

Yiqing Xu

List of Publications by Citations

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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.

32
papers

511
citations

14
h-index

21
g-index

39
ext. papers

720
ext. citations

3.1
avg, IF

4.47
L-index

#	Paper	IF	Citations
32	Multi-Stage Feature Constraints Learning for Age Estimation. <i>IEEE Transactions on Information Forensics and Security</i> , 2020 , 15, 2417-2428	8	53
31	Least squares twin bounded support vector machines based on L1-norm distance metric for classification. <i>Pattern Recognition</i> , 2018 , 74, 434-447	7.7	52
30	Non-intrusive load disaggregation based on deep dilated residual network. <i>Electric Power Systems Research</i> , 2019 , 170, 277-285	3.5	50
29	Organellar genome assembly methods and comparative analysis of horticultural plants. <i>Horticulture Research</i> , 2018 , 5, 3	7.7	44
28	Analysis of the Complete Mitochondrial Genome Sequence of the Diploid Cotton by Comparative Genomics Approaches. <i>BioMed Research International</i> , 2016 , 2016, 5040598	3	25
27	Genome-wide identification and characterization of WRKY gene family in <i>Salix suchowensis</i> . <i>PeerJ</i> , 2016 , 4, e2437	3.1	25
26	Water Areas Segmentation from Remote Sensing Images Using a Separable Residual SegNet Network. <i>ISPRS International Journal of Geo-Information</i> , 2020 , 9, 256	2.9	24
25	Cloud/shadow segmentation based on global attention feature fusion residual network for remote sensing imagery. <i>International Journal of Remote Sensing</i> , 2021 , 42, 2022-2045	3.1	24
24	DAU-Net: a novel water areas segmentation structure for remote sensing image. <i>International Journal of Remote Sensing</i> , 2021 , 42, 2594-2621	3.1	24
23	Assembly and comparative analysis of complete mitochondrial genome sequence of an economic plant. <i>PeerJ</i> , 2017 , 5, e3148	3.1	23
22	Dilated residual attention network for load disaggregation. <i>Neural Computing and Applications</i> , 2019 , 31, 8931-8953	4.8	17
21	River segmentation based on separable attention residual network. <i>Journal of Applied Remote Sensing</i> , 2019 , 14, 1	1.4	17
20	L1-Norm GEPSVM Classifier Based on an Effective Iterative Algorithm for Classification. <i>Neural Processing Letters</i> , 2018 , 48, 273-298	2.4	17
19	Dilated multi-scale cascade forest for satellite image classification. <i>International Journal of Remote Sensing</i> , 2020 , 41, 7779-7800	3.1	16
18	VGSC: A Web-Based Vector Graph Toolkit of Genome Synteny and Collinearity. <i>BioMed Research International</i> , 2016 , 2016, 7823429	3	14
17	Portfolio trading system of digital currencies: A deep reinforcement learning with multidimensional attention gating mechanism. <i>Neurocomputing</i> , 2020 , 402, 171-182	5.4	12
16	Assembly and analysis of the complete L. (Salicaceae) mitochondrial genome sequence. <i>SpringerPlus</i> , 2016 , 5, 1894		10

15	Weighted Densely Connected Convolutional Networks for Reinforcement Learning. <i>International Journal of Pattern Recognition and Artificial Intelligence</i> , 2020 , 34, 2052001	1.1	10
14	TCDNet: Trilateral Change Detection Network for Google Earth Image. <i>Remote Sensing</i> , 2020 , 12, 2669	5	9
13	The complete mitochondrial genome of. <i>Mitochondrial DNA Part B: Resources</i> , 2016 , 1, 122-123	0.5	7
12	Characterization and Analysis of the Mitochondrial Genome of Common Bean () by Comparative Genomic Approaches. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	5
11	Land use/land cover recognition in arid zone using A multi-dimensional multi-grained residual Forest?. <i>Computers and Geosciences</i> , 2020 , 144, 104557	4.5	5
10	The complete chloroplast genome sequence of an economic plant. <i>Mitochondrial DNA Part B: Resources</i> , 2017 , 2, 483-485	0.5	4
9	VGSC2: Second generation vector graph toolkit of genome synteny and collinearity. <i>Genomics</i> , 2020 , 112, 286-288	4.3	4
8	The Whole Genome Assembly and Comparative Genomic Research of <i>Thellungiella parvula</i> (Extremophile Crucifer) Mitochondrion. <i>International Journal of Genomics</i> , 2016 , 2016, 5283628	2.5	3
7	Density-based semi-supervised online sequential extreme learning machine. <i>Neural Computing and Applications</i> , 2020 , 32, 7747-7758	4.8	3
6	The complete chloroplast genome of. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 4188-4189	0.5	2
5	The complete chloroplast genome of. <i>Mitochondrial DNA Part B: Resources</i> , 2019 , 4, 4089-4090	0.5	2
4	Desert classification based on a multi-scale residual network with an attention mechanism. <i>Geosciences Journal</i> , 2021 , 25, 387-399	1.4	2
3	The complete mitochondrial genome sequence of an alpine plant. <i>Mitochondrial DNA Part B: Resources</i> , 2018 , 3, 725-727	0.5	2
2	GEsture: an online hand-drawing tool for gene expression pattern search. <i>PeerJ</i> , 2018 , 6, e4927	3.1	1
1	Non-Intrusive Load Disaggregation Based on a Multi-Scale Attention Residual Network. <i>Applied Sciences (Switzerland)</i> , 2020 , 10, 9132	2.6	0