

# Yanglan Gan

## List of Publications by Year in descending order

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Version: 2024-02-01

37  
papers

282  
citations

933264

10  
h-index

1058333

14  
g-index

37  
all docs

37  
docs citations

37  
times ranked

256  
citing authors

#	ARTICLE	IF	CITATIONS
1	Deep structural clustering for single-cell RNA-seq data jointly through autoencoder and graph neural network. <i>Briefings in Bioinformatics</i> , 2022, 23, .	3.2	22
2	DeepTSQP: Temporal-aware service QoS prediction via deep neural network and feature integration. <i>Knowledge-Based Systems</i> , 2022, 241, 108062.	4.0	20
3	DeepLTSC: Long-Tail Service Classification via Integrating Category Attentive Deep Neural Network and Feature Augmentation. <i>IEEE Transactions on Network and Service Management</i> , 2022, 19, 922-935.	3.2	6
4	Deep semi-supervised learning with contrastive learning and partial label propagation for image data. <i>Knowledge-Based Systems</i> , 2022, 245, 108602.	4.0	12
5	Energy efficient and reliable routing in wireless body area networks based on reinforcement learning and fuzzy logic. <i>Wireless Networks</i> , 2022, 28, 2669-2693.	2.0	9
6	Towards the optimality of service instance selection in mobile edge computing. <i>Knowledge-Based Systems</i> , 2021, 217, 106831.	4.0	15
7	Data tag replacement algorithm for data integrity verification in cloud storage. <i>Computers and Security</i> , 2021, 103, 102205.	4.0	7
8	Inferring gene regulatory network from single-cell transcriptomic data by integrating multiple prior networks. <i>Computational Biology and Chemistry</i> , 2021, 93, 107512.	1.1	2
9	TriPCE: A Novel Tri-Clustering Algorithm for Identifying Pan-Cancer Epigenetic Patterns. <i>Frontiers in Genetics</i> , 2020, 10, 1298.	1.1	3
10	Identification of Differential Gene Groups From Single-Cell Transcriptomes Using Network Entropy. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 588041.	1.8	3
11	NDMF: Neighborhood-Integrated Deep Matrix Factorization for Service QoS Prediction. <i>IEEE Transactions on Network and Service Management</i> , 2020, 17, 2717-2730.	3.2	23
12	LMA: label-based multi-head attentive model for long-tail web service classification. <i>International Journal of Computational Science and Engineering</i> , 2020, 23, 158.	0.4	0
13	Distance-aware Edge User Allocation with QoE Optimization. , 2020, , .		6
14	DeepWSC: A Novel Framework with Deep Neural Network for Web Service Clustering. , 2019, , .		9
15	Genome-wide analysis of epigenetic dynamics across human developmental stages and tissues. <i>BMC Genomics</i> , 2019, 20, 221.	1.2	3
16	Augmenting Labeled Probabilistic Topic Model for Web Service Classification. <i>International Journal of Web Services Research</i> , 2019, 16, 93-113.	0.5	9
17	DEEPCEN: a convolutional neural network based method for super-enhancer prediction. <i>BMC Bioinformatics</i> , 2019, 20, 598.	1.2	12
18	Towards the optimality of QoS-aware web service composition with uncertainty. <i>International Journal of Web and Grid Services</i> , 2019, 15, 1.	0.4	3

#	ARTICLE	IF	CITATIONS
19	Incremental processing for string similarity join. <i>International Journal of Computational Science and Engineering</i> , 2019, 20, 255.	0.4	0
20	Identifying influential individuals in microblogging networks using graph partitioning. <i>Expert Systems With Applications</i> , 2018, 102, 70-82.	4.4	10
21	A rapid detection algorithm of corrupted data in cloud storage. <i>Journal of Parallel and Distributed Computing</i> , 2018, 111, 115-125.	2.7	6
22	Identification of cancer subtypes from single-cell RNA-seq data using a consensus clustering method. <i>BMC Medical Genomics</i> , 2018, 11, 117.	0.7	33
23	iHMS: a database integrating human histone modification data across developmental stages and tissues. <i>BMC Bioinformatics</i> , 2017, 18, 103.	1.2	4
24	Towards Uncertain QoS-Aware Service Composition via Multi-Objective Optimization. , 2017, , .		11
25	Clustering-based uncertain QoS prediction of web services via collaborative filtering. <i>International Journal of Web and Grid Services</i> , 2017, 13, 403.	0.4	0
26	Dynamic epigenetic mode analysis using spatial temporal clustering. <i>BMC Bioinformatics</i> , 2016, 17, 537.	1.2	4
27	An integrative analysis of nucleosome occupancy and positioning using diverse sequence dependent properties. <i>Neurocomputing</i> , 2016, 206, 35-41.	3.5	0
28	UCLAO* and BHUC: Two Novel Planning Algorithms for Uncertain Web Service Composition. , 2016, , .		0
29	Computing uncertain skyline of Web services via interval number. , 2016, , .		3
30	Service composition and user modeling for personalized recommendation in cloud computing. , 2014, , .		1
31	Towards automated choreography of Web services using planning in large scale service repositories. <i>Applied Intelligence</i> , 2014, 41, 383-404.	3.3	6
32	Dynamic composition of Web services using efficient planners in large-scale service repository. <i>Knowledge-Based Systems</i> , 2014, 62, 98-112.	4.0	18
33	Towards optimal discovery of web services for multiple QoS constraints and preferences. <i>International Journal of Web Engineering and Technology</i> , 2014, 9, 277.	0.1	1
34	A novel approach to annotating web service based on interface concept mapping and semantic expansion. <i>Soft Computing</i> , 2011, 15, 929-938.	2.1	2
35	Discovering pattern-based subspace clusters by pattern tree. <i>Knowledge-Based Systems</i> , 2009, 22, 569-579.	4.0	14
36	Inferring Gene Regulatory Networks From Single-Cell Transcriptomic Data Using Bidirectional RNN. <i>Frontiers in Oncology</i> , 0, 12, .	1.3	3

#	ARTICLE	IF	CITATIONS
37	Entropy-based inference of transition states and cellular trajectory for single-cell transcriptomics. Briefings in Bioinformatics, 0, , .	3.2	2