## Arzucan Ozgur

List of Publications by Year in descending order

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361413 254184 2,371 50 20 43 citations h-index g-index papers 51 51 51 2658 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	DeepDTA: deep drug–target binding affinity prediction. Bioinformatics, 2018, 34, i821-i829.	4.1	717
2	Identifying gene-disease associations using centrality on a literature mined gene-interaction network. Bioinformatics, 2008, 24, i277-i285.	4.1	314
3	PHISTO: pathogen–host interaction search tool. Bioinformatics, 2013, 29, 1357-1358.	4.1	145
4	A comparative study of SMILES-based compound similarity functions for drug-target interaction prediction. BMC Bioinformatics, 2016, 17, 128.	2.6	99
5	BIOSSES: a semantic sentence similarity estimation system for the biomedical domain. Bioinformatics, 2017, 33, i49-i58.	4.1	97
6	A review on computational systems biology of pathogenââ,¬â€œhost interactions. Frontiers in Microbiology, 2015, 6, 235.	3.5	93
7	GLASS: a comprehensive database for experimentally validated GPCR-ligand associations. Bioinformatics, 2015, 31, 3035-3042.	4.1	92
8	Michigan molecular interactions r2: from interacting proteins to pathways. Nucleic Acids Research, 2009, 37, D642-D646.	14.5	85
9	Exploring chemical space using natural language processing methodologies for drug discovery. Drug Discovery Today, 2020, 25, 689-705.	6.4	68
10	Introducing meta-services for biomedical information extraction. Genome Biology, 2008, 9, S6.	8.8	61
11	Mining of vaccine-associated IFN- $\hat{l}^3$ gene interaction networks using the Vaccine Ontology. Journal of Biomedical Semantics, 2011, 2, S8.	1.6	54
12	Classification of Beta-Lactamases and Penicillin Binding Proteins Using Ligand-Centric Network Models. PLoS ONE, 2015, 10, e0117874.	2.5	41
13	Literature-Based Discovery of IFN- <mml:math xmlns:mml="http://www.w3.org/1998/Math/MathML"><mml:mi>γ</mml:mi></mml:math> and Vaccine-Mediated Gene Interaction Networks. Journal of Biomedicine and Biotechnology, 2010, 2010, 1-13	3.0	34
14	A novel methodology on distributed representations of proteins using their interacting ligands. Bioinformatics, 2018, 34, i295-i303.	4.1	31
15	Improving Named Entity Recognition for Morphologically Rich Languages Using Word Embeddings. , 2014, , .		30
16	Overview of the BioCreative VI Precision Medicine Track: mining protein interactions and mutations for precision medicine. Database: the Journal of Biological Databases and Curation, 2019, 2019, .	3.0	30
17	BioCreative V BioC track overview: collaborative biocurator assistant task for BioGRID. Database: the Journal of Biological Databases and Curation, 2016, 2016, baw121.	3.0	28
18	Identification of fever and vaccine-associated gene interaction networks using ontology-based literature mining. Journal of Biomedical Semantics, 2012, 3, 18.	1.6	26

#	Article	IF	Citations
19	Development and application of an interaction network ontology for literature mining of vaccine-associated gene-gene interactions. Journal of Biomedical Semantics, 2015, 6, 2.	1.6	23
20	CO-OCCURRENCE NETWORK OF REUTERS NEWS. International Journal of Modern Physics C, 2008, 19, 689-702.	1.7	22
21	Linking entities through an ontology using word embeddings and syntactic re-ranking. BMC Bioinformatics, 2019, 20, 156.	2.6	22
22	Identifying Topics in Microblogs Using Wikipedia. PLoS ONE, 2016, 11, e0151885.	2.5	15
23	Bayesian Pathway Analysis of Cancer Microarray Data. PLoS ONE, 2014, 9, e102803.	2.5	13
24	Ontology-based literature mining of E. coli vaccine-associated gene interaction networks. Journal of Biomedical Semantics, 2017, 8, 12.	1.6	13
25	Machine learning-based identification and rule-based normalization of adverse drug reactions in drug labels. BMC Bioinformatics, 2019, 20, 707.	2.6	13
26	ChemBoost: A Chemical Language Based Approach for Protein – Ligand Binding Affinity Prediction. Molecular Informatics, 2021, 40, e2000212.	2.5	12
27	U-Compare bio-event meta-service: compatible BioNLP event extraction services. BMC Bioinformatics, 2011, 12, 481.	2.6	10
28	Literature Mining and Ontology based Analysis of Host-Brucella Gene–Gene Interaction Network. Frontiers in Microbiology, 2015, 6, 1386.	3.5	9
29	A novel gene selection method for gene expression data for the task of cancer type classification. Biology Direct, 2021, 16, 7.	4.6	9
30	Efficient indexing technique for XML-based electronic product catalogs. Electronic Commerce Research and Applications, 2006, 5, 66-77.	5.0	7
31	Detection and categorization of bacteria habitats using shallow linguistic analysis. BMC Bioinformatics, 2015, 16, S5.	2.6	7
32	Question Analysis for a Closed Domain Question Answering System. Lecture Notes in Computer Science, 2015, , 468-482.	1.3	7
33	The Interaction Network Ontology-supported modeling and mining of complex interactions represented with multiple keywords in biomedical literature. BioData Mining, 2016, 9, 41.	4.0	7
34	A closed-domain question answering framework using reliable resources to assist students. Natural Language Engineering, 2018, 24, 725-762.	2.5	7
35	Ontology-based literature mining and class effect analysis of adverse drug reactions associated with neuropathy-inducing drugs. Journal of Biomedical Semantics, 2018, 9, 17.	1.6	7
36	Ontology-Based Categorization of Bacteria and Habitat Entities using Information Retrieval Techniques. , $2016,  ,  .$		7

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#	Article	IF	CITATIONS
37	Turkish Treebanking: Unifying and Constructing Efforts. , 2019, , .		7
38	Automatic query generation using word embeddings for retrieving passages describing experimental methods. Database: the Journal of Biological Databases and Curation, 2017, 2017, baw166.	3.0	6
39	Segmenting hashtags and analyzing their grammatical structure. Journal of the Association for Information Science and Technology, 2018, 69, 675-686.	2.9	6
40	Resources for Turkish dependency parsing: introducing the BOUN Treebank and the BoAT annotation tool. Language Resources and Evaluation, 2022, 56, 259-307.	2.7	6
41	Automated Neuroanatomical Relation Extraction: A Linguistically Motivated Approach with a PVT Connectivity Graph Case Study. Frontiers in Neuroinformatics, 2016, 10, 39.	2.5	5
42	Social Network of Co-occurrence in News Articles. Lecture Notes in Computer Science, 2004, , 688-695.	1.3	5
43	Improving the Annotations in the Turkish Universal Dependency Treebank. , 2019, , .		5
44	Statistical representation models for mutation information within genomic data. BMC Bioinformatics, 2019, 20, 324.	2.6	4
45	Classification of Skewed and Homogenous Document Corpora with Class-Based and Corpus-Based Keywords. , 2006, , 91-101.		4
46	N-gram Parsing for Jointly Training a Discriminative Constituency Parser. Polibits, 0, 47, 5-12.	0.0	2
47	Rule-based focus extraction in Turkish question answering systems. , 2014, , .		1
48	Identifying Image Related Sentences in News Articles. , 2019, , .		1
49	Cluster-based mention typing for named entity disambiguation. Natural Language Engineering, 2020, , $1-37$ .	2.5	1
50	Editorial: Machine Learning Methodologies to Study Molecular Interactions. Frontiers in Molecular Biosciences, 2021, 8, 806474.	3.5	1