Forrest W Crawford

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Predicting daily COVID-19 case rates from SARS-CoV-2 RNA concentrations across a diversity of wastewater catchments. FEMS Microbes, 2022, 2, xtab022.	0.8	19
2	Impact of close interpersonal contact on COVID-19 incidence: Evidence from 1 year of mobile device data. Science Advances, 2022, 8, eabi5499.	4.7	19
3	Vaccination with BNT162b2 reduces transmission of SARS-CoV-2 to household contacts in Israel. Science, 2022, 375, 1151-1154.	6.0	109
4	SARS-CoV-2 Attack Rate and Population Immunity in Southern New England, March 2020 to May 2021. JAMA Network Open, 2022, 5, e2214171.	2.8	5
5	Violence and Discrimination Against Men Who Have Sex With Men in Lebanon: The Role of International Displacement and Migration. Journal of Interpersonal Violence, 2021, 36, 10267-10284.	1.3	8
6	Repeat SARS-CoV-2 testing models for residential college populations. Health Care Management Science, 2021, 24, 305-318.	1.5	29
7	Estimates of people who injected drugs within the last 12 months in Belgium based on a capture-recapture and multiplier method. Drug and Alcohol Dependence, 2021, 219, 108436.	1.6	2
8	Identification of causal intervention effects under contagion. Journal of Causal Inference, 2021, 9, 9-38.	0.5	4
9	One year of modeling and forecasting COVID-19 transmission to support policymakers in Connecticut. Scientific Reports, 2021, 11, 20271.	1.6	18
10	COVID-19 Testing and Case Rates and Social Contact Among Residential College Students in Connecticut During the 2020-2021 Academic Year. JAMA Network Open, 2021, 4, e2140602.	2.8	22
11	Costâ€effectiveness of expanding the capacity of opioid agonist treatment in Ukraine: dynamic modeling analysis. Addiction, 2020, 115, 437-450.	1.7	15
12	Transmission Modeling with Regression Adjustment for Analyzing Household-based Studies of Infectious Disease. Epidemiology, 2020, 31, 238-247.	1.2	1
13	Using sigLASSO to optimize cancer mutation signatures jointly with sampling likelihood. Nature Communications, 2020, 11, 3575.	5.8	28
14	Collective communication and behaviour in response to uncertain â€~Danger' in network experiments. Proceedings of the Royal Society A: Mathematical, Physical and Engineering Sciences, 2020, 476, 20190685.	1.0	6
15	Creating a Framework for Conducting Randomized Clinical Trials during Disease Outbreaks. New England Journal of Medicine, 2020, 382, 1366-1369.	13.9	63
16	Estimation of Excess Deaths Associated With the COVID-19 Pandemic in the United States, March to May 2020. JAMA Internal Medicine, 2020, 180, 1336.	2.6	374
17	Divergent estimates of HIV incidence among people who inject drugs in Ukraine. International Journal of Drug Policy, 2019, 73, 156-162.	1.6	2
18	Interpretation of the Individual Effect Under Treatment Spillover. American Journal of Epidemiology, 2019, 188, 1407-1409.	1.6	6

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19	Empirical evidence of recruitment bias in a network study of people who inject drugs. American Journal of Drug and Alcohol Abuse, 2019, 45, 460-469.	1.1	2
20	Suspected heroin-related overdoses incidentsÂin Cincinnati, Ohio: AÂspatiotemporal analysis. PLoS Medicine, 2019, 16, e1002956.	3.9	17
21	An Adaptive Approach to Locating Mobile HIV Testing Services. Medical Decision Making, 2018, 38, 262-272.	1.2	6
22	Risk ratios for contagious outcomes. Journal of the Royal Society Interface, 2018, 15, 20170696.	1.5	16
23	Estimating dose-specific cell division and apoptosis rates from chemo-sensitivity experiments. Scientific Reports, 2018, 8, 2705.	1.6	5
24	Computational methods for birthâ€death processes. Wiley Interdisciplinary Reviews: Computational Statistics, 2018, 10, e1423.	2.1	12
25	Exposure, hazard, and survival analysis of diffusion on social networks. Statistics in Medicine, 2018, 37, 2561-2585.	0.8	9
26	Hidden Population Size Estimation From Respondent-Driven Sampling: A Network Approach. Journal of the American Statistical Association, 2018, 113, 755-766.	1.8	41
27	Identification of Homophily and Preferential Recruitment in Respondent-Driven Sampling. American Journal of Epidemiology, 2018, 187, 153-160.	1.6	29
28	Birth/birth-death processes and their computable transition probabilities with biological applications. Journal of Mathematical Biology, 2018, 76, 911-944.	0.8	28
29	HIV Testing and awareness of HIV status among people who inject drugs in greater Kuala Lumpur, Malaysia AIDS Care - Psychological and Socio-Medical Aspects of AIDS/HIV, 2018, 30, 59-64.	0.6	9
30	Dynamics of the HIV outbreak and response in Scott County, IN, USA, 2011–15: a modelling study. Lancet HIV,the, 2018, 5, e569-e577.	2.1	84
31	Direct likelihood-based inference for discretely observed stochastic compartmental models of infectious disease. Annals of Applied Statistics, 2018, 12, .	0.5	8
32	Computer-assisted detection of tardus parvus waveforms on Doppler ultrasound. Ultrasound, 2018, 26, 81-92.	0.3	3
33	HIV Risk, Prevalence, and Access to Care Among Men Who Have Sex with Men in Lebanon. AIDS Research and Human Retroviruses, 2017, 33, 1149-1154.	0.5	23
34	Using data from respondent-driven sampling studies to estimate the number of people who inject drugs: Application to the Kohtla-J¤ve region of Estonia. PLoS ONE, 2017, 12, e0185711.	1.1	4
35	Predictors of mortality within prison and after release among persons living with HIV in Indonesia. Research and Reports in Tropical Medicine, 2017, Volume 8, 25-35.	2.8	10
36	The burden of typhoid fever in low- and middle-income countries: A meta-regression approach. PLoS Neglected Tropical Diseases, 2017, 11, e0005376.	1.3	212

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37	The Graphical Structure of Respondent-driven Sampling. Sociological Methodology, 2016, 46, 187-211.	1.4	17
38	Coupling bounds for approximating birth–death processes by truncation. Statistics and Probability Letters, 2016, 109, 30-38.	0.4	4
39	Sex, lies and self-reported counts: Bayesian mixture models for heaping in longitudinal count data via birth–death processes. Annals of Applied Statistics, 2015, 9, 572-596.	0.5	20
40	Did Large-Scale Vaccination Drive Changes in the Circulating Rotavirus Population in Belgium?. Scientific Reports, 2015, 5, 18585.	1.6	36
41	Nonparametric identification for respondent-driven sampling. Statistics and Probability Letters, 2015, 106, 100-102.	0.4	7
42	Markov counting models for correlated binary responses. Biostatistics, 2015, 16, 427-440.	0.9	1
43	Combining List Experiment and Direct Question Estimates of Sensitive Behavior Prevalence. Journal of Survey Statistics and Methodology, 2015, 3, 43-66.	0.5	54
44	Obesity not associated with severity among hospitalized adults with seasonal influenza virus infection. Infection, 2015, 43, 569-575.	2.3	35
45	Historical Biogeography Using Species Geographical Ranges. Systematic Biology, 2015, 64, 1059-1073.	2.7	46
46	HIV Prevalence Among People Who Inject Drugs in Greater Kuala Lumpur Recruited Using Respondent-Driven Sampling. AIDS and Behavior, 2015, 19, 2347-2357.	1.4	18
47	On the distribution of interspecies correlation for Markov models of character evolution on Yule trees. Journal of Theoretical Biology, 2015, 364, 275-283.	0.8	10
48	Estimation for General Birth-Death Processes. Journal of the American Statistical Association, 2014, 109, 730-747.	1.8	32
49	Diversity, Disparity, and Evolutionary Rate Estimation for Unresolved Yule Trees. Systematic Biology, 2013, 62, 439-455.	2.7	17
50	Transition probabilities for general birth–death processes with applications in ecology, genetics, and evolution. Journal of Mathematical Biology, 2012, 65, 553-580.	0.8	60