

# Minxian Wang

## List of Publications by Year in descending order

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Version: 2024-02-01

33  
papers

1,570  
citations

471509

17  
h-index

414414

32  
g-index

39  
all docs

39  
docs citations

39  
times ranked

2723  
citing authors

#	ARTICLE	IF	CITATIONS
1	Rare coding variants in 35 genes associate with circulating lipid levelsâ€”A multi-ancestry analysis of 170,000 exomes. <i>American Journal of Human Genetics</i> , 2022, 109, 81-96.	6.2	24
2	Association of Habitual Alcohol Intake With Risk of Cardiovascular Disease. <i>JAMA Network Open</i> , 2022, 5, e223849.	5.9	136
3	Association of the Interaction Between Familial Hypercholesterolemia Variants and Adherence to a Healthy Lifestyle With Risk of Coronary Artery Disease. <i>JAMA Network Open</i> , 2022, 5, e222687.	5.9	17
4	DGAT2 Inhibition Potentiates Lipid Droplet Formation To Reduce Cytotoxicity in APOL1 Kidney Risk Variants. <i>Journal of the American Society of Nephrology: JASN</i> , 2022, 33, 889-907.	6.1	15
5	Association of Pathogenic DNA Variants Predisposing to Cardiomyopathy With Cardiovascular Disease Outcomes and All-Cause Mortality. <i>JAMA Cardiology</i> , 2022, 7, 723.	6.1	15
6	Inherited basis of visceral, abdominal subcutaneous and gluteofemoral fat depots. <i>Nature Communications</i> , 2022, 13, .	12.8	43
7	Lp(a) (Lipoprotein[a]) Concentrations and Incident Atherosclerotic Cardiovascular Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2021, 41, 465-474.	2.4	104
8	Genetic Predictor to Identify Individuals With High Lipoprotein(a) Concentrations. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003182.	3.6	10
9	De novo TRIM8 variants impair its protein localization to nuclear bodies and cause developmental delay, epilepsy, and focal segmental glomerulosclerosis. <i>American Journal of Human Genetics</i> , 2021, 108, 357-367.	6.2	14
10	Quantifying and Understanding the Higher Risk of Atherosclerotic Cardiovascular Disease Among South Asian Individuals. <i>Circulation</i> , 2021, 144, 410-422.	1.6	72
11	Rare, Damaging DNA Variants in <i>CORIN</i> and Risk of Coronary Artery Disease: Insights From Functional Genomics and Large-Scale Sequencing Analyses. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003399.	3.6	10
12	Polygenic basis and biomedical consequences of telomere length variation. <i>Nature Genetics</i> , 2021, 53, 1425-1433.	21.4	145
13	Machine learning enables new insights into genetic contributions to liver fat accumulation. <i>Cell Genomics</i> , 2021, 1, 100066.	6.5	34
14	Evaluating the Potential of T Cell Receptor Repertoires in Predicting the Prognosis of Resectable Non-Small Cell Lung Cancers. <i>Molecular Therapy - Methods and Clinical Development</i> , 2020, 18, 73-83.	4.1	24
15	Validation of a Genome-Wide Polygenic Score for Coronary Artery Disease in South Asians. <i>Journal of the American College of Cardiology</i> , 2020, 76, 703-714.	2.8	76
16	Polygenic background modifies penetrance of monogenic variants for tier 1 genomic conditions. <i>Nature Communications</i> , 2020, 11, 3635.	12.8	277
17	Analysis of cardiac magnetic resonance imaging in 36,000 individuals yields genetic insights into dilated cardiomyopathy. <i>Nature Communications</i> , 2020, 11, 2254.	12.8	140
18	Association of Rare Pathogenic DNA Variants for Familial Hypercholesterolemia, Hereditary Breast and Ovarian Cancer Syndrome, and Lynch Syndrome With Disease Risk in Adults According to Family History. <i>JAMA Network Open</i> , 2020, 3, e203959.	5.9	75

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19	Phosphorylation of ACTN4 Leads to Podocyte Vulnerability and Proteinuric Glomerulosclerosis. <i>Journal of the American Society of Nephrology: JASN</i> , 2020, 31, 1479-1495.	6.1	19
20	Genome-Wide Polygenic Score and Cardiovascular Outcomes With Evacetrapib in Patients With High-Risk Vascular Disease. <i>Circulation Genomic and Precision Medicine</i> , 2020, 13, e002767.	3.6	9
21	Autosomal Dominant Tubulointerstitial Kidney Disease—Uromodulin Misclassified as Focal Segmental Glomerulosclerosis or Hereditary Glomerular Disease. <i>Kidney International Reports</i> , 2020, 5, 519-529.	0.8	14
22	Contributions of Rare Gene Variants to Familial and Sporadic FSGS. <i>Journal of the American Society of Nephrology: JASN</i> , 2019, 30, 1625-1640.	6.1	42
23	Rare Genetic Variants Associated With Sudden Cardiac Death in Adults. <i>Journal of the American College of Cardiology</i> , 2019, 74, 2623-2634.	2.8	27
24	APOL1-Associated Kidney Disease in Brazil. <i>Kidney International Reports</i> , 2019, 4, 923-929.	0.8	24
25	Volanesorsen, Familial Chylomicronemia Syndrome, and Thrombocytopenia. <i>New England Journal of Medicine</i> , 2019, 381, 2582-2584.	27.0	21
26	Disease-causing mutation in $\alpha$ -actinin-4 promotes podocyte detachment through maladaptation to periodic stretch. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 1517-1522.	7.1	51
27	<i>UBD</i> modifies <i>APOL1</i> -induced kidney disease risk. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 3446-3451.	7.1	52
28	Composition and Interactions of Hepatitis B Virus Quasispecies Defined the Virological Response During Telbivudine Therapy. <i>Scientific Reports</i> , 2015, 5, 17123.	3.3	14
29	A probabilistic method for testing and estimating selection differences between populations. <i>Genome Research</i> , 2015, 25, 1903-1909.	5.5	10
30	Detecting Recent Positive Selection with High Accuracy and Reliability by Conditional Coalescent Tree. <i>Molecular Biology and Evolution</i> , 2014, 31, 3068-3080.	8.9	20
31	Resolving ambiguity in the phylogenetic relationship of genotypes A, B, and C of hepatitis B virus. <i>BMC Evolutionary Biology</i> , 2013, 13, 120.	3.2	2
32	A Monte Carlo Permutation Test for Random Mating Using Genome Sequences. <i>PLoS ONE</i> , 2013, 8, e71496.	2.5	1
33	South Asian Patient Population Genetics Reveal Strong Founder Effects and High Rates of Homozygosity — New Resources for Precision Medicine. <i>SSRN Electronic Journal</i> , 0, , .	0.4	2