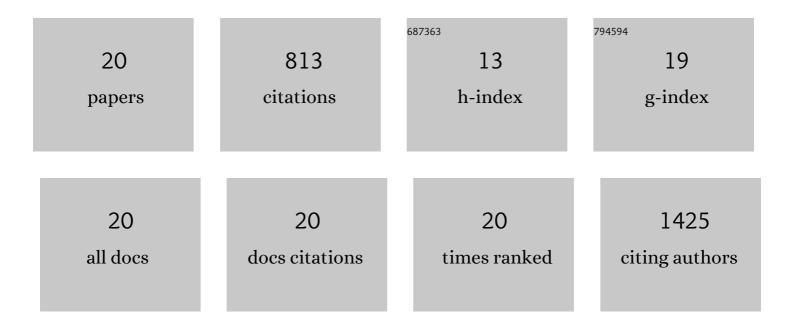
Yao Yang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Development and Analytical Validation of a 29 Gene Clinical Pharmacogenetic Genotyping Panel: Multiâ€Ethnic Allele and Copy Number Variant Detection. Clinical and Translational Science, 2021, 14, 204-213.	3.1	16
2	Differential flora in the microenvironment of lung tumor and paired adjacent normal tissues. Carcinogenesis, 2020, 41, 1094-1103.	2.8	11
3	Phased Haplotype Resolution of the SLC6A4 Promoter Using Long-Read Single Molecule Real-Time (SMRT) Sequencing. Genes, 2020, 11, 1333.	2.4	5
4	Integrated CYP2D6 interrogation for multiethnic copy number and tandem allele detection. Pharmacogenomics, 2019, 20, 9-20.	1.3	9
5	Cytogenomic identification and long-read single molecule real-time (SMRT) sequencing of a Bardet–Biedl Syndrome 9 (BBS9) deletion. Npj Genomic Medicine, 2018, 3, 3.	3.8	97
6	Interplay between the lung microbiome and lung cancer. Cancer Letters, 2018, 415, 40-48.	7.2	188
7	A network-based signature to predict the survival of non-smoking lung adenocarcinoma. Cancer Management and Research, 2018, Volume 10, 2683-2693.	1.9	9
8	Sequencing the <i>CYP2D6</i> gene: from variant allele discovery to clinical pharmacogenetic testing. Pharmacogenomics, 2017, 18, 673-685.	1.3	105
9	DNA Methylation Profiling Using Long-Read Single Molecule Real-Time Bisulfite Sequencing (SMRT-BS). Methods in Molecular Biology, 2017, 1654, 125-134.	0.9	21
10	Institutional profile: translational pharmacogenomics at the Icahn School of Medicine at Mount Sinai. Pharmacogenomics, 2017, 18, 1381-1386.	1.3	20
11	Long-Read Single Molecule Real-Time Full Gene Sequencing of Cytochrome P450-2D6. Human Mutation, 2016, 37, 315-323.	2.5	86
12	Genetic variation at glucose and insulin trait loci and response to glucose–insulin–potassium (GIK) therapy: the IMMEDIATE trial. Pharmacogenomics Journal, 2015, 15, 55-62.	2.0	6
13	The pharmacogenetic control of antiplatelet response: candidate genes and <i>CYP2C19</i> . Expert Opinion on Drug Metabolism and Toxicology, 2015, 11, 1599-1617.	3.3	22
14	Quantitative and multiplexed DNA methylation analysis using long-read single-molecule real-time bisulfite sequencing (SMRT-BS). BMC Genomics, 2015, 16, 350.	2.8	68
15	Delivery Mode Shaped the Gut Microbiome in Chinese Newborns. Journal of Pediatric Gastroenterology and Nutrition, 2015, 60, 149-150.	1.8	0
16	Genetic Modifiers of Cardiorespiratory Fitness Response to Lifestyle Intervention. Medicine and Science in Sports and Exercise, 2014, 46, 302-311.	0.4	30
17	Pharmacogenetics in Jewish populations. Drug Metabolism and Drug Interactions, 2014, 29, 221-233.	0.3	14
18	An Allele-Specific PCR System for Rapid Detection and Discrimination of the CYP2C19â^—4A, â^—4B, and â^—17 Alleles. Journal of Molecular Diagnostics, 2013, 15, 783-789.	2.8	17

#	Article	IF	CITATIONS
19	Frequency of the cholesteryl ester storage disease common <i>LIPA</i> E8SJM mutation (c.894G>A) in various racial and ethnic groups. Hepatology, 2013, 58, 958-965.	7.3	85
20	Identification of methylated regions with peak search based on Poisson model from massively parallel methylated DNA immunoprecipitationâ€sequencing data. Electrophoresis, 2010, 31, 3537-3544.	2.4	4