

Ying Hong Li

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8918526/publications.pdf>

Version: 2024-02-01

21
papers

2,332
citations

471477

17
h-index

713444

21
g-index

21
all docs

21
docs citations

21
times ranked

3493
citing authors

#	ARTICLE	IF	CITATIONS
1	Therapeutic target database 2020: enriched resource for facilitating research and early development of targeted therapeutics. <i>Nucleic Acids Research</i> , 2020, 48, D1031-D1041.	14.5	488
2	Therapeutic target database update 2018: enriched resource for facilitating bench-to-clinic research of targeted therapeutics. <i>Nucleic Acids Research</i> , 2018, 46, D1121-D1127.	14.5	462
3	NOREVA: normalization and evaluation of MS-based metabolomics data. <i>Nucleic Acids Research</i> , 2017, 45, W162-W170.	14.5	305
4	Therapeutic target database update 2016: enriched resource for bench to clinical drug target and targeted pathway information. <i>Nucleic Acids Research</i> , 2016, 44, D1069-D1074.	14.5	278
5	Clinical trials, progression-speed differentiating features and swiftness rule of the innovative targets of first-in-class drugs. <i>Briefings in Bioinformatics</i> , 2020, 21, 649-662.	6.5	139
6	Therapeutic target database update 2014: a resource for targeted therapeutics. <i>Nucleic Acids Research</i> , 2014, 42, D1118-D1123.	14.5	116
7	SVM-Prot 2016: A Web-Server for Machine Learning Prediction of Protein Functional Families from Sequence Irrespective of Similarity. <i>PLoS ONE</i> , 2016, 11, e0155290.	2.5	98
8	Identification of the inhibitory mechanism of FDA approved selective serotonin reuptake inhibitors: an insight from molecular dynamics simulation study. <i>Physical Chemistry Chemical Physics</i> , 2016, 18, 3260-3271.	2.8	66
9	The Human Kinome Targeted by FDA Approved Multi-Target Drugs and Combination Products: A Comparative Study from the Drug-Target Interaction Network Perspective. <i>PLoS ONE</i> , 2016, 11, e0165737.	2.5	51
10	MMEASE: Online meta-analysis of metabolomic data by enhanced metabolite annotation, marker selection and enrichment analysis. <i>Journal of Proteomics</i> , 2021, 232, 104023.	2.4	50
11	Exploring the Inhibitory Mechanism of Approved Selective Norepinephrine Reuptake Inhibitors and Reboxetine Enantiomers by Molecular Dynamics Study. <i>Scientific Reports</i> , 2016, 6, 26883.	3.3	46
12	Comparison of FDA Approved Kinase Targets to Clinical Trial Ones: Insights from Their System Profiles and Drug-Target Interaction Networks. <i>BioMed Research International</i> , 2016, 2016, 1-9.	1.9	36
13	Assessing the Performances of Protein Function Prediction Algorithms from the Perspectives of Identification Accuracy and False Discovery Rate. <i>International Journal of Molecular Sciences</i> , 2018, 19, 183.	4.1	35
14	Prediction of the binding mode and resistance profile for a dual-target pyrrolyl diketo acid scaffold against HIV-1 integrase and reverse-transcriptase-associated ribonuclease H. <i>Physical Chemistry Chemical Physics</i> , 2018, 20, 23873-23884.	2.8	31
15	Determining the Balance Between Drug Efficacy and Safety by the Network and Biological System Profile of Its Therapeutic Target. <i>Frontiers in Pharmacology</i> , 2018, 9, 1245.	3.5	28
16	ExoBCD: a comprehensive database for exosomal biomarker discovery in breast cancer. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	23
17	Comparison of computational model and X-ray crystal structure of human serotonin transporter: potential application for the pharmacology of human monoamine transporters. <i>Molecular Simulation</i> , 2017, 43, 1089-1098.	2.0	21
18	Consistent Biomarkers and Related Pathogenesis Underlying Asthma Revealed by Systems Biology Approach. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4037.	4.1	20

#	ARTICLE	IF	CITATIONS
19	What Makes Species Productive of Anti-Cancer Drugs? Clues from Drugs'™ Species Origin, Druglikeness, Target and Pathway. <i>Anti-Cancer Agents in Medicinal Chemistry</i> , 2019, 19, 194-203.	1.7	16
20	Identification of Potential Biomarkers for Anti-PD-1 Therapy in Melanoma by Weighted Correlation Network Analysis. <i>Genes</i> , 2020, 11, 435.	2.4	14
21	A Cancer Associated Fibroblasts-Related Six-Gene Panel for Anti-PD-1 Therapy in Melanoma Driven by Weighted Correlation Network Analysis and Supervised Machine Learning. <i>Frontiers in Medicine</i> , 2022, 9, 880326.	2.6	9