

Stefan C Kremer

List of Publications by Year in Descending Order

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Version: 2024-04-28

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

31
papers

523
citations

12
h-index

22
g-index

39
ext. papers

726
ext. citations

3.8
avg. IF

4.33
L-index

#	Paper	IF	Citations
31	Bulk arthropod abundance, biomass and diversity estimation using deep learning for computer vision. <i>Methods in Ecology and Evolution</i> , 2022 , 13, 346-357	7.7	0
30	Long-term TE persistence even without beneficial insertion. <i>BMC Genomics</i> , 2021 , 22, 260	4.5	
29	Transposable element persistence via potential genome-level ecosystem engineering. <i>BMC Genomics</i> , 2020 , 21, 367	4.5	6
28	2020 ,		14
27	Three critical factors affecting automated image species recognition performance for camera traps. <i>Ecology and Evolution</i> , 2020 , 10, 3503-3517	2.8	26
26	Past, present and future approaches using computer vision for animal re-identification from camera trap data. <i>Methods in Ecology and Evolution</i> , 2019 , 10, 461-470	7.7	61
25	Network intrusion detection system based on recursive feature addition and bigram technique. <i>Computers and Security</i> , 2018 , 73, 137-155	4.9	53
24	Yes! There are resilient generalizations (or "laws") in ecology. <i>Quarterly Review of Biology</i> , 2016 , 91, 119-34	3.4	12
23	Prediction of Protein Coding Regions Using a Wide-Range Wavelet Window Method. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2016 , 13, 742-53	3	20
22	Applying ecological models to communities of genetic elements: the case of neutral theory. <i>Molecular Ecology</i> , 2015 , 24, 3232-42	5.7	23
21	A new Canadian interdisciplinary Ph.D. in computational sciences. <i>Journal of Computational Science</i> , 2015 , 9, 82-87	3.4	2
20	A dynamic representation-based, de novo method for protein-coding region prediction and biological information detection 2015 , 46, 10-18		6
19	A survey of QoS/QoE mechanisms in heterogeneous wireless networks. <i>Physical Communication</i> , 2014 , 13, 61-72	2.2	39
18	An Accurate, Fast Embedded Feature Selection for SVMs 2014 ,		6
17	A novel application of ecological analyses to assess transposable element distributions in the genome of the domestic cow, <i>Bos taurus</i> . <i>Genome</i> , 2013 , 56, 521-33	2.4	4
16	Distinguishing ecological from evolutionary approaches to transposable elements. <i>Biological Reviews</i> , 2013 , 88, 573-84	13.5	16
15	Protein secondary structure prediction using support vector machines and a codon encoding scheme 2012 ,		2

14	Gene prediction based on DNA spectral analysis: a literature review. <i>Journal of Computational Biology</i> , 2011 , 18, 639-76	1.7	46
13	Protein coding region prediction based on the adaptive representation method 2011 ,		3
12	Amino acid encoding schemes for machine learning methods 2011 ,		7
11	Neural grammar networks for toxicology 2010 ,		1
10	Theoretical justification of computing the 3-base periodicity using nucleotide distribution variance. <i>BioSystems</i> , 2010 , 101, 185-6	1.9	6
9	Neural Grammar Networks in QSAR Chemistry 2009 ,		2
8	A new distance distribution paradigm to detect the variability of the influenza-A virus in high dimensional spaces 2009 ,		1
7	Neural Grammar Networks. <i>Studies in Computational Intelligence</i> , 2009 , 67-96	0.8	1
6	New directions in fuzzy automata. <i>International Journal of Approximate Reasoning</i> , 2005 , 38, 175-214	3.6	66
5	A taxonomy for spatiotemporal connectionist networks revisited: the unsupervised case. <i>Neural Computation</i> , 2003 , 15, 1255-320	2.9	55
4	Spatiotemporal Connectionist Networks: A Taxonomy and Review. <i>Neural Computation</i> , 2001 , 13, 249-306	0.9	43
3	Cell Boundary Detection and Volume Approximation of Confocal Microscope Images for Bioinformatics. <i>Microscopy and Microanalysis</i> , 2000 , 6, 816-817	0.5	
2	Genomic Environments and Their Influence on Transposable Element Communities		1
1	Similarity learning networks for animal individual re-identification: an ecological perspective. <i>Mammalian Biology</i> , 1	1.6	0