

BlaÅ¾ Å krlj

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

329
citations

759233

12
h-index

996975

15
g-index

40
all docs

40
docs citations

40
times ranked

287
citing authors

#	ARTICLE	IF	CITATIONS
1	tax2vec: Constructing Interpretable Features from Taxonomies for Short Text Classification. <i>Computer Speech and Language</i> , 2021, 65, 101104.	4.3	27
2	Prioritisation of Compounds for 3CLpro Inhibitor Development on SARS-CoV-2 Variants. <i>Molecules</i> , 2021, 26, 3003.	3.8	23
3	Propositionalization and embeddings: two sides of the same coin. <i>Machine Learning</i> , 2020, 109, 1465-1507.	5.4	20
4	RaKUn: Rank-based Keyword Extraction via Unsupervised Learning and Meta Vertex Aggregation. <i>Lecture Notes in Computer Science</i> , 2019, , 311-323.	1.3	17
5	Towards Robust Text Classification with Semantics-Aware Recurrent Neural Architecture. <i>Machine Learning and Knowledge Extraction</i> , 2019, 1, 575-589.	5.0	16
6	To BAN or Not to BAN: Bayesian Attention Networks for Reliable Hate Speech Detection. <i>Cognitive Computation</i> , 2022, 14, 353-371.	5.2	16
7	ProBiS-Dock Database: A Web Server and Interactive Web Repository of Small Ligand-Protein Binding Sites for Drug Design. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 4097-4107.	5.4	16
8	GenProBiS: web server for mapping of sequence variants to protein binding sites. <i>Nucleic Acids Research</i> , 2017, 45, W253-W259.	14.5	14
9	Py3plex toolkit for visualization and analysis of multilayer networks. <i>Applied Network Science</i> , 2019, 4, .	1.5	14
10	Zero-Shot Learning for Cross-Lingual News Sentiment Classification. <i>Applied Sciences (Switzerland)</i> , 2020, 10, 5993.	2.5	14
11	A comprehensive comparison of molecular feature representations for use in predictive modeling. <i>Computers in Biology and Medicine</i> , 2021, 130, 104197.	7.0	13
12	BoBER: web interface to the base of bioisosterically exchangeable replacements. <i>Journal of Cheminformatics</i> , 2017, 9, 62.	6.1	12
13	Prediction Uncertainty Estimation for Hate Speech Classification. <i>Lecture Notes in Computer Science</i> , 2019, , 286-298.	1.3	12
14	CBSSD: community-based semantic subgroup discovery. <i>Journal of Intelligent Information Systems</i> , 2019, 53, 265-304.	3.9	11
15	Embedding-based Silhouette community detection. <i>Machine Learning</i> , 2020, 109, 2161-2193.	5.4	11
16	Differential Response of Grapevine to Infection with <i>Candidatus Phytoplasma solani</i> ™ in Early and Late Growing Season through Complex Regulation of mRNA and Small RNA Transcriptomes. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3531.	4.1	10
17	autoBOT: evolving neuro-symbolic representations for explainable low resource text classification. <i>Machine Learning</i> , 2021, 110, 989-1028.	5.4	8
18	Fuzzy Jaccard Index: A robust comparison of ordered lists. <i>Applied Soft Computing Journal</i> , 2021, 113, 107849.	7.2	7

#	ARTICLE	IF	CITATIONS
19	Community-Based Semantic Subgroup Discovery. Lecture Notes in Computer Science, 2018, , 182-196.	1.3	7
20	Computational identification of non-synonymous polymorphisms within regions corresponding to protein interaction sites. Computers in Biology and Medicine, 2016, 79, 30-35.	7.0	6
21	Py3plex: A Library for Scalable Multilayer Network Analysis and Visualization. Studies in Computational Intelligence, 2019, , 757-768.	0.9	6
22	Layer entanglement in multiplex, temporal multiplex, and coupled multilayer networks. Applied Network Science, 2020, 5, .	1.5	6
23	Identification of COVID-19 Related Fake News via Neural Stacking. Communications in Computer and Information Science, 2021, , 177-188.	0.5	6
24	Identification of Sequence Variants within Experimentally Validated Protein Interaction Sites Provides New Insights into Molecular Mechanisms of Disease Development. Molecular Informatics, 2017, 36, 1700017.	2.5	4
25	Insights from Ion Binding Site Network Analysis into Evolution and Functions of Proteins. Molecular Informatics, 2018, 37, e1700144.	2.5	4
26	SNoRe: Scalable Unsupervised Learning of Symbolic Node Representations. IEEE Access, 2020, 8, 212568-212588.	4.2	4
27	ProBiS-Dock: A Hybrid Multitemplate Homology Flexible Docking Algorithm Enabled by Protein Binding Site Comparison. Journal of Chemical Information and Modeling, 2022, 62, 1573-1584.	5.4	4
28	New Cross-Talks between Pathways Involved in Grapevine Infection with <i>Candidatus Phytoplasma solani</i> ™ Revealed by Temporal Network Modelling. Plants, 2021, 10, 646.	3.5	3
29	ReliefE: feature ranking in high-dimensional spaces via manifold embeddings. Machine Learning, 2022, 111, 273-317.	5.4	3
30	Investigating cross-lingual training for offensive language detection. PeerJ Computer Science, 2021, 7, e559.	4.5	3
31	Semantic Reasoning from Model-Agnostic Explanations. , 2021, , .		2
32	JSI at the FinSim-2 task: Ontology-Augmented Financial Concept Classification. , 2021, , .		2
33	Deep node ranking for neuro-symbolic structural node embedding and classification. International Journal of Intelligent Systems, 2022, 37, 914-943.	5.7	2
34	Targeted End-to-End Knowledge Graph Decomposition. Lecture Notes in Computer Science, 2018, , 157-171.	1.3	2
35	Relational tree ensembles and feature rankings. Knowledge-Based Systems, 2022, 251, 109254.	7.1	2
36	Ensemble- and distance-based feature ranking for unsupervised learning. International Journal of Intelligent Systems, 2021, 36, 3068-3086.	5.7	1

#	ARTICLE	IF	CITATIONS
37	Patterns of Multiplex Layer Entanglement Across Real and Synthetic Networks. Studies in Computational Intelligence, 2020, , 671-683.	0.9	1
38	PubMed-Scale Chemical Concept Embeddings Reconstruct Physical Protein Interaction Networks. Frontiers in Research Metrics and Analytics, 2021, 6, 644614.	1.9	0
39	Unsupervised Feature Ranking via Attribute Networks. Lecture Notes in Computer Science, 2021, , 334-343.	1.3	0
40	Prioritization of COVID-19-Related Literature via Unsupervised Keyphrase Extraction and Document Representation Learning. Lecture Notes in Computer Science, 2021, , 204-217.	1.3	0