

# Wei Q Deng

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10596344/publications.pdf>

Version: 2024-02-01

10  
papers

659  
citations

1478505

6  
h-index

1588992

8  
g-index

13  
all docs

13  
docs citations

13  
times ranked

1198  
citing authors

#	ARTICLE	IF	CITATIONS
1	gJLS2: an R package for generalized joint location and scale analysis in X-inclusive genome-wide association studies. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	0
2	Analytical strategies to include the X-chromosome in variance heterogeneity analyses: Evidence for trait-specific polygenic variance structure. <i>Genetic Epidemiology</i> , 2019, 43, 815-830.	1.3	14
3	A robust method to estimate regional polygenic correlation under misspecified linkage disequilibrium structure. <i>Genetic Epidemiology</i> , 2018, 42, 636-647.	1.3	3
4	Ranking and characterization of established BMI and lipid associated loci as candidates for gene-environment interactions. <i>PLoS Genetics</i> , 2017, 13, e1006812.	3.5	24
5	A method to estimate the contribution of regional genetic associations to complex traits from summary association statistics. <i>Scientific Reports</i> , 2016, 6, 27644.	3.3	5
6	Response to Letter Regarding Article "Temporal Relationship Between Subclinical Atrial Fibrillation and Embolic Events". <i>Circulation</i> , 2015, 131, e337-8.	1.6	0
7	Contribution of Large Region Joint Associations to Complex Traits Genetics. <i>PLoS Genetics</i> , 2015, 11, e1005103.	3.5	10
8	Meta-analysis of SNPs involved in variance heterogeneity using Levene's test for equal variances. <i>European Journal of Human Genetics</i> , 2014, 22, 427-430.	2.8	7
9	Temporal Relationship Between Subclinical Atrial Fibrillation and Embolic Events. <i>Circulation</i> , 2014, 129, 2094-2099.	1.6	579
10	A fast algorithm to optimize SNP prioritization for gene-gene and gene-environment interactions. <i>Genetic Epidemiology</i> , 2011, 35, 729-738.	1.3	17