

# Chuan Qiu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/575730/publications.pdf>

Version: 2024-02-01

46  
papers

1,269  
citations

430874

18  
h-index

377865

34  
g-index

47  
all docs

47  
docs citations

47  
times ranked

2293  
citing authors

#	ARTICLE	IF	CITATIONS
1	Pathway-based metabolomics study of sarcopenia-related traits in two US cohorts. <i>Aging</i> , 2022, 14, 2101-2112.	3.1	5
2	Integrative analysis of multi-omics data to detect the underlying molecular mechanisms for obesity in vivo in humans. <i>Human Genomics</i> , 2022, 16, 15.	2.9	6
3	A bi-directional Mendelian randomization study of the sarcopenia-related traits and osteoporosis. <i>Aging</i> , 2022, , 5681-5698.	3.1	7
4	Activation of RAGE-dependent endoplasmic reticulum stress associates with exacerbated postmyocardial infarction ventricular arrhythmias in diabetes. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2021, 320, E539-E550.	3.5	6
5	Identification of novel functional CpG-SNPs associated with Type 2 diabetes and birth weight. <i>Aging</i> , 2021, 13, 10619-10658.	3.1	5
6	A generalized kernel machine approach to identify higher-order composite effects in multi-view datasets, with application to adolescent brain development and osteoporosis. <i>Journal of Biomedical Informatics</i> , 2021, 120, 103854.	4.3	2
7	Rivaroxaban Suppresses Atherosclerosis by Inhibiting FXa-Induced Macrophage M1 Polarization-Mediated Phenotypic Conversion of Vascular Smooth Muscle Cells. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 739212.	2.4	12
8	Multi-omics Data Integration for Identifying Osteoporosis Biomarkers and Their Biological Interaction and Causal Mechanisms. <i>iScience</i> , 2020, 23, 100847.	4.1	48
9	Identification of pleiotropic genes between risk factors of stroke by multivariate metaCCA analysis. <i>Molecular Genetics and Genomics</i> , 2020, 295, 1173-1185.	2.1	5
10	Mendelian Randomization Identifies CpG Methylation Sites With Mediation Effects for Genetic Influences on BMD in Peripheral Blood Monocytes. <i>Frontiers in Genetics</i> , 2020, 11, 60.	2.3	9
11	Identification of novel functional CpG-SNPs associated with type 2 diabetes and coronary artery disease. <i>Molecular Genetics and Genomics</i> , 2020, 295, 607-619.	2.1	11
12	Diabetes mellitus exacerbates postmyocardial infarction heart failure by reducing sarcolipin promoter methylation. <i>ESC Heart Failure</i> , 2020, 7, 1935-1948.	3.1	16
13	Novel Prognostic Model for Gastric Cancer using 13 Co-Expression Long Non-Coding RNAs (LncRNAs). <i>Medical Science Monitor</i> , 2020, 26, e923295.	1.1	1
14	Ear Crease Features Are Associated with Complexity of Coronary Lesions. <i>Medical Science Monitor</i> , 2020, 26, e923343.	1.1	3
15	Integrative genomic analysis predicts novel functional enhancer-SNPs for bone mineral density. <i>Human Genetics</i> , 2019, 138, 167-185.	3.8	4
16	MIR-21-3p Plays a Crucial Role in Metabolism Alteration of Renal Tubular Epithelial Cells during Sepsis Associated Acute Kidney Injury via AKT/CDK2-FOXO1 Pathway. <i>BioMed Research International</i> , 2019, 2019, 1-12.	1.9	25
17	Matrine alleviates AGEs- induced cardiac dysfunctions by attenuating calcium overload via reducing ryanodine receptor 2 activity. <i>European Journal of Pharmacology</i> , 2019, 842, 118-124.	3.5	24
18	Matrine suppresses cardiac fibrosis by inhibiting the TGF $\beta$ <sup>1</sup> /Smad pathway in experimental diabetic cardiomyopathy. <i>Molecular Medicine Reports</i> , 2018, 17, 1775-1781.	2.4	41

#	ARTICLE	IF	CITATIONS
19	Integrative functional analysis of super enhancer SNPs for coronary artery disease. <i>Journal of Human Genetics</i> , 2018, 63, 627-638.	2.3	29
20	Matrine suppresses AGE-induced HAEC injury by inhibiting ROS-mediated NLRP3 inflammasome activation. <i>European Journal of Pharmacology</i> , 2018, 822, 207-211.	3.5	25
21	Novel <i>ASK1</i> inhibitor <i>AGI-1067</i> improves AGE-induced cardiac dysfunction by inhibiting <i>MKK3/p38</i> <i>MAPK</i> and <i>NF-<math>\kappa</math>B</i> apoptotic signaling. <i>FEBS Open Bio</i> , 2018, 8, 1445-1456.	2.3	5
22	Meta-Analysis of Genome-Wide Association Studies Identifies Novel Functional CpG-SNPs Associated with Bone Mineral Density at Lumbar Spine. <i>International Journal of Genomics</i> , 2018, 2018, 1-11.	1.6	9
23	Matrine blocks AGEs- induced HSCMCs phenotypic conversion via suppressing Dll4-Notch pathway. <i>European Journal of Pharmacology</i> , 2018, 835, 126-131.	3.5	16
24	Metabolomic profiles associated with bone mineral density in US Caucasian women. <i>Nutrition and Metabolism</i> , 2018, 15, 57.	3.0	51
25	A joint analysis of metabolomic profiles associated with muscle mass and strength in Caucasian women. <i>Aging</i> , 2018, 10, 2624-2635.	3.1	18
26	Matrine attenuates cardiac fibrosis by affecting ATF6 signaling pathway in diabetic cardiomyopathy. <i>European Journal of Pharmacology</i> , 2017, 804, 21-30.	3.5	40
27	Matrine-type Alkaloids Inhibit Advanced Glycation End Products Induced Reactive Oxygen Species-Mediated Apoptosis of Aortic Endothelial Cells In Vivo and In Vitro by Targeting MKK3 and p38MAPK Signaling. <i>Journal of the American Heart Association</i> , 2017, 6, .	3.7	26
28	Adenosine triphosphate-sensitive potassium channels and cardiomyopathies (Review). <i>Molecular Medicine Reports</i> , 2016, 13, 1447-1454.	2.4	6
29	Clinical Epigenetics and Epigenomics. <i>Translational Bioinformatics</i> , 2016, , 269-293.	0.0	0
30	Matrine pretreatment improves cardiac function in rats with diabetic cardiomyopathy via suppressing ROS/TLR-4 signaling pathway. <i>Acta Pharmacologica Sinica</i> , 2015, 36, 323-333.	6.1	70
31	Integrative Analysis of Transcriptomic and Epigenomic Data to Reveal Regulation Patterns for BMD Variation. <i>PLoS ONE</i> , 2015, 10, e0138524.	2.5	25
32	P638PERK- a potential molecular regulator of calcium homeostasis related with arrhythmia in diabetic cardiomyopathy. <i>Cardiovascular Research</i> , 2014, 103, S116.2-S116.	3.8	0
33	Protein kinase RNA-like endoplasmic reticulum kinase (PERK)/calcineurin signaling is a novel pathway regulating intracellular calcium accumulation which might be involved in ventricular arrhythmias in diabetic cardiomyopathy. <i>Cellular Signalling</i> , 2014, 26, 2591-2600.	3.6	57
34	Selenium Attenuates Adriamycin-Induced Cardiac Dysfunction via Restoring Expression of ATP-Sensitive Potassium Channels in Rats. <i>Biological Trace Element Research</i> , 2013, 153, 220-228.	3.5	10
35	Characterization of the DNA methylome and its interindividual variation in human peripheral blood monocytes. <i>Epigenomics</i> , 2013, 5, 255-269.	2.1	19
36	Selenium Attenuates High Glucose-Induced ROS/TLR-4 Involved Apoptosis of Rat Cardiomyocyte. <i>Biological Trace Element Research</i> , 2013, 156, 262-270.	3.5	28

#	ARTICLE	IF	CITATIONS
37	Protein kinase RNA- like endoplasmic reticulum kinase (PERK) signaling pathway plays a major role in reactive oxygen species (ROS)- mediated endoplasmic reticulum stress- induced apoptosis in diabetic cardiomyopathy. <i>Cardiovascular Diabetology</i> , 2013, 12, 158.	6.8	169
38	Copy Number Variation on Chromosome 10q26.3 for Obesity Identified by a Genome-Wide Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2013, 98, E191-E195.	3.6	19
39	ASSA13-06-6â€¦Prevention of Cardiac Remodelling by Gene Silencing of Toll-Like Receptor-4 in Mice with Diabetic Cardiomyopathy. <i>Heart</i> , 2013, 99, A35.1-A35.	2.9	0
40	A novel replicated association between FXYP6 gene and schizophrenia. <i>Biochemical and Biophysical Research Communications</i> , 2011, 405, 118-121.	2.1	10
41	Genetics of osteoporotic fracture. <i>Orthopedic Research and Reviews</i> , 2011, Volume 3, 11-21.	1.1	3
42	Population-based and family-based association studies of ZNF804A locus and schizophrenia. <i>Molecular Psychiatry</i> , 2011, 16, 360-361.	7.9	58
43	Genome-Wide Association Study Identifies ALDH7A1 as a Novel Susceptibility Gene for Osteoporosis. <i>PLoS Genetics</i> , 2010, 6, e1000806.	3.5	101
44	The regulation-of-autophagy pathway may influence Chinese stature variation: evidence from elder adults. <i>Journal of Human Genetics</i> , 2010, 55, 441-447.	2.3	20
45	Is the EFNB2 locus associated with schizophrenia? Single nucleotide polymorphisms and haplotypes analysis. <i>Psychiatry Research</i> , 2010, 180, 5-9.	3.3	16
46	Genome-wide Copy-Number-Variation Study Identified a Susceptibility Gene, UGT2B17, for Osteoporosis. <i>American Journal of Human Genetics</i> , 2008, 83, 663-674.	6.2	209