

# Nima S Hejazi

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5806633/publications.pdf>

Version: 2024-02-01

15  
papers

217  
citations

1874746  
5  
h-index

1427216  
11  
g-index

15  
all docs

15  
docs citations

15  
times ranked

351  
citing authors

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