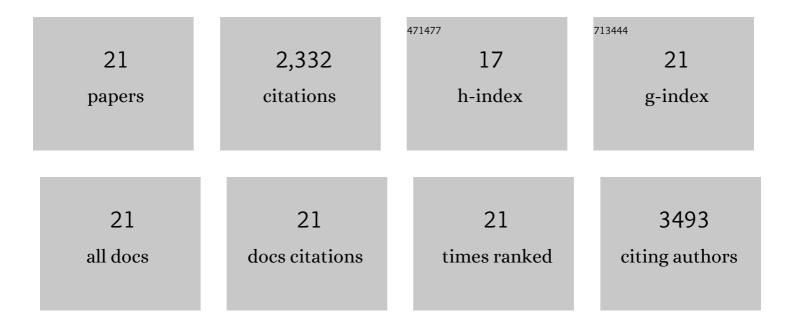
## Ying Hong Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8918526/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Therapeutic target database 2020: enriched resource for facilitating research and early development of targeted therapeutics. Nucleic Acids Research, 2020, 48, D1031-D1041.	14.5	488
2	Therapeutic target database update 2018: enriched resource for facilitating bench-to-clinic research of targeted therapeutics. Nucleic Acids Research, 2018, 46, D1121-D1127.	14.5	462
3	NOREVA: normalization and evaluation of MS-based metabolomics data. Nucleic Acids Research, 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. Nucleic Acids Research, 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. Briefings in Bioinformatics, 2020, 21, 649-662.	6.5	139
6	Therapeutic target database update 2014: a resource for targeted therapeutics. Nucleic Acids Research, 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. PLoS ONE, 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. Physical Chemistry Chemical Physics, 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. PLoS ONE, 2016, 11, e0165737.	2.5	51
10	MMEASE: Online meta-analysis of metabolomic data by enhanced metabolite annotation, marker selection and enrichment analysis. Journal of Proteomics, 2021, 232, 104023.	2.4	50
11	Exploring the Inhibitory Mechanism of Approved Selective Norepinephrine Reuptake Inhibitors and Reboxetine Enantiomers by Molecular Dynamics Study. Scientific Reports, 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. BioMed Research International, 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. International Journal of Molecular Sciences, 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. Physical Chemistry Chemical Physics, 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. Frontiers in Pharmacology, 2018, 9, 1245.	3.5	28
16	ExoBCD: a comprehensive database for exosomal biomarker discovery in breast cancer. Briefings in Bioinformatics, 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. Molecular Simulation, 2017, 43, 1089-1098.	2.0	21
18	Consistent Biomarkers and Related Pathogenesis Underlying Asthma Revealed by Systems Biology Approach. International Journal of Molecular Sciences, 2019, 20, 4037.	4.1	20

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#	Article	IF	CITATIONS
19	What Makes Species Productive of Anti-Cancer Drugs? Clues from Drugs' Species Origin, Druglikeness, Target and Pathway. Anti-Cancer Agents in Medicinal Chemistry, 2019, 19, 194-203.	1.7	16
20	Identification of Potential Biomarkers for Anti-PD-1 Therapy in Melanoma by Weighted Correlation Network Analysis. Genes, 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. Frontiers in Medicine, 2022, 9, 880326.	2.6	9