.	1.4	0
135	CardioScape-II: the need to map cardiovascular funding patterns in Europe. <i>Cardiovascular Research</i> , 2020, 116, 879-881.	3.8	0