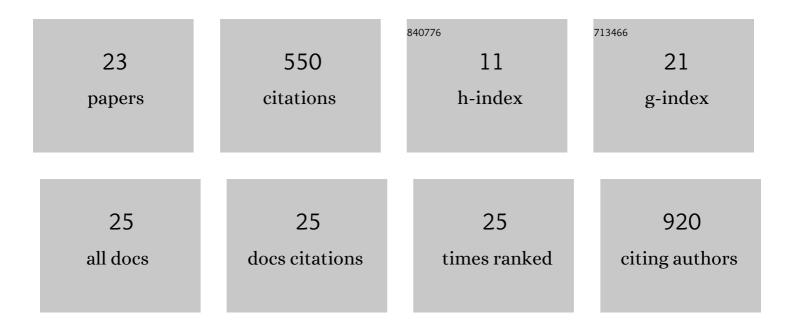
Tong Hao

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Construction of Eriocheir sinensis Protein-protein Interaction Network and Extraction of Molting Sub-network. , 2022, , .		0
2	Reconstruction of Litopenaeus vannamei Genome-Scale Metabolic Network Model and Nutritional Requirements Analysis of Different Shrimp Commercial Varieties. Frontiers in Genetics, 2021, 12, 658109.	2.3	5
3	Reconstruction of Eriocheir sinensis Y-organ Genome-Scale Metabolic Network and Differential Analysis After Eyestalk Ablation. Frontiers in Genetics, 2020, 11, 532492.	2.3	3
4	Multi-View Hierarchical Fusion Network for 3D Object Retrieval and Classification. IEEE Access, 2019, 7, 153021-153030.	4.2	10
5	The Protein–Protein Interaction Network of Litopenaeus vannamei Haemocytes. Frontiers in Physiology, 2019, 10, 156.	2.8	9
6	Mitosis event recognition and detection based on evolution of feature in time domain. Machine Vision and Applications, 2018, 29, 1249-1256.	2.7	9
7	Multiple person tracking based on slow feature analysis. Multimedia Tools and Applications, 2018, 77, 3623-3637.	3.9	8
8	Reconstruction and Analysis of a Genome-Scale Metabolic Network for Eriocheir Sinensis Hepatopancreas. IEEE Access, 2018, 6, 79235-79244.	4.2	7
9	Analyzing of Molecular Networks for Human Diseases and Drug Discovery. Current Topics in Medicinal Chemistry, 2018, 18, 1007-1014.	2.1	18
10	The Genome-Scale Integrated Networks in Microorganisms. Frontiers in Microbiology, 2018, 9, 296.	3.5	26
11	Identification of exosomes and its signature miRNAs of male and female Cynoglossus semilaevis. Scientific Reports, 2017, 7, 860.	3.3	30
12	A unified framework for cross-modality 3D model retrieval. Multimedia Tools and Applications, 2017, 76, 20217-20230.	3.9	1
13	Pathway-Consensus Approach to Metabolic Network Reconstruction for Pseudomonas putida KT2440 by Systematic Comparison of Published Models. PLoS ONE, 2017, 12, e0169437.	2.5	29
14	Reconstruction and Application of Protein–Protein Interaction Network. International Journal of Molecular Sciences, 2016, 17, 907.	4.1	67
15	Multiple/Single-View Human Action Recognition via Part-Induced Multitask Structural Learning. IEEE Transactions on Cybernetics, 2015, 45, 1194-1208.	9.5	100
16	Changes in the Organics Metabolism in the Hepatopancreas Induced by Eyestalk Ablation of the Chinese Mitten Crab Eriocheir sinensis Determined via Transcriptome and DGE Analysis. PLoS ONE, 2014, 9, e95827.	2.5	27
17	Cell type-independent mitosis event detection via hidden-state conditional neural fields. , 2014, , .		13
18	In silico metabolic engineering of Bacillus subtilis for improved production of riboflavin, Egl-237, (R,R)-2,3-butanediol and isobutanol. Molecular BioSystems, 2013, 9, 2034.	2.9	42

Tong Hao

#	Article	IF	CITATIONS
19	Partwise bagâ€ofâ€wordsâ€based multiâ€ŧask learning for human action recognition. Electronics Letters, 2013, 49, 803-805.	1.0	11
20	Nonnegative Mixed-Norm Convex Optimization for Mitotic Cell Detection in Phase Contrast Microscopy. Computational and Mathematical Methods in Medicine, 2013, 2013, 1-10.	1.3	16
21	The reconstruction and analysis of tissue specific human metabolic networks. Molecular BioSystems, 2012, 8, 663-670.	2.9	25
22	Compartmentalization of the Edinburgh Human Metabolic Network. BMC Bioinformatics, 2010, 11, 393.	2.6	92
23	Identification of exosomes and its signature miRNAs of male and female Cynoglossus semilaevis. , 0, .		1