

# Qian Wang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5084076/publications.pdf>

Version: 2024-02-01

10  
papers

789  
citations

933447

10  
h-index

1372567

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

2243  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. <i>JAMA Psychiatry</i> , 2017, 74, 1234.	11.0	74
2	Genetic factor common to schizophrenia and HIV infection is associated with risky sexual behavior: antagonistic vs. synergistic pleiotropic SNPs enriched for distinctly different biological functions. <i>Human Genetics</i> , 2017, 136, 75-83.	3.8	17
3	The Interplay Between Risky Sexual Behaviors and Alcohol Dependence: Genome-Wide Association and Neuroimaging Support for LHPP as a Risk Gene. <i>Neuropsychopharmacology</i> , 2017, 42, 598-605.	5.4	40
4	Systematic tissue-specific functional annotation of the human genome highlights immune-related DNA elements for late-onset Alzheimer's disease. <i>PLoS Genetics</i> , 2017, 13, e1006933.	3.5	96
5	Integrative Tissue-Specific Functional Annotations in the Human Genome Provide Novel Insights on Many Complex Traits and Improve Signal Prioritization in Genome Wide Association Studies. <i>PLoS Genetics</i> , 2016, 12, e1005947.	3.5	94
6	Genome-wide Association Study of Cannabis Dependence Severity, Novel Risk Variants, and Shared Genetic Risks. <i>JAMA Psychiatry</i> , 2016, 73, 472.	11.0	148
7	Genome-wide Association Studies of Posttraumatic Stress Disorder in 2 Cohorts of US Army Soldiers. <i>JAMA Psychiatry</i> , 2016, 73, 695.	11.0	158
8	Implications of pleiotropy: challenges and opportunities for mining Big Data in biomedicine. <i>Frontiers in Genetics</i> , 2015, 6, 229.	2.3	41
9	A review of study designs and statistical methods for genomic epidemiology studies using next generation sequencing. <i>Frontiers in Genetics</i> , 2015, 6, 149.	2.3	48
10	Pervasive pleiotropy between psychiatric disorders and immune disorders revealed by integrative analysis of multiple GWAS. <i>Human Genetics</i> , 2015, 134, 1195-1209.	3.8	72