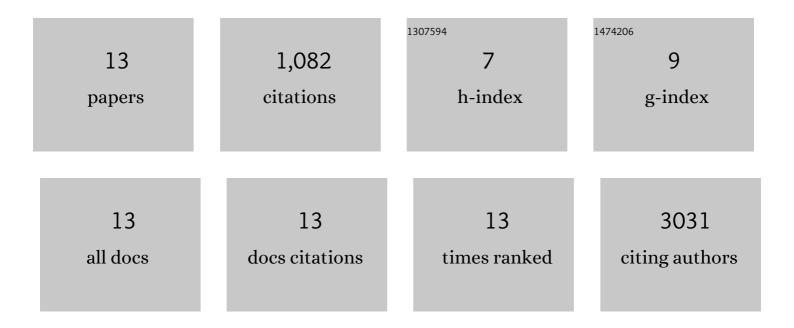
Wei Cheng

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/218315/publications.pdf Version: 2024-02-01



WELCHENC

#	Article	IF	CITATIONS
1	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. Nature Genetics, 2017, 49, 27-35.	21.4	838
2	Flexible and robust co-regularized multi-domain graph clustering. , 2013, , .		48
3	Co-Regularized Deep Multi-Network Embedding. , 2018, , .		35
4	Scaling Up Kernel SVM on Limited Resources: A Low-Rank Linearization Approach. IEEE Transactions on Neural Networks and Learning Systems, 2019, 30, 369-378.	11.3	35
5	Graph-regularized dual Lasso for robust eQTL mapping. Bioinformatics, 2014, 30, i139-i148.	4.1	33
6	Searching Dimension Incomplete Databases. IEEE Transactions on Knowledge and Data Engineering, 2014, 26, 725-738.	5.7	25
7	Robust Multi-Network Clustering via Joint Cross-Domain Cluster Alignment. , 2015, 2015, 291-300.		19
8	ComClus: A Self-Grouping Framework for Multi-Network Clustering. IEEE Transactions on Knowledge and Data Engineering, 2018, 30, 435-448.	5.7	14
9	HICC: an entropy splitting-based framework for hierarchical co-clustering. Knowledge and Information Systems, 2016, 46, 343-367.	3.2	12
10	Fast and robust group-wise eQTL mapping using sparse graphical models. BMC Bioinformatics, 2015, 16, 2.	2.6	11
11	<inline-formula> <tex-math notation="LaTeX">\$mathcal{DBSDA}\$ </tex-math> </inline-formula> : Lowering the Bound of Misclassification Rate for Sparse Linear Discriminant Analysis via Model Debiasing. IEEE Transactions on Neural Networks and Learning Systems, 2019, 30, 707-717.	11.3	7
12	Sparse regression models for unraveling group and individual associations in eQTL mapping. BMC Bioinformatics, 2016, 17, 136.	2.6	4
13	Measuring Opinion Relevance in Latent Topic Space. , 2011, , .		1