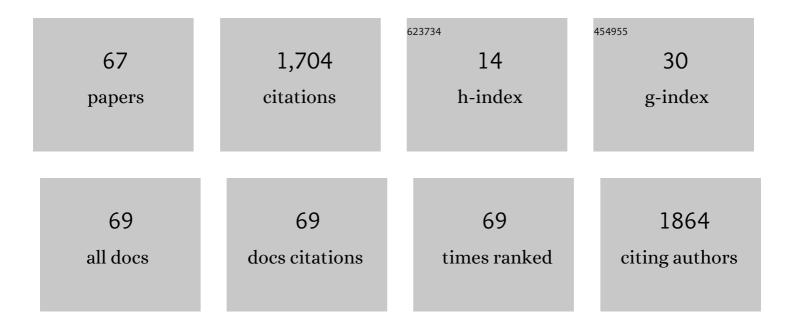
## Hung-Yu Kao

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6649902/publications.pdf

Version: 2024-02-01



ΗμΝΟ-ΥμΚλΟ

#	Article	IF	CITATIONS
1	Comparative effectiveness of antidepressants on geriatric depression: Real-world evidence from a population-based study. Journal of Affective Disorders, 2022, 296, 609-615.	4.1	14
2	Differences in the prophylactic effect of serum lithium levels on depression and mania in bipolar disorder: A dose-response meta-analysis. European Neuropsychopharmacology, 2022, 58, 20-29.	0.7	13
3	Month of birth and the risk of narcolepsy: a systematic review and meta-analysis. Journal of Clinical Sleep Medicine, 2022, 18, 1113-1120.	2.6	0
4	An Innovative Wearable Device For Monitoring Continuous Body Surface Temperature (HEARThermo): Instrument Validation Study. JMIR MHealth and UHealth, 2021, 9, e19210.	3.7	5
5	Month of birth and mental disorders: A populationâ€based study and validation using global metaâ€analysis. Acta Psychiatrica Scandinavica, 2021, 144, 153-167.	4.5	9
6	Differences in Psychiatric Comorbidities and Gender Distribution among Three Clusters of Personality Disorders: A Nationwide Population-Based Study. Journal of Clinical Medicine, 2021, 10, 3294.	2.4	4
7	Lithium concentration and recurrence risk during maintenance treatment of bipolar disorder: Multicenter cohort and metaâ€analysis. Acta Psychiatrica Scandinavica, 2021, 144, 368-378.	4.5	17
8	Predicting Serum Levels of Lithium-Treated Patients: A Supervised Machine Learning Approach. Biomedicines, 2021, 9, 1558.	3.2	6
9	True or False: Does the Deep Learning Model Learn to Detect Rumors?. , 2021, , .		1
10	Continuous temperature monitoring by a wearable device for early detection of febrile events in the SARS-CoV-2 outbreak in Taiwan, 2020. Journal of Microbiology, Immunology and Infection, 2020, 53, 503-504.	3.1	21
11	C-3PO: Click-sequence-aware deeP neural network (DNN)-based Pop-uPs recOmmendation. Soft Computing, 2019, 23, 11793-11799.	3.6	3
12	\$\$k^{-}\$\$ kanonymization of multiple shortest paths. Soft Computing, 2017, 21, 4215-4226.	3.6	4
13	Special issue on soft computing for knowledge management and web applications. Soft Computing, 2017, 21, 281-282.	3.6	0
14	CDRnN: A high performance chemical-disease recognizer in biomedical literature. , 2017, , .		2
15	Data-Driven and Deep Learning Methodology for Deceptive Advertising and Phone Scams Detection. , 2017, , .		4
16	AuDis: an automatic CRF-enhanced disease normalization in biomedical text. Database: the Journal of Biological Databases and Curation, 2016, 2016, baw091.	3.0	17
17	GNormPlus: An Integrative Approach for Tagging Genes, Gene Families, and Protein Domains. BioMed Research International, 2015, 2015, 1-7.	1.9	155
18	Text Mining for Translational Bioinformatics. BioMed Research International, 2015, 2015, 1-2.	1.9	4

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#	Article	IF	CITATIONS
19	Identification of item features in microblogging data. , 2015, , .		Ο
20	LDA based semi-supervised learning from streaming short text. , 2015, , .		5
21	On anonymizing transactions with sensitive items. Applied Intelligence, 2014, 41, 1043-1058.	5.3	52
22	[K1, K2]-anonymization of Shortest Paths. , 2014, , .		1
23	Text mining tools for assisting literature curation. , 2014, , .		2
24	User preference space partition and product filters for reverse top-k queries. , 2014, , .		0
25	IT2FS-based ontology with soft-computing mechanism for malware behavior analysis. Soft Computing, 2014, 18, 267-284.	3.6	22
26	Using hidden Markov models to predict DNA-binding proteins with sequence and structure information. Soft Computing, 2014, 18, 2365-2376.	3.6	7
27	SHORTEST PATHS ANONYMIZATION ON WEIGHTED GRAPHS. International Journal of Software Engineering and Knowledge Engineering, 2013, 23, 65-79.	0.8	6
28	An Adaptive Page Clustering Based Weighting Method for Information Retrieval. , 2013, , .		0
29	Genealogical-based method for ontology self-extension in MeSH. , 2013, , .		0
30	Introduction to the 23rd International Conference on Genome Informatics (GIW2012). Gene, 2013, 518, 1.	2.2	0
31	PubTator: a web-based text mining tool for assisting biocuration. Nucleic Acids Research, 2013, 41, W518-W522.	14.5	523
32	tmVar: a text mining approach for extracting sequence variants in biomedical literature. Bioinformatics, 2013, 29, 1433-1439.	4.1	197
33	CoIN: a network analysis for document triage. Database: the Journal of Biological Databases and Curation, 2013, 2013, bat076.	3.0	4
34	Unsupervised corpus distillation for represented indicator measurement on focus species detection. International Journal of Data Mining and Bioinformatics, 2013, 8, 413.	0.1	1
35	Large-Scale Event Extraction from Literature with Multi-Level Gene Normalization. PLoS ONE, 2013, 8, e55814.	2.5	83
36	Using a Search Engine-Based Mutually Reinforcing Approach to Assess the Semantic Relatedness of Biomedical Terms. PLoS ONE, 2013, 8, e77868.	2.5	2

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37	Accelerating literature curation with text-mining tools: a case study of using PubTator to curate genes in PubMed abstracts. Database: the Journal of Biological Databases and Curation, 2012, 2012, bas041-bas041.	3.0	83
38	The Retrieval of Important News Stories by Influence Propagation among Communities and Categories. , 2012, , .		0
39	Link Prediction in a Bipartite Network Using Wikipedia Revision Information. , 2012, , .		13
40	Confining Edge Types in K-anonymization of Shortest Paths. , 2012, , .		1
41	TWMAN+: A Type-2 fuzzy ontology model for malware behavior analysis. , 2012, , .		5
42	SR4GN: A Species Recognition Software Tool for Gene Normalization. PLoS ONE, 2012, 7, e38460.	2.5	71
43	Evolutional Dependency Parse Trees for Biological Relation Extraction. , 2011, , .		1
44	K-anonymity on sensitive transaction items. , 2011, , .		1
45	Augmented Transitive Relationships in Direct Protein-Protein Interaction Prediction. , 2011, , .		0
46	Measuring Semantic Relatedness Using Wikipedia Revision Information in a Signed Network. , 2011, , .		0
47	Applying FML and Fuzzy Ontologies to malware behavioural analysis. , 2011, , .		12
48	Malware behavioral analysis system: TWMAN. , 2011, , .		14
49	The gene normalization task in BioCreative III. BMC Bioinformatics, 2011, 12, S2.	2.6	101
50	Cross-species gene normalization by species inference. BMC Bioinformatics, 2011, 12, S5.	2.6	61
51	CAIS: Community Based Annotation Insight Search in a Folksonomy Network. , 2011, , .		0
52	Protein surface search in DNA-binding protein prediction by Delaunay triangulation modeling. , 2010, , .		1
53	Anonymizing Set-Valued Social Data. , 2010, , .		7
54	Represented indicator measurement and corpus distillation on focus species detection. , 2010, , .		1

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#	Article	IF	CITATIONS
55	A Categorized Sentiment Analysis of Chinese Reviews by Mining Dependency in Product Features and Opinions from Blogs. , 2010, , .		7
56	Incrementally Mining High Utility Itemsets in Dynamic Databases. , 2010, , .		2
57	Blog Popularity Mining by Social Interconnection Analysis. IEEE Internet Computing, 2010, , .	3.3	10
58	Hiding Sensitive Association Rules on Stars. , 2010, , .		2
59	Multi-table association rules hiding. , 2010, , .		2
60	Entropy-Based Visual Tree Evaluation on Block Extraction. , 2009, , .		3
61	Normalizing Biomedical Name Entities by Similarity-Based Inference Network and De-ambiguity Mining. , 2009, , .		2
62	A Fast PageRank Convergence Method based on the Cluster Prediction. , 2007, , .		9
63	The Mining and Extraction of Primary Informative Blocks and Data Objects from Systematic Web Pages. , 2006, , .		9
64	WISDOM: Web intrapage informative structure mining based on document object model. IEEE Transactions on Knowledge and Data Engineering, 2005, 17, 614-627.	5.7	37
65	Mining web informative structures and contents based on entropy analysis. IEEE Transactions on Knowledge and Data Engineering, 2004, 16, 41-55.	5.7	65
66	Clustering for Web information hierarchy mining. , 0, , .		0
67	Web information hierarchy and importance mining based on DOM information distillation. , 0, , .		0