## Yiqing Xu

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6205948/publications.pdf

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	471061	476904
927	17	29
citations	h-index	g-index
39	39	761
docs citations	times ranked	citing authors
	citations 39	927 17 citations h-index  39 39

#	Article	IF	CITATIONS
1	Multi-Stage Feature Constraints Learning for Age Estimation. IEEE Transactions on Information Forensics and Security, 2020, 15, 2417-2428.	4.5	85
2	Least squares twin bounded support vector machines based on L1-norm distance metric for classification. Pattern Recognition, 2018, 74, 434-447.	5.1	80
3	Non-intrusive load disaggregation based on deep dilated residual network. Electric Power Systems Research, 2019, 170, 277-285.	2.1	77
4	Analysis of the Complete Mitochondrial Genome Sequence of the Diploid Cotton <i>Gossypium raimondii</i> by Comparative Genomics Approaches. BioMed Research International, 2016, 2016, 1-18.	0.9	65
5	Water Areas Segmentation from Remote Sensing Images Using a Separable Residual SegNet Network. ISPRS International Journal of Geo-Information, 2020, 9, 256.	1.4	58
6	Assembly and comparative analysis of complete mitochondrial genome sequence of an economic plant <i>Salix suchowensis</i> <li>PeerJ, 2017, 5, e3148.</li>	0.9	55
7	Organellar genome assembly methods and comparative analysis of horticultural plants. Horticulture Research, 2018, 5, 3.	2.9	53
8	Genome-wide identification and characterization of WRKY gene family in <i>Salix suchowensis</i> PeerJ, 2016, 4, e2437.	0.9	52
9	DAU-Net: a novel water areas segmentation structure for remote sensing image. International Journal of Remote Sensing, 2021, 42, 2594-2621.	1.3	49
10	Characterization and Analysis of the Mitochondrial Genome of Common Bean (Phaseolus vulgaris) by Comparative Genomic Approaches. International Journal of Molecular Sciences, 2020, 21, 3778.	1.8	44
11	Cloud/shadow segmentation based on global attention feature fusion residual network for remote sensing imagery. International Journal of Remote Sensing, 2021, 42, 2022-2045.	1.3	43
12	Dilated residual attention network for load disaggregation. Neural Computing and Applications, 2019, 31, 8931-8953.	3.2	37
13	Portfolio trading system of digital currencies: A deep reinforcement learning with multidimensional attention gating mechanism. Neurocomputing, 2020, 402, 171-182.	3.5	30
14	L1-Norm GEPSVM Classifier Based on an Effective Iterative Algorithm for Classification. Neural Processing Letters, 2018, 48, 273-298.	2.0	23
15	Dilated multi-scale cascade forest for satellite image classification. International Journal of Remote Sensing, 2020, 41, 7779-7800.	1.3	23
16	VGSC: A Web-Based Vector Graph Toolkit of Genome Synteny and Collinearity. BioMed Research International, 2016, 2016, 1-7.	0.9	22
17	River segmentation based on separable attention residual network. Journal of Applied Remote Sensing, 2019, 14, 1.	0.6	19
18	TCDNet: Trilateral Change Detection Network for Google Earth Image. Remote Sensing, 2020, 12, 2669.	1.8	15

#	Article	IF	CITATIONS
19	Assembly and analysis of the complete Salix purpurea L. (Salicaceae) mitochondrial genome sequence. SpringerPlus, 2016, 5, 1894.	1.2	14
20	The complete mitochondrial genome of Medicago truncatula. Mitochondrial DNA Part B: Resources, 2016, 1, 122-123.	0.2	11
21	Weighted Densely Connected Convolutional Networks for Reinforcement Learning. International Journal of Pattern Recognition and Artificial Intelligence, 2020, 34, 2052001.	0.7	11
22	Land use/land cover recognition in arid zone using A multi-dimensional multi-grained residual Forestâ <sup>*</sup> †. Computers and Geosciences, 2020, 144, 104557.	2.0	10
23	The complete chloroplast genome sequence of an economic plant <i>Coffea canephora</i> Mitochondrial DNA Part B: Resources, 2017, 2, 483-485.	0.2	7
24	The Whole Genome Assembly and Comparative Genomic Research of <i>Thellungiella parvula &lt; /i&gt; (Extremophile Crucifer) Mitochondrion. International Journal of Genomics, 2016, 2016, 1-13.</i>	0.8	6
25	VGSC2: Second generation vector graph toolkit of genome synteny and collinearity. Genomics, 2020, 112, 286-288.	1.3	6
26	The complete mitochondrial genome sequence of an alpine plant <i>Arabis alpina</i> . Mitochondrial DNA Part B: Resources, 2018, 3, 725-727.	0.2	5
27	Density-based semi-supervised online sequential extreme learning machine. Neural Computing and Applications, 2020, 32, 7747-7758.	3.2	5
28	Desert classification based on a multi-scale residual network with an attention mechanism. Geosciences Journal, 2021, 25, 387-399.	0.6	3
29	Non-Intrusive Load Disaggregation Based on a Multi-Scale Attention Residual Network. Applied Sciences (Switzerland), 2020, 10, 9132.	1.3	3
30	The complete chloroplast genome of Actinidia macrosperma. Mitochondrial DNA Part B: Resources, 2019, 4, 4188-4189.	0.2	2
31	The complete chloroplast genome of Actinidia fulvicoma. Mitochondrial DNA Part B: Resources, 2019, 4, 4089-4090.	0.2	2
32	The complete chloroplast genome of <i>Actinidia valvata</i> (Actinidiaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 1607-1608.	0.2	2
33	GEsture: an online hand-drawing tool for gene expression pattern search. PeerJ, 2018, 6, e4927.	0.9	2
34	DNA motif prediction of Shrub Willow (suchowensis) through comparative genetic approach. , 2017, , .		0