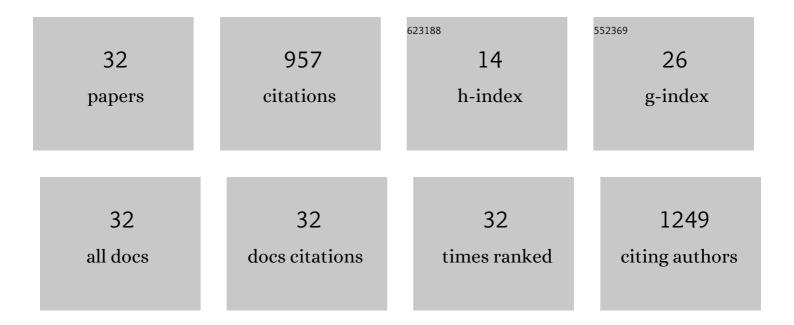
Yitian Zhou

List of Publications by Year in descending order

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ΥιτιλΝ ΖΗΟΙΙ

#	Article	IF	CITATIONS
1	Pharmacogenomics of statins: lipid response and other outcomes in Brazilian cohorts. Pharmacological Reports, 2022, 74, 47-66.	1.5	7
2	Population pharmacogenomics: an update on ethnogeographic differences and opportunities for precision public health. Human Genetics, 2022, 141, 1113-1136.	1.8	17
3	Novel loss-of-function variant in DENND5A impedes melanosomal cargo transport and predisposes to familial cutaneous melanoma. Genetics in Medicine, 2022, 24, 157-169.	1.1	0
4	Computational Methods and Approaches in Pharmacogenomic Research. , 2022, , 53-83.		1
5	Organotypic and Microphysiological Human Tissue Models for Drug Discovery and Development—Current State-of-the-Art and Future Perspectives. Pharmacological Reviews, 2022, 74, 141-206.	7.1	23
6	Identification and functional validation of novel pharmacogenomic variants using a next-generation sequencing-based approach for clinical pharmacogenomics. Pharmacological Research, 2022, 176, 106087.	3.1	9
7	Pharmacogenomics of Antihypertensive Drugs in Brazil: Recent Progress and Clinical Implications. Endocrine, Metabolic and Immune Disorders - Drug Targets, 2022, 22, 1263-1275.	0.6	1
8	Genetic Variant ABCC1 rs45511401 Is Associated with Increased Response to Statins in Patients with Familial Hypercholesterolemia. Pharmaceutics, 2022, 14, 944.	2.0	4
9	Genome-wide association meta-analysis identifies 48 risk variants and highlights the role of the stria vascularis in hearing loss. American Journal of Human Genetics, 2022, 109, 1077-1091.	2.6	27
10	Global Frequencies of Clinically Important <i>HLA</i> Alleles and Their Implications For the Costâ€Effectiveness of Preemptive Pharmacogenetic Testing. Clinical Pharmacology and Therapeutics, 2021, 109, 160-174.	2.3	35
11	Pharmacogenetics of tamoxifen therapy in Asian populations: from genetic polymorphism to clinical outcomes. European Journal of Clinical Pharmacology, 2021, 77, 1095-1111.	0.8	8
12	Pharmacogenomics in the era of next generation sequencing – from byte to bedside. Drug Metabolism Reviews, 2021, 53, 253-278.	1.5	19
13	Computational Tools to Assess the Functional Consequences of Rare and Noncoding Pharmacogenetic Variability. Clinical Pharmacology and Therapeutics, 2021, 110, 626-636.	2.3	16
14	Editorial: Population Pharmacogenomics (PGx): From Variant Identification to Clinical Implementation. Frontiers in Genetics, 2021, 12, 736626.	1.1	1
15	Cardiovascular Pharmacogenomics: An Update on Clinical Studies of Antithrombotic Drugs in Brazilian Patients. Molecular Diagnosis and Therapy, 2021, 25, 735-755.	1.6	3
16	Rare genetic variability in human drug target genes modulates drug response and can guide precision medicine. Science Advances, 2021, 7, eabi6856.	4.7	16
17	Pharmacogenomic network analysis of the gene-drug interaction landscape underlying drug disposition. Computational and Structural Biotechnology Journal, 2020, 18, 52-58.	1.9	3
18	Population-scale predictions of DPD and TPMT phenotypes using a quantitative pharmacogene-specific ensemble classifier. British Journal of Cancer, 2020, 123, 1782-1789.	2.9	23

Υιτιαν Ζηου

#	Article	IF	CITATIONS
19	Impact of variants in ATP-binding cassetteÂtransporters on breast cancer treatment. Pharmacogenomics, 2020, 21, 1299-1310.	0.6	9
20	Ethnogeographic and inter-individual variability of human ABC transporters. Human Genetics, 2020, 139, 623-646.	1.8	34
21	Germline variant burden in multidrug resistance transporters is a therapyâ€specific predictor of survival in breast cancer patients. International Journal of Cancer, 2020, 146, 2475-2487.	2.3	20
22	In Vitro Functional Characterization and in Silico Prediction of Rare Genetic Variation in the Bile Acid and Drug Transporter, Na ⁺ -Taurocholate Cotransporting Polypeptide (NTCP,) Tj ETQq0 0 0 rgBT /Ov	verbosik 10	Tf 1540 617 Td
23	CYP3A5 is unlikely to mediate anticancer drug resistance in hepatocellular carcinoma. Pharmacogenomics, 2019, 20, 1085-1092.	0.6	2
24	Comprehensive Evaluation of Organotypic and Microphysiological Liver Models for Prediction of Drug-Induced Liver Injury. Frontiers in Pharmacology, 2019, 10, 1093.	1.6	62
25	Novel genetic and epigenetic factors of importance for inter-individual differences in drug disposition, response and toxicity. , 2019, 197, 122-152.		83
26	An optimized prediction framework to assess the functional impact of pharmacogenetic variants. Pharmacogenomics Journal, 2019, 19, 115-126.	0.9	109
27	Prediction of Drug-Induced Hepatotoxicity Using Long-Term Stable Primary Hepatic 3D Spheroid Cultures in Chemically Defined Conditions. Toxicological Sciences, 2018, 163, 655-665.	1.4	140
28	Computational Methods for the Pharmacogenetic Interpretation of Next Generation Sequencing Data. Frontiers in Pharmacology, 2018, 9, 1437.	1.6	62
29	Integrating rare genetic variants into pharmacogenetic drug response predictions. Human Genomics, 2018, 12, 26.	1.4	166
30	Comprehensive overview of the pharmacogenetic diversity in Ashkenazi Jews. Journal of Medical Genetics, 2018, 55, 617-627.	1.5	24
31	Global genetic diversity of human apolipoproteins and effects on cardiovascular disease risk. Journal of Lipid Research, 2018, 59, 1987-2000.	2.0	19
32	Prediction of drug-induced hepatotoxicity using long-term stable primary hepatic 3D spheroid cultures. Proceedings for Annual Meeting of the Japanese Pharmacological Society, 2018, WCP2018, PO4-9-16.	0.0	0