

Ling Luo

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

17
papers

351
citations

8
h-index

18
g-index

22
ext. papers

560
ext. citations

6.7
avg, IF

3.57
L-index

#	Paper	IF	Citations
17	PhenoRerank: a re-ranking model for phenotypic concept recognition pre-trained on human phenotype ontology.. <i>Journal of Biomedical Informatics</i> , 2022 , 104059	10.2	
16	LitSuggest: a web-based system for literature recommendation and curation using machine learning. <i>Nucleic Acids Research</i> , 2021 , 49, W352-W358	20.1	9
15	Artificial Intelligence in Action: Addressing the COVID-19 Pandemic with Natural Language Processing. <i>Annual Review of Biomedical Data Science</i> , 2021 , 4, 313-339	5.6	11
14	PhenoTagger: A Hybrid Method for Phenotype Concept Recognition using Human Phenotype Ontology. <i>Bioinformatics</i> , 2021 ,	7.2	1
13	A neural network-based joint learning approach for biomedical entity and relation extraction from biomedical literature. <i>Journal of Biomedical Informatics</i> , 2020 , 103, 103384	10.2	15
12	KGHC: a knowledge graph for hepatocellular carcinoma. <i>BMC Medical Informatics and Decision Making</i> , 2020 , 20, 135	3.6	3
11	Exploiting sequence labeling framework to extract document-level relations from biomedical texts. <i>BMC Bioinformatics</i> , 2020 , 21, 125	3.6	8
10	Overview of the BioCreative VI Precision Medicine Track: mining protein interactions and mutations for precision medicine. <i>Database: the Journal of Biological Databases and Curation</i> , 2019 , 2019,	5	18
9	HMNPPID-human malignant neoplasm protein-protein interaction database. <i>Human Genomics</i> , 2019 , 13, 44	6.8	1
8	An attention-based BiLSTM-CRF approach to document-level chemical named entity recognition. <i>Bioinformatics</i> , 2018 , 34, 1381-1388	7.2	130
7	A multi-task learning based approach to biomedical entity relation extraction 2018 ,		3
6	A neural network approach to chemical and gene/protein entity recognition in patents. <i>Journal of Cheminformatics</i> , 2018 , 10, 65	8.6	5
5	Document triage for identifying protein-protein interactions affected by mutations: a neural network ensemble approach. <i>Database: the Journal of Biological Databases and Curation</i> , 2018 , 2018,	5	3
4	Disease named entity recognition from biomedical literature using a novel convolutional neural network. <i>BMC Medical Genomics</i> , 2017 , 10, 73	3.7	21
3	An attention-based effective neural model for drug-drug interactions extraction. <i>BMC Bioinformatics</i> , 2017 , 18, 445	3.6	41
2	ML-CNN: A novel deep learning based disease named entity recognition architecture 2016 ,		3
1	Drug drug interaction extraction from biomedical literature using syntax convolutional neural network. <i>Bioinformatics</i> , 2016 , 32, 3444-3453	7.2	78

