Qian Wang

List of Publications by Year
in descending order

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	Genome-wide Association Studies of Posttraumatic Stress Disorder in 2 Cohorts of US Army Soldiers. JAMA Psychiatry, 2016, 73, 695.	11.0	158
2	Genome-wide Association Study of Cannabis Dependence Severity, Novel Risk Variants, and Shared Genetic Risks. JAMA Psychiatry, 2016, 73, 472.	11.0	148
3	Systematic tissue-specific functional annotation of the human genome highlights immune-related DNA elements for late-onset Alzheimer's disease. PLoS Genetics, 2017, 13, e1006933.	3.5	96
4	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. PLoS Genetics, 2016, 12, e1005947.	3.5	94
5	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. JAMA Psychiatry, 2017, 74, 1234.	11.0	74
6	Pervasive pleiotropy between psychiatric disorders and immune disorders revealed by integrative analysis of multiple GWAS. Human Genetics, 2015, 134, 1195-1209.	3.8	72
7	A review of study designs and statistical methods for genomic epidemiology studies using next generation sequencing. Frontiers in Genetics, 2015, 6, 149.	2.3	48
8	Implications of pleiotropy: challenges and opportunities for mining Big Data in biomedicine. Frontiers in Genetics, 2015, 6, 229.	2.3	41
9	The Interplay Between Risky Sexual Behaviors and Alcohol Dependence: Genome-Wide Association and Neuroimaging Support for LHPP as a Risk Gene. Neuropsychopharmacology, 2017, 42, 598-605.	5.4	40
10	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. Human Genetics, 2017, 136, 75-83.	3.8	17