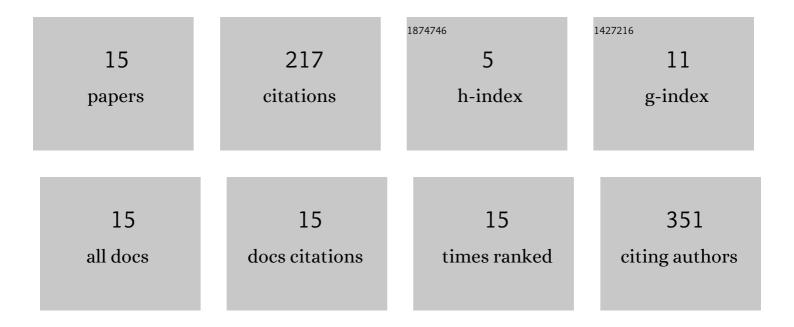
Nima S Hejazi

List of Publications by Year in descending order

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NIMA S HEIAZI

#	Article	IF	CITATIONS
1	Nonparametric causal mediation analysis for stochastic interventional (in)direct effects. Biostatistics, 2023, 24, 686-707.	0.9	3
2	Biodigester Cookstove Interventions and Child Diarrhea in Semirural Nepal: A Causal Analysis of Daily Observations. Environmental Health Perspectives, 2022, 130, 17002.	2.8	5
3	medoutcon: Nonparametric efficient causal mediation analysis with machine learning in R. Journal of Open Source Software, 2022, 7, 3979.	2.0	2
4	Evaluating the robustness of targeted maximum likelihood estimators via realistic simulations in nutrition intervention trials. Statistics in Medicine, 2022, 41, 2132-2165.	0.8	2
5	Efficient nonparametric inference on the effects of stochastic interventions under twoâ€phase sampling, with applications to vaccine efficacy trials. Biometrics, 2021, 77, 1241-1253.	0.8	15
6	Vaccine Efficacy of ALVAC-HIV and Bivalent Subtype C gp120–MF59 in Adults. New England Journal of Medicine, 2021, 384, 1089-1100.	13.9	144
7	cvCovEst: Cross-validated covariance matrix estimator selection and evaluation in R. Journal of Open Source Software, 2021, 6, 3273.	2.0	3
8	Exploring high-dimensional biological data with sparse contrastive principal component analysis. Bioinformatics, 2020, 36, 3422-3430.	1.8	20
9	hal9001: Scalable highly adaptive lasso regression in R. Journal of Open Source Software, 2020, 5, 2526.	2.0	9
10	scPCA: A toolbox for sparse contrastive principal component analysis in R. Journal of Open Source Software, 2020, 5, 2079.	2.0	0
11	txshift: Efficient estimation of the causal effects of stochastic interventions in R. Journal of Open Source Software, 2020, 5, 2447.	2.0	6
12	origami: A Generalized Framework for Cross-Validation in R. Journal of Open Source Software, 2018, 3, 512.	2.0	6
13	adaptest: Data-Adaptive Statistics for High-Dimensional Testing in R. Journal of Open Source Software, 2018, 3, 161.	2.0	0
14	biotmle: Targeted Learning for Biomarker Discovery. Journal of Open Source Software, 2017, 2, 295.	2.0	1
15	methyvim: Targeted, robust, and model-free differential methylation analysis in R. F1000Research, 0, 7, 1424.	0.8	1