

Samuel Soubeyrand

List of Publications by Year in descending order

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Version: 2024-02-01

54
papers

1,047
citations

516215

16
h-index

525886

27
g-index

62
all docs

62
docs citations

62
times ranked

1406
citing authors

#	ARTICLE	IF	CITATIONS
1	One Health concepts and challenges for surveillance, forecasting, and mitigation of plant disease beyond the traditional scope of crop production. <i>Plant Pathology</i> , 2022, 71, 86-97.	1.2	18
2	Strain diversity and spatial distribution are linked to epidemic dynamics in host populations. <i>American Naturalist</i> , 2022, 199, 59-74.	1.0	8
3	Spatial statistics and stochastic partial differential equations: A mechanistic viewpoint. <i>Spatial Statistics</i> , 2022, , 100591.	0.9	2
4	When the average hides the risk of Bt-corn pollen on non-target Lepidoptera: Application to Aglais io in Catalonia. <i>Ecotoxicology and Environmental Safety</i> , 2021, 207, 111215.	2.9	11
5	Equilibrium and sensitivity analysis of a spatio-temporal host-vector epidemic model. <i>Nonlinear Analysis: Real World Applications</i> , 2021, 57, 103194.	0.9	3
6	Extension of the spatially and temporally explicit "brisk" model to assess potential adverse effects of Bt-maize pollen on non-target Lepidoptera at landscape level. <i>EFSA Supporting Publications</i> , 2021, 18, 6443E.	0.3	2
7	Inferring long-distance connectivity shaped by air-mass movement for improved experimental design in aerobiology. <i>Scientific Reports</i> , 2021, 11, 11093.	1.6	12
8	COVID-19 mortality dynamics: The future modelled as a (mixture of) past(s). <i>PLoS ONE</i> , 2020, 15, e0238410.	1.1	8
9	Towards unified and real-time analyses of outbreaks at country-level during pandemics. <i>One Health</i> , 2020, 11, 100187.	1.5	11
10	Impact of Lockdown on the Epidemic Dynamics of COVID-19 in France. <i>Frontiers in Medicine</i> , 2020, 7, 274.	1.2	52
11	Using Early Data to Estimate the Actual Infection Fatality Ratio from COVID-19 in France. <i>Biology</i> , 2020, 9, 97.	1.3	84
12	A parsimonious approach for spatial transmission and heterogeneity in the COVID-19 propagation. <i>Royal Society Open Science</i> , 2020, 7, 201382.	1.1	23
13	Analyzing the Influence of Landscape Aggregation on Disease Spread to Improve Management Strategies. <i>Phytopathology</i> , 2019, 109, 1198-1207.	1.1	5
14	Quick inference for log Gaussian Cox processes with non-stationary underlying random fields. <i>Spatial Statistics</i> , 2019, 33, 100388.	0.9	0
15	Dating and localizing an invasion from post-introduction data and a coupled reaction-diffusion-absorption model. <i>Journal of Mathematical Biology</i> , 2019, 79, 765-789.	0.8	24
16	Inferring epidemiological links from deep sequencing data: a statistical learning approach for human, animal and plant diseases. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20180258.	1.8	14
17	Improving Management Strategies of Plant Diseases Using Sequential Sensitivity Analyses. <i>Phytopathology</i> , 2019, 109, 1184-1197.	1.1	17
18	Identifying Lookouts for Epidemio-Surveillance: Application to the Emergence of <i>Xylella fastidiosa</i> in France. <i>Phytopathology</i> , 2019, 109, 265-276.	1.1	37

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19	A Spatio-temporal Exposure-hazard Model for Assessing Biological Risk and Impact. <i>Risk Analysis</i> , 2019, 39, 54-70.	1.5	11
20	Using sensitivity analysis to identify key factors for the propagation of a plant epidemic. <i>Royal Society Open Science</i> , 2018, 5, 171435.	1.1	18
21	Spatial exposure-hazard and landscape models for assessing the impact of GM crops on non-target organisms. <i>Science of the Total Environment</i> , 2018, 624, 470-479.	3.9	10
22	Inferring pathogen dynamics from temporal count data: the emergence of <i>Xylella fastidiosa</i> in France is probably not recent. <i>New Phytologist</i> , 2018, 219, 824-836.	3.5	63
23	Assessing the Aerial Interconnectivity of Distant Reservoirs of <i>Sclerotinia sclerotiorum</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 2257.	1.5	14
24	Mapping Rainfall Feedback to Reveal the Potential Sensitivity of Precipitation to Biological Aerosols. <i>Bulletin of the American Meteorological Society</i> , 2017, 98, 1109-1118.	1.7	26
25	On parameter estimation for doubly inhomogeneous cluster point processes. <i>Spatial Statistics</i> , 2017, 20, 191-205.	0.9	4
26	Exploiting Genetic Information to Trace Plant Virus Dispersal in Landscapes. <i>Annual Review of Phytopathology</i> , 2017, 55, 139-160.	3.5	19
27	<scp>PESO</scp>; a modelling framework to help improve management strategies for epidemics – application to sharka. <i>EPPO Bulletin</i> , 2017, 47, 231-236.	0.6	2
28	Testing Differences Between Pathogen Compositions with Small Samples and Sparse Data. <i>Phytopathology</i> , 2017, 107, 1199-1208.	1.1	7
29	When Group Dispersal and Allee Effect Shape Metapopulation Dynamics. <i>Annales Zoologici Fennici</i> , 2017, 54, 123-138.	0.2	2
30	Monte Carlo testing in spatial statistics, with applications to spatial residuals. <i>Spatial Statistics</i> , 2016, 18, 40-53.	0.9	6
31	Rearranging agricultural landscapes towards habitat quality optimisation: In silico application to pest regulation. <i>Ecological Complexity</i> , 2016, 28, 113-122.	1.4	13
32	Assessing the Mismatch Between Incubation and Latent Periods for Vector-Borne Diseases: The Case of Sharka. <i>Phytopathology</i> , 2015, 105, 1408-1416.	1.1	20
33	Sharka Epidemiology and Worldwide Management Strategies: Learning Lessons to Optimize Disease Control in Perennial Plants. <i>Annual Review of Phytopathology</i> , 2015, 53, 357-378.	3.5	76
34	Evolution of dispersal in asexual populations: to be independent, clumped or grouped?. <i>Evolutionary Ecology</i> , 2015, 29, 947-963.	0.5	11
35	Weak convergence of posteriors conditional on maximum pseudo-likelihood estimates and implications in ABC. <i>Statistics and Probability Letters</i> , 2015, 107, 84-92.	0.4	7
36	Regression-Based Ranking of Pathogen Strains with Respect to Their Contribution to Natural Epidemics. <i>PLoS ONE</i> , 2014, 9, e86591.	1.1	4

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37	Long-Distance Wind-Dispersal of Spores in a Fungal Plant Pathogen: Estimation of Anisotropic Dispersal Kernels from an Extensive Field Experiment. <i>PLoS ONE</i> , 2014, 9, e103225.	1.1	94
38	Parameter estimation for reaction-diffusion models of biological invasions. <i>Population Ecology</i> , 2014, 56, 427-434.	0.7	33
39	A Nonstationary Cylinder-Based Model Describing Group Dispersal in a Fragmented Habitat. <i>Stochastic Models</i> , 2014, 30, 48-67.	0.3	5
40	Analysis of fragmented time directionality in time series to elucidate feedbacks in climate data. <i>Environmental Modelling and Software</i> , 2014, 61, 78-86.	1.9	10
41	A Bayesian approach for inferring the dynamics of partially observed endemic infectious diseases from space-time-genetic data. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20133251.	1.2	76
42	Approximate Bayesian computation with functional statistics. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2013, 12, 17-37.	0.2	8
43	Goodness-of-fit test of the mark distribution in a point process with non-stationary marks. <i>Statistics and Computing</i> , 2012, 22, 931-943.	0.8	0
44	A statistical-reaction-diffusion approach for analyzing expansion processes. <i>Journal of Theoretical Biology</i> , 2011, 274, 43-51.	0.8	30
45	Patchy patterns due to group dispersal. <i>Journal of Theoretical Biology</i> , 2011, 271, 87-99.	0.8	16
46	Aggregation patterns in hierarchy/proximity spaces. <i>Ecological Complexity</i> , 2010, 7, 21-31.	1.4	1
47	Exploring Spatial and Multitype Assemblages of Species Abundances. <i>Biometrical Journal</i> , 2009, 51, 979-995.	0.6	3
48	Autoinfection in wheat leaf rust epidemics. <i>New Phytologist</i> , 2008, 177, 1001-1011.	3.5	28
49	Modelling the spread in space and time of an airborne plant disease. <i>Journal of the Royal Statistical Society Series C: Applied Statistics</i> , 2008, 57, 253-272.	0.5	21
50	Accounting for roughness of circular processes: Using Gaussian random processes to model the anisotropic spread of airborne plant disease. <i>Theoretical Population Biology</i> , 2008, 73, 92-103.	0.5	27
51	Model-based estimation of the link between the daily survival probability and a time-varying covariate, application to mosquitofish survival data. <i>Mathematical Biosciences</i> , 2007, 210, 508-522.	0.9	4
52	Residual-based specification of a hidden random field included in a hierarchical spatial model. <i>Computational Statistics and Data Analysis</i> , 2007, 51, 6404-6422.	0.7	3
53	A Frailty Model to Assess Plant Disease Spread from Individual Count Data. <i>Journal of Data Science</i> , 2007, 5, 67-83.	0.5	9
54	Residual-based specification of the random-effects distribution for cluster data. <i>Statistical Methodology</i> , 2006, 3, 464-482.	0.5	4