

David Savage

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3166653/publications.pdf>

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12
papers

580
citations

759233

12
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

718
citing authors

#	ARTICLE	IF	CITATIONS
1	Requirements, design and implementation of a general model of biological invasion. <i>Ecological Modelling</i> , 2014, 272, 394-409.	2.5	16
2	Orientation and speed of wind gusts causing abscission of wind-dispersed seeds influences dispersal distance. <i>Functional Ecology</i> , 2014, 28, 973-981.	3.6	22
3	Anomaly detection in online social networks. <i>Social Networks</i> , 2014, 39, 62-70.	2.1	198
4	Temporal Patterns of Ascospore Release in <i>Leptosphaeria maculans</i> Vary Depending on Geographic Region and Time of Observation. <i>Microbial Ecology</i> , 2013, 65, 584-592.	2.8	20
5	Mobile traps are better than stationary traps for surveillance of airborne fungal spores. <i>Crop Protection</i> , 2012, 36, 23-30.	2.1	22
6	Seasonal and Diurnal Patterns of Spore Release Can Significantly Affect the Proportion of Spores Expected to Undergo Long-Distance Dispersal. <i>Microbial Ecology</i> , 2012, 63, 578-585.	2.8	45
7	Can mechanistically parameterised, anisotropic dispersal kernels provide a reliable estimate of wind-assisted dispersal?. <i>Ecological Modelling</i> , 2011, 222, 1673-1682.	2.5	18
8	Timing of propagule release significantly alters the deposition area of resulting aerial dispersal. <i>Diversity and Distributions</i> , 2010, 16, 288-299.	4.1	33
9	A neurotropic herpesvirus infecting the gastropod, abalone, shares ancestry with oyster herpesvirus and a herpesvirus associated with the amphioxus genome. <i>Virology Journal</i> , 2010, 7, 308.	3.4	63
10	SSRPrimer and SSR Taxonomy Tree: Biome SSR discovery. <i>Nucleic Acids Research</i> , 2006, 34, W656-W659.	14.5	62
11	SNPServer: a real-time SNP discovery tool. <i>Nucleic Acids Research</i> , 2005, 33, W493-W495.	14.5	56
12	New Computational Tools for Brassica Genome Research. <i>Comparative and Functional Genomics</i> , 2004, 5, 276-280.	2.0	25