

Wei Zheng

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

280
papers

10,553
citations

52
h-index

92
g-index

295
ext. papers

13,352
ext. citations

7
avg. IF

6.49
L-index

#	Paper	IF	Citations
280	SARS-CoV-2 Nucleocapsid Protein TR-FRET Assay Amenable to High Throughput Screening.. <i>ACS Pharmacology and Translational Science</i> , 2022 , 5, 8-19	5.9	0
279	Targeting the Fusion Process of SARS-CoV-2 Infection by Small Molecule Inhibitors.. <i>MBio</i> , 2022 , e03238218	7.8	1
278	A high throughput screening assay for inhibitors of SARS-CoV-2 pseudotyped particle entry.. <i>SLAS Discovery</i> , 2022 ,	3.4	4
277	Domain knowledge-based security bug reports prediction. <i>Knowledge-Based Systems</i> , 2022 , 241, 108293	7.3	7
276	iPS-derived neural stem cells for disease modeling and evaluation of therapeutics for mucopolysaccharidosis type II.. <i>Experimental Cell Research</i> , 2022 , 412, 113007	4.2	0
275	Continuous Encoding for Overlapping Community Detection in Attributed Network.. <i>IEEE Transactions on Cybernetics</i> , 2022 , PP,	10.2	1
274	c-Abl Activation Linked to Autophagy-Lysosomal Dysfunction Contributes to Neurological Impairment in Niemann-Pick Type A Disease.. <i>Frontiers in Cell and Developmental Biology</i> , 2022 , 10, 844297	5.7	0
273	Mitoxantrone modulates a heparan sulfate-spike complex to inhibit SARS-CoV-2 infection.. <i>Scientific Reports</i> , 2022 , 12, 6294	4.9	0
272	Progressive assembly of multi-domain protein structures from cryo-EM density maps. <i>Nature Computational Science</i> , 2022 , 2, 265-275		0
271	Decoding the link of microbiome niches with homologous sequences enables accurately targeted protein structure prediction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	2
270	A Comparative Study of Class Rebalancing Methods for Security Bug Report Classification. <i>IEEE Transactions on Reliability</i> , 2021 , 70, 1658-1670	4.6	28
269	Mining of high throughput screening database reveals AP-1 and autophagy pathways as potential targets for COVID-19 therapeutics. <i>Scientific Reports</i> , 2021 , 11, 6725	4.9	12
268	Induction of interferon signaling and allograft inflammatory factor 1 in macrophages in a mouse model of breast cancer metastases. <i>Wellcome Open Research</i> , 2021 , 6, 52	4.8	2
267	Identification of Antifungal Compounds against Multidrug-Resistant <i>Candida auris</i> Utilizing a High-Throughput Drug-Repurposing Screen. <i>Antimicrobial Agents and Chemotherapy</i> , 2021 , 65,	5.9	4
266	Deducing high-accuracy protein contact-maps from a triplet of coevolutionary matrices through deep residual convolutional networks. <i>PLoS Computational Biology</i> , 2021 , 17, e1008865	5	20
265	SEN1-mediated deSUMOylation of JAK2 regulates its kinase activity and platinum drug resistance. <i>Cell Death and Disease</i> , 2021 , 12, 341	9.8	3
264	Saracatinib is an efficacious clinical candidate for fibrodysplasia ossificans progressiva. <i>JCI Insight</i> , 2021 , 6,	9.9	6

263	mRNA therapy restores euglycemia and prevents liver tumors in murine model of glycogen storage disease. <i>Nature Communications</i> , 2021 , 12, 3090	17.4	3
262	Generation of an induced pluripotent stem cell line (TRNDi030-A) from a patient with Farber disease carrying a homozygous p. Y36C (c. 107 A>G) mutation in ASA1. <i>Stem Cell Research</i> , 2021 , 53, 102387	1.6	1
261	Application of niclosamide and analogs as small molecule inhibitors of Zika virus and SARS-CoV-2 infection. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2021 , 40, 127906	2.9	3
260	Viral Proteases as Targets for Coronavirus Disease 2019 Drug Development. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2021 , 378, 166-172	4.7	6
259	Improving high-impact bug report prediction with combination of interactive machine learning and active learning. <i>Information and Software Technology</i> , 2021 , 133, 106530	3.4	25
258	High-throughput screening assays for SARS-CoV-2 drug development: Current status and future directions. <i>Drug Discovery Today</i> , 2021 , 26, 2439-2444	8.8	4
257	Generation of Alagille syndrome derived induced pluripotent stem cell line carrying heterozygous mutation in the JAGGED-1 gene at splicing site (Chr20: 10,629,709C>A) before exon 11. <i>Stem Cell Research</i> , 2021 , 53, 102366	1.6	0
256	Mechanism for DPY30 and ASH2L intrinsically disordered regions to modulate the MLL/SET1 activity on chromatin. <i>Nature Communications</i> , 2021 , 12, 2953	17.4	2
255	Drug combination therapy for emerging viral diseases. <i>Drug Discovery Today</i> , 2021 , 26, 2367-2376	8.8	11
254	Induction of interferon signaling and allograft inflammatory factor 1 in macrophages in a mouse model of breast cancer metastases. <i>Wellcome Open Research</i> , 2021 , 6, 52	4.8	2
253	Enrichment of NPC1-deficient cells with the lipid LBPA stimulates autophagy, improves lysosomal function, and reduces cholesterol storage. <i>Journal of Biological Chemistry</i> , 2021 , 297, 100813	5.4	4
252	Generation of an induced pluripotent stem cell line (TRNDi031-A) from a patient with Alagille syndrome type 1 carrying a heterozygous p. C312X (c. 936T>A) mutation in JAGGED-1. <i>Stem Cell Research</i> , 2021 , 54, 102447	1.6	0
251	Discovery of Small Molecule Entry Inhibitors Targeting the Fusion Peptide of SARS-CoV-2 Spike Protein. <i>ACS Medicinal Chemistry Letters</i> , 2021 , 12, 1267-1274	4.3	5
250	Folding non-homologous proteins by coupling deep-learning contact maps with I-TASSER assembly simulations. <i>Cell Reports Methods</i> , 2021 , 1,		40
249	Generation of an induced pluripotent stem cell line (TRNDi012-B) from Fibrodysplasia Ossificans Progressiva (FOP) patient carrying a heterozygous mutation c. 617G>A in the ACVR1 gene. <i>Stem Cell Research</i> , 2021 , 54, 102424	1.6	
248	Effects of SARS-CoV-2 mutations on protein structures and intraviral protein-protein interactions. <i>Journal of Medical Virology</i> , 2021 , 93, 2132-2140	19.7	48
247	The SARS-CoV-2 Cytopathic Effect Is Blocked by Lysosome Alkalinizing Small Molecules. <i>ACS Infectious Diseases</i> , 2021 , 7, 1389-1408	5.5	39
246	Structural interaction between DISC1 and ATF4 underlying transcriptional and synaptic dysregulation in an iPSC model of mental disorders. <i>Molecular Psychiatry</i> , 2021 , 26, 1346-1360	15.1	13

245	Functions of Essential Genes and a Scale-Free Protein Interaction Network Revealed by Structure-Based Function and Interaction Prediction for a Minimal Genome. <i>Journal of Proteome Research</i> , 2021 , 20, 1178-1189	5.6	7
244	Data Quality Matters: A Case Study on Data Label Correctness for Security Bug Report Prediction. <i>IEEE Transactions on Software Engineering</i> , 2021 , 1-1	3.5	22
243	An Integrated Systems Biology Approach Identifies the Proteasome as A Critical Host Machinery for ZIKV and DENV Replication. <i>Genomics, Proteomics and Bioinformatics</i> , 2021 , 19, 108-122	6.5	3
242	Biological activity-based modeling identifies antiviral leads against SARS-CoV-2. <i>Nature Biotechnology</i> , 2021 , 39, 747-753	44.5	14
241	An induced pluripotent stem cell line (NCATS-CL9075) from a patient carrying compound heterozygote mutations, p.R390P and p.L318P, in the NGLY1 gene. <i>Stem Cell Research</i> , 2021 , 54, 102400 ^{1.6}		
240	Therapeutics Development for Alagille Syndrome. <i>Frontiers in Pharmacology</i> , 2021 , 12, 704586	5.6	3
239	Protein structure prediction using deep learning distance and hydrogen-bonding restraints in CASP14. <i>Proteins: Structure, Function and Bioinformatics</i> , 2021 , 89, 1734-1751	4.2	9
238	Improving fragment-based ab initio protein structure assembly using low-accuracy contact-map predictions. <i>Nature Communications</i> , 2021 , 12, 5011	17.4	6
237	High-throughput protein modification quantitation analysis using intact protein MRM and its application on hENGase inhibitor screening. <i>Talanta</i> , 2021 , 231, 122384	6.2	1
236	Protein inter-residue contact and distance prediction by coupling complementary coevolution features with deep residual networks in CASP14. <i>Proteins: Structure, Function and Bioinformatics</i> , 2021 , 89, 1911-1921	4.2	4
235	Protein structural features predict responsiveness to pharmacological chaperone treatment for three lysosomal storage disorders. <i>PLoS Computational Biology</i> , 2021 , 17, e1009370	5	0
234	Hybrid Approach Reveals Novel Inhibitors of Multiple SARS-CoV-2 Variants. <i>ACS Pharmacology and Translational Science</i> , 2021 , 4, 1675-1688	5.9	2
233	Generation of two gene corrected human isogenic iPSC lines (NCATS-CL6104 and NCATS-CL6105) from a patient line (NCATS-CL6103) carrying a homozygous p.R401X mutation in the NGLY1 gene using CRISPR/Cas9. <i>Stem Cell Research</i> , 2021 , 56, 102554	1.6	1
232	Disease modeling for Mucopolysaccharidosis type IIIB using patient derived induced pluripotent stem cells. <i>Experimental Cell Research</i> , 2021 , 407, 112785	4.2	2
231	Discovery and characterization of potent And-1 inhibitors for cancer treatment.. <i>Clinical and Translational Medicine</i> , 2021 , 11, e627	5.7	3
230	Zika Virus-Induced Neuronal Apoptosis via Increased Mitochondrial Fragmentation. <i>Frontiers in Microbiology</i> , 2020 , 11, 598203	5.7	14
229	Landscape of variable domain of heavy-chain-only antibody repertoire from alpaca. <i>Immunology</i> , 2020 , 161, 53-65	7.8	3
228	Protein Structure and Sequence Reanalysis of 2019-nCoV Genome Refutes Snakes as Its Intermediate Host and the Unique Similarity between Its Spike Protein Insertions and HIV-1. <i>Journal of Proteome Research</i> , 2020 , 19, 1351-1360	5.6	166

227	Generation and characterization of four Chediak-Higashi Syndrome (CHS) induced pluripotent stem cell (iPSC) lines. <i>Stem Cell Research</i> , 2020 , 47, 101883	1.6	0
226	Impact of mRNA chemistry and manufacturing process on innate immune activation. <i>Science Advances</i> , 2020 , 6, eaaz6893	14.3	73
225	Invalid bug reports complicate the software aging situation. <i>Software Quality Journal</i> , 2020 , 28, 195-220	1.2	3
224	FUpred: detecting protein domains through deep-learning-based contact map prediction. <i>Bioinformatics</i> , 2020 , 36, 3749-3757	7.2	15
223	Modeling CNS Involvement in Pompe Disease Using Neural Stem Cells Generated from Patient-Derived Induced Pluripotent Stem Cells. <i>Cells</i> , 2020 , 10,	7.9	2
222	CircRNA-SORE mediates sorafenib resistance in hepatocellular carcinoma by stabilizing YBX1. <i>Signal Transduction and Targeted Therapy</i> , 2020 , 5, 298	2.1	72
221	DeepMSA: constructing deep multiple sequence alignment to improve contact prediction and fold-recognition for distant-homology proteins. <i>Bioinformatics</i> , 2020 , 36, 2105-2112	7.2	60
220	SSIPE: accurately estimating protein-protein binding affinity change upon mutations using evolutionary profiles in combination with an optimized physical energy function. <i>Bioinformatics</i> , 2020 , 36, 2429-2437	7.2	21
219	CVE-assisted large-scale security bug report dataset construction method. <i>Journal of Systems and Software</i> , 2020 , 160, 110456	3.3	10
218	A cell-based, infectious-free, platform to identify inhibitors of lassa virus ribonucleoprotein (vRNP) activity. <i>Antiviral Research</i> , 2020 , 173, 104667	10.8	6
217	Four induced pluripotent stem cell lines (TRNDi021-C, TRNDi023-D, TRNDi024-D and TRNDi025-A) generated from fibroblasts of four healthy individuals. <i>Stem Cell Research</i> , 2020 , 49, 102011	1.6	1
216	Development of a High-Throughput Homogeneous AlphaLISA Drug Screening Assay for the Detection of SARS-CoV-2 Nucleocapsid. <i>ACS Pharmacology and Translational Science</i> , 2020 , 3, 1233-1241	5.9	4
215	RNA-Dependent RNA Polymerase as a Target for COVID-19 Drug Discovery. <i>SLAS Discovery</i> , 2020 , 25, 1141-1151	3.4	64
214	Heparan sulfate assists SARS-CoV-2 in cell entry and can be targeted by approved drugs in vitro. <i>Cell Discovery</i> , 2020 , 6, 80	22.3	86
213	Carbon Dots for Efficient Small Interfering RNA Delivery and Gene Silencing in Plants. <i>Plant Physiology</i> , 2020 , 184, 647-657	6.6	40
212	Drug Discovery Strategies for SARS-CoV-2. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2020 , 375, 127-138	4.7	51
211	Pharmacological clearance of misfolded rhodopsin for the treatment of RHO-associated retinitis pigmentosa. <i>FASEB Journal</i> , 2020 , 34, 10146-10167	0.9	0
210	Human recombinant lysosomal Hexosaminidases produced in <i>Pichia pastoris</i> efficiently reduced lipid accumulation in Tay-Sachs fibroblasts. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2020 , 184, 885-895	3.1	0

209	Identifying SARS-CoV-2 Entry Inhibitors through Drug Repurposing Screens of SARS-S and MERS-S Pseudotyped Particles. <i>ACS Pharmacology and Translational Science</i> , 2020 , 3, 1165-1175	5.9	42
208	Identification of SARS-CoV-2 3CL Protease Inhibitors by a Quantitative High-Throughput Screening. <i>ACS Pharmacology and Translational Science</i> , 2020 , 3, 1008-1016	5.9	76
207	Cell-Based No-Wash Fluorescence Assays for Compound Screens Using a Fluorescence Cytometry Plate Reader. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2020 , 374, 500-511	4.7	
206	Human Pluripotent Stem Cell-Derived Neural Cells and Brain Organoids Reveal SARS-CoV-2 Neurotropism Predominates in Choroid Plexus Epithelium. <i>Cell Stem Cell</i> , 2020 , 27, 937-950.e9	18	151
205	Torin 2 Derivative, NCATS-SM3710, Has Potent Multistage Antimalarial Activity through Inhibition of Phosphatidylinositol 4-Kinase (PI4KIII). <i>ACS Pharmacology and Translational Science</i> , 2020 , 3, 948-964	5.9	8
204	Two-Level Protein Methylation Prediction using structure model-based features. <i>Scientific Reports</i> , 2020 , 10, 6008	4.9	4
203	The Human DNA Mismatch Repair Protein MSH3 Contains Nuclear Localization and Export Signals That Enable Nuclear-Cytosolic Shuttling in Response to Inflammation. <i>Molecular and Cellular Biology</i> , 2020 , 40,	4.8	9
202	Drug Repurposing Screen for Compounds Inhibiting the Cytopathic Effect of SARS-CoV-2. <i>Frontiers in Pharmacology</i> , 2020 , 11, 592737	5.6	39
201	Improving therapy of severe infections through drug repurposing of synergistic combinations. <i>Current Opinion in Pharmacology</i> , 2019 , 48, 92-98	5.1	27
200	Induced pluripotent stem cells for neural drug discovery. <i>Drug Discovery Today</i> , 2019 , 24, 992-999	8.8	43
199	-Tocopherol Effect on Endocytosis and Its Combination with Enzyme Replacement Therapy for Lysosomal Disorders: A New Type of Drug Interaction?. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2019 , 370, 823-833	4.7	2
198	Pharmacological analysis of CFTR variants of cystic fibrosis using stem cell-derived organoids. <i>Drug Discovery Today</i> , 2019 , 24, 2126-2138	8.8	9
197	An induced pluripotent stem cell line (TRNDi009-C) from a Niemann-Pick disease type A patient carrying a heterozygous p.L302P (c.905 T>C) mutation in the SMPD1 gene. <i>Stem Cell Research</i> , 2019 , 38, 101461	1.6	4
196	Identification of Ezetimibe and Pranlukast as Pharmacological Chaperones for the Treatment of the Rare Disease Mucopolysaccharidosis Type IVA. <i>Journal of Medicinal Chemistry</i> , 2019 , 62, 6175-6189	8.3	13
195	Generation of an induced pluripotent stem cell line (TRNDi008-A) from a Hunter syndrome patient carrying a hemizygous 208insC mutation in the IDS gene. <i>Stem Cell Research</i> , 2019 , 37, 101451	1.6	4
194	17-Hydroxy Wortmannin Restores TRAIL α Response by Ameliorating Increased Beclin 1 Level and Autophagy Function in TRAIL-Resistant Colon Cancer Cells. <i>Molecular Cancer Therapeutics</i> , 2019 , 18, 1265-1277 ¹	6.1	
193	Generation of an induced pluripotent stem cell line (TRNDi004-I) from a Niemann-Pick disease type B patient carrying a heterozygous mutation of p.L43_A44delA in the SMPD1 gene. <i>Stem Cell Research</i> , 2019 , 37, 101436	1.6	1
192	I-TASSER gateway: A protein structure and function prediction server powered by XSEDE. <i>Future Generation Computer Systems</i> , 2019 , 99, 73-85	7.5	35

191	A human induced pluripotent stem cell line (TRNDi007-B) from an infantile onset Pompe patient carrying p.R854X mutation in the GAA gene. <i>Stem Cell Research</i> , 2019 , 37, 101435	1.6	7
190	High-Throughput Zika Viral Titer Assay for Rapid Screening of Antiviral Drugs. <i>Assay and Drug Development Technologies</i> , 2019 , 17, 128-139	2.1	6
189	An induced pluripotent stem cell line (TRNDi006-A) from a MPS IIIB patient carrying homozygous mutation of p.Glu153Lys in the NAGLU gene. <i>Stem Cell Research</i> , 2019 , 37, 101427	1.6	4
188	Identification, design and synthesis of novel pyrazolopyridine influenza virus nonstructural protein 1 antagonists. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2019 , 29, 1113-1119	2.9	4
187	Towards understanding bugs in an open source cloud management stack: An empirical study of OpenStack software bugs. <i>Journal of Systems and Software</i> , 2019 , 151, 210-223	3.3	6
186	Generation of an induced pluripotent stem cell line (TRNDi005-A) from a Mucopolysaccharidosis Type IVA (MPS IVA) patient carrying compound heterozygous p.R61W and p.WT405del mutations in the GALNS gene. <i>Stem Cell Research</i> , 2019 , 36, 101408	1.6	3
185	Advancing precision medicine with personalized drug screening. <i>Drug Discovery Today</i> , 2019 , 24, 272-278.8	14	
184	LOMETS2: improved meta-threading server for fold-recognition and structure-based function annotation for distant-homology proteins. <i>Nucleic Acids Research</i> , 2019 , 47, W429-W436	20.1	76
183	ERK Regulates HIF1 β Mediated Platinum Resistance by Directly Targeting PHD2 in Ovarian Cancer. <i>Clinical Cancer Research</i> , 2019 , 25, 5947-5960	12.9	19
182	An induced pluripotent stem cell line (TRNDi010-C) from a patient carrying a homozygous p.R401X mutation in the NGLY1 gene. <i>Stem Cell Research</i> , 2019 , 39, 101496	1.6	1
181	Deep-learning contact-map guided protein structure prediction in CASP13. <i>Proteins: Structure, Function and Bioinformatics</i> , 2019 , 87, 1149-1164	4.2	108
180	Detecting distant-homology protein structures by aligning deep neural-network based contact maps. <i>PLoS Computational Biology</i> , 2019 , 15, e1007411	5	25
179	Lung Mammary Metastases but Not Primary Tumors Induce Accumulation of Atypical Large Platelets and Their Chemokine Expression. <i>Cell Reports</i> , 2019 , 29, 1747-1755.e4	10.6	7
178	Phosphocyclocreatine is the dominant form of cyclocreatine in control and creatine transporter deficiency patient fibroblasts. <i>Pharmacology Research and Perspectives</i> , 2019 , 7, e00525	3.1	4
177	Cryo-EM structure of the human MLL1 core complex bound to the nucleosome. <i>Nature Communications</i> , 2019 , 10, 5540	17.4	24
176	Quantitative Chemotherapeutic Profiling of Gynecologic Cancer Cell Lines Using Approved Drugs and Bioactive Compounds. <i>Translational Oncology</i> , 2019 , 12, 441-452	4.9	9
175	Generation of an induced pluripotent stem cell line (TRNDi003-A) from a Noonan syndrome with multiple lentigines (NSML) patient carrying a p.Q510P mutation in the PTPN11 gene. <i>Stem Cell Research</i> , 2019 , 34, 101374	1.6	6
174	Generation of an induced pluripotent stem cell line (TRNDi002-B) from a patient carrying compound heterozygous p.Q208X and p.G310G mutations in the NGLY1 gene. <i>Stem Cell Research</i> , 2019 , 34, 101362	1.6	5

173	MetaGO: Predicting Gene Ontology of Non-homologous Proteins Through Low-Resolution Protein Structure Prediction and Protein-Protein Network Mapping. <i>Journal of Molecular Biology</i> , 2018 , 430, 2256-2265	6.5	32
172	Systemic Medication Associations with Presumed Advanced or Uncontrolled Primary Open-Angle Glaucoma. <i>Ophthalmology</i> , 2018 , 125, 984-993	7.3	29
171	Astrocytes as targets for drug discovery. <i>Drug Discovery Today</i> , 2018 , 23, 673-680	8.8	27
170	Neural stem cells for disease modeling and evaluation of therapeutics for infantile (CLN1/PPT1) and late infantile (CLN2/TPP1) neuronal ceroid lipofuscinoses. <i>Orphanet Journal of Rare Diseases</i> , 2018 , 13, 54	4.2	23
169	DUOXA1-mediated ROS production promotes cisplatin resistance by activating ATR-Chk1 pathway in ovarian cancer. <i>Cancer Letters</i> , 2018 , 428, 104-116	9.9	41
168	Drug repurposing screens and synergistic drug-combinations for infectious diseases. <i>British Journal of Pharmacology</i> , 2018 , 175, 181-191	8.6	111
167	Repurposing a novel parathyroid hormone analogue to treat hypoparathyroidism. <i>British Journal of Pharmacology</i> , 2018 , 175, 262-271	8.6	11
166	A large-scale comparative assessment of methods for residue-residue contact prediction. <i>Briefings in Bioinformatics</i> , 2018 , 19, 219-230	13.4	20
165	Small Molecules Identified from a Quantitative Drug Combinational Screen Resensitize Cisplatin Response in Drug-Resistant Ovarian Cancer Cells. <i>Translational Oncology</i> , 2018 , 11, 1053-1064	4.9	5
164	Patient iPSC-derived neural stem cells exhibit phenotypes in concordance with the clinical severity of mucopolysaccharidosis I. <i>Human Molecular Genetics</i> , 2018 , 27, 3612-3626	5.6	17
163	Pluripotent Stem Cell Platforms for Drug Discovery. <i>Trends in Molecular Medicine</i> , 2018 , 24, 805-820	11.5	24
162	Emetine inhibits Zika and Ebola virus infections through two molecular mechanisms: inhibiting viral replication and decreasing viral entry. <i>Cell Discovery</i> , 2018 , 4, 31	22.3	81
161	Drugging SUMOylation for neuroprotection and oncotherapy. <i>Neural Regeneration Research</i> , 2018 , 13, 415-416	4.5	8
160	Repurposing Screen Identifies Unconventional Drugs With Activity Against Multidrug Resistant. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018 , 8, 438	5.9	18
159	Zika Virus: Origins, Pathological Action, and Treatment Strategies. <i>Frontiers in Microbiology</i> , 2018 , 9, 3253-7	3.7	24
158	Quantitative high-throughput screening identifies cytoprotective molecules that enhance SUMO conjugation via the inhibition of SUMO-specific protease (SENP)2. <i>FASEB Journal</i> , 2018 , 32, 1677-1691	0.9	22
157	Canvass: A Crowd-Sourced, Natural-Product Screening Library for Exploring Biological Space. <i>ACS Central Science</i> , 2018 , 4, 1727-1741	16.8	26
156	"Real-Time" High-Throughput Drug and Synergy Testing for Multidrug-Resistant Bacterial Infection: A Case Report. <i>Frontiers in Medicine</i> , 2018 , 5, 267	4.9	3

155	Neural stem cells for disease modeling and evaluation of therapeutics for Tay-Sachs disease. <i>Orphanet Journal of Rare Diseases</i> , 2018 , 13, 152	4.2	20
154	Memory-Enhanced Dynamic Multi-Objective Evolutionary Algorithm Based on Lp Decomposition. <i>Applied Sciences (Switzerland)</i> , 2018 , 8, 1673	2.6	12
153	Discovery of 3-(4-sulfamoylnaphthyl)pyrazolo[1,5-a]pyrimidines as potent and selective ALK2 inhibitors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2018 , 28, 3356-3362	2.9	11
152	Metarrestin, a perinucleolar compartment inhibitor, effectively suppresses metastasis. <i>Science Translational Medicine</i> , 2018 , 10,	17.5	34
151	Optimization of the first small-molecule relaxin/insulin-like family peptide receptor (RXFP1) agonists: Activation results in an antifibrotic gene expression profile. <i>European Journal of Medicinal Chemistry</i> , 2018 , 156, 79-92	6.8	7
150	A Novel Method for Drug Screen to Regulate G Protein-Coupled Receptors in the Metabolic Network of Alzheimer's Disease. <i>BioMed Research International</i> , 2018 , 2018, 5486403	3	2
149	Methyl- β -cyclodextrin restores impaired autophagy flux in Niemann-Pick C1-deficient cells through activation of AMPK. <i>Autophagy</i> , 2017 , 13, 1435-1451	10.2	52
148	Development of an Aryloxazole Class of Hepatitis C Virus Inhibitors Targeting the Entry Stage of the Viral Replication Cycle. <i>Journal of Medicinal Chemistry</i> , 2017 , 60, 6364-6383	8.3	9
147	PrAS: Prediction of amidation sites using multiple feature extraction. <i>Computational Biology and Chemistry</i> , 2017 , 66, 57-62	3.6	6
146	BindProfX: Assessing Mutation-Induced Binding Affinity Change by Protein Interface Profiles with Pseudo-Counts. <i>Journal of Molecular Biology</i> , 2017 , 429, 426-434	6.5	65
145	What DKK1 tells where to metastasize. <i>Nature Cell Biology</i> , 2017 , 19, 1146-1148	23.4	1
144	Efficient Synthesis of 1,9-Substituted Benzo[h][1,6]naphthyridin-2(1H)-ones and Evaluation of their Plasmodium falciparum Gametocytocidal Activities. <i>ACS Combinatorial Science</i> , 2017 , 19, 748-754	3.9	4
143	Identification of 4-phenylquinolin-2(1H)-one as a specific allosteric inhibitor of Akt. <i>Scientific Reports</i> , 2017 , 7, 11673	4.9	5
142	Drug discovery and development for rare genetic disorders. <i>American Journal of Medical Genetics, Part A</i> , 2017 , 173, 2307-2322	2.5	41
141	Neural stem cells for disease modeling of Wolman disease and evaluation of therapeutics. <i>Orphanet Journal of Rare Diseases</i> , 2017 , 12, 120	4.2	14
140	Synergistic drug combination effectively blocks Ebola virus infection. <i>Antiviral Research</i> , 2017 , 137, 165-172	17.28	58
139	An Improved MOEA/D with Optimal DE Schemes for Many-Objective Optimization Problems. <i>Algorithms</i> , 2017 , 10, 86	1.8	1
138	Identification of small-molecule inhibitors of Zika virus infection and induced neural cell death via a drug repurposing screen. <i>Nature Medicine</i> , 2016 , 22, 1101-1107	50.5	458

137	Molecular signatures associated with ZIKV exposure in human cortical neural progenitors. <i>Nucleic Acids Research</i> , 2016 , 44, 8610-8620	20.1	119
136	A New Glucocerebrosidase Chaperone Reduces β Synuclein and Glycolipid Levels in iPSC-Derived Dopaminergic Neurons from Patients with Gaucher Disease and Parkinsonism. <i>Journal of Neuroscience</i> , 2016 , 36, 7441-52	6.6	150
135	Rapid antimicrobial susceptibility test for identification of new therapeutics and drug combinations against multidrug-resistant bacteria. <i>Emerging Microbes and Infections</i> , 2016 , 5, e116	18.9	45
134	Inhibiting macrophage PI3K to enhance immunotherapy. <i>Cell Research</i> , 2016 , 26, 1267-1268	24.7	14
133	Identification of Multiple Cryptococcal Fungicidal Drug Targets by Combined Gene Dosing and Drug Affinity Responsive Target Stability Screening. <i>MBio</i> , 2016 , 7,	7.8	11
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