Senay Kafkas

List of Publications by Year in descending order

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933447 940533 19 291 10 16 citations h-index g-index papers 24 24 24 481 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Assessment of NER solutions against the first and second CALBC Silver Standard Corpus. Journal of Biomedical Semantics, 2011, 2, S11.	1.6	39
2	DeepViral: prediction of novel virus–host interactions from protein sequences and infectious disease phenotypes. Bioinformatics, 2021, 37, 2722-2729.	4.1	35
3	Database Citation in Full Text Biomedical Articles. PLoS ONE, 2013, 8, e63184.	2.5	33
4	Database citation in supplementary data linked to Europe PubMed Central full text biomedical articles. Journal of Biomedical Semantics, 2015, 6, 1.	1.6	27
5	<scp><i>EMC10</i></scp> homozygous variant identified in a family with global developmental delay, mild intellectual disability, and speech delay. Clinical Genetics, 2020, 98, 555-561.	2.0	24
6	Literature evidence in open targets - a target validation platform. Journal of Biomedical Semantics, 2017, 8, 20.	1.6	16
7	Evaluating gold standard corpora against gene/protein tagging solutions and lexical resources. Journal of Biomedical Semantics, 2013, 4, 28.	1.6	15
8	Combining lexical and context features for automatic ontology extension. Journal of Biomedical Semantics, 2020, 11, 1.	1.6	14
9	Section level search functionality in Europe PMC. Journal of Biomedical Semantics, 2015, 6, 7.	1.6	13
10	PathoPhenoDB, linking human pathogens to their phenotypes in support of infectious disease research. Scientific Data, 2019, 6, 79.	5. 3	13
11	Ontology based text mining of gene-phenotype associations: application to candidate gene prediction. Database: the Journal of Biological Databases and Curation, 2019, 2019, .	3.0	10
12	What is the right sequencing approach? Solo VS extended family analysis in consanguineous populations. BMC Medical Genomics, 2020, 13, 103.	1.5	10
13	Entity Recognition in Parallel Multi-lingual Biomedical Corpora: The CLEF-ER Laboratory Overview. Lecture Notes in Computer Science, 2013, , 353-367.	1.3	9
14	Ontology based mining of pathogen–disease associations from literature. Journal of Biomedical Semantics, 2019, 10, 15.	1.6	7
15	Linking common human diseases to their phenotypes; development of a resource for human phenomics. Journal of Biomedical Semantics, 2021, 12, 17.	1.6	6
16	Monitoring named entity recognition: the League Table. Journal of Biomedical Semantics, 2013, 4, 19.	1.6	4
17	Functional variation of alternative splice forms in their protein interaction networks: a literature mining approach. BMC Bioinformatics, 2010, 11 , .	2.6	3
18	Diversity in the Interactions of Isoforms Linked to Clustered Transcripts: A Systematic Literature Analysis. Journal of Proteomics and Bioinformatics, 2011, 4, .	0.4	2

#	Article	IF	CITATIONS
19	Usage of cell nomenclature in biomedical literature. BMC Bioinformatics, 2017, 18, 561.	2.6	1