

# Yiqing Xu

## List of Publications by Year in Descending Order

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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	Cloud/shadow segmentation based on global attention feature fusion residual network for remote sensing imagery. <i>International Journal of Remote Sensing</i> , <b>2021</b> , 42, 2022-2045	3.1	24
31	Desert classification based on a multi-scale residual network with an attention mechanism. <i>Geosciences Journal</i> , <b>2021</b> , 25, 387-399	1.4	2
30	DAU-Net: a novel water areas segmentation structure for remote sensing image. <i>International Journal of Remote Sensing</i> , <b>2021</b> , 42, 2594-2621	3.1	24
29	Characterization and Analysis of the Mitochondrial Genome of Common Bean () by Comparative Genomic Approaches. <i>International Journal of Molecular Sciences</i> , <b>2020</b> , 21,	6.3	5
28	Water Areas Segmentation from Remote Sensing Images Using a Separable Residual SegNet Network. <i>ISPRS International Journal of Geo-Information</i> , <b>2020</b> , 9, 256	2.9	24
27	Multi-Stage Feature Constraints Learning for Age Estimation. <i>IEEE Transactions on Information Forensics and Security</i> , <b>2020</b> , 15, 2417-2428	8	53
26	Portfolio trading system of digital currencies: A deep reinforcement learning with multidimensional attention gating mechanism. <i>Neurocomputing</i> , <b>2020</b> , 402, 171-182	5.4	12
25	Non-Intrusive Load Disaggregation Based on a Multi-Scale Attention Residual Network. <i>Applied Sciences (Switzerland)</i> , <b>2020</b> , 10, 9132	2.6	0
24	Land use/land cover recognition in arid zone using A multi-dimensional multi-grained residual Forest?. <i>Computers and Geosciences</i> , <b>2020</b> , 144, 104557	4.5	5
23	Dilated multi-scale cascade forest for satellite image classification. <i>International Journal of Remote Sensing</i> , <b>2020</b> , 41, 7779-7800	3.1	16
22	TCDNet: Trilateral Change Detection Network for Google Earth Image. <i>Remote Sensing</i> , <b>2020</b> , 12, 2669	5	9
21	VGSC2: Second generation vector graph toolkit of genome synteny and collinearity. <i>Genomics</i> , <b>2020</b> , 112, 286-288	4.3	4
20	Density-based semi-supervised online sequential extreme learning machine. <i>Neural Computing and Applications</i> , <b>2020</b> , 32, 7747-7758	4.8	3
19	Weighted Densely Connected Convolutional Networks for Reinforcement Learning. <i>International Journal of Pattern Recognition and Artificial Intelligence</i> , <b>2020</b> , 34, 2052001	1.1	10
18	Non-intrusive load disaggregation based on deep dilated residual network. <i>Electric Power Systems Research</i> , <b>2019</b> , 170, 277-285	3.5	50
17	Dilated residual attention network for load disaggregation. <i>Neural Computing and Applications</i> , <b>2019</b> , 31, 8931-8953	4.8	17
16	River segmentation based on separable attention residual network. <i>Journal of Applied Remote Sensing</i> , <b>2019</b> , 14, 1	1.4	17

15	The complete chloroplast genome of. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 4188-4189	0.5	2
14	The complete chloroplast genome of. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 4089-4090	0.5	2
13	Organellar genome assembly methods and comparative analysis of horticultural plants. <i>Horticulture Research</i> , <b>2018</b> , 5, 3	7.7	44
12	Least squares twin bounded support vector machines based on L1-norm distance metric for classification. <i>Pattern Recognition</i> , <b>2018</b> , 74, 434-447	7.7	52
11	GEsture: an online hand-drawing tool for gene expression pattern search. <i>PeerJ</i> , <b>2018</b> , 6, e4927	3.1	1
10	L1-Norm GEPSVM Classifier Based on an Effective Iterative Algorithm for Classification. <i>Neural Processing Letters</i> , <b>2018</b> , 48, 273-298	2.4	17
9	The complete mitochondrial genome sequence of an alpine plant. <i>Mitochondrial DNA Part B: Resources</i> , <b>2018</b> , 3, 725-727	0.5	2
8	The complete chloroplast genome sequence of an economic plant. <i>Mitochondrial DNA Part B: Resources</i> , <b>2017</b> , 2, 483-485	0.5	4
7	Assembly and comparative analysis of complete mitochondrial genome sequence of an economic plant. <i>PeerJ</i> , <b>2017</b> , 5, e3148	3.1	23
6	Analysis of the Complete Mitochondrial Genome Sequence of the Diploid Cotton by Comparative Genomics Approaches. <i>BioMed Research International</i> , <b>2016</b> , 2016, 5040598	3	25
5	Assembly and analysis of the complete L. (Salicaceae) mitochondrial genome sequence. <i>SpringerPlus</i> , <b>2016</b> , 5, 1894		10
4	The complete mitochondrial genome of. <i>Mitochondrial DNA Part B: Resources</i> , <b>2016</b> , 1, 122-123	0.5	7
3	Genome-wide identification and characterization of WRKY gene family in <i>Salix suchowensis</i> . <i>PeerJ</i> , <b>2016</b> , 4, e2437	3.1	25
2	The Whole Genome Assembly and Comparative Genomic Research of <i>Thellungiella parvula</i> (Extremophile Crucifer) Mitochondrion. <i>International Journal of Genomics</i> , <b>2016</b> , 2016, 5283628	2.5	3
1	VGSC: A Web-Based Vector Graph Toolkit of Genome Synteny and Collinearity. <i>BioMed Research International</i> , <b>2016</b> , 2016, 7823429	3	14