

# Justin Zobel

## List of Publications by Year in descending order

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Version: 2024-02-01

63  
papers

5,990  
citations

331259

21  
h-index

214527

47  
g-index

70  
all docs

70  
docs citations

70  
times ranked

6549  
citing authors

#	ARTICLE	IF	CITATIONS
1	Benchmarks for measurement of duplicate detection methods in nucleotide databases. Database: the Journal of Biological Databases and Curation, 2023, 2023, .	1.4	10
2	Measurement of clustering effectiveness for document collections. Information Retrieval, 2022, 25, 239-268.	1.6	5
3	Immediate Text Search on Streams Using Apoptotic Indexes. Lecture Notes in Computer Science, 2022, , 157-169.	1.0	2
4	Exploring automatic inconsistency detection for literature-based gene ontology annotation. Bioinformatics, 2022, 38, i273-i281.	1.8	0
5	Automatic consistency assurance for literature-based gene ontology annotation. BMC Bioinformatics, 2021, 22, 565.	1.2	2
6	Document Clustering vs Topic Models: A Case Study. , 2021, , .		1
7	Quality Matters: Biocuration Experts on the Impact of Duplication and Other Data Quality Issues in Biological Databases. Genomics, Proteomics and Bioinformatics, 2020, 18, 91-103.	3.0	14
8	GeneMates: an R package for detecting horizontal gene co-transfer between bacteria using gene-gene associations controlled for population structure. BMC Genomics, 2020, 21, 658.	1.2	9
9	Generation of Synthetic Query Auto Completion Logs. Lecture Notes in Computer Science, 2020, , 621-635.	1.0	6
10	Exploring effective approaches for haplotype block phasing. BMC Bioinformatics, 2019, 20, 540.	1.2	24
11	Abstraction of query auto-completion logs for anonymity-preserving analysis. Information Retrieval, 2019, 22, 499-524.	1.6	4
12	Automated assessment of biological database assertions using the scientific literature. BMC Bioinformatics, 2019, 20, 216.	1.2	3
13	Modeling User Actions in Job Search. Lecture Notes in Computer Science, 2019, , 652-664.	1.0	3
14	Search Effectiveness in Nonredundant Sequence Databases: Assessments and Solutions. Journal of Computational Biology, 2019, 26, 605-617.	0.8	1
15	Can machine translation systems be evaluated by the crowd alone. Natural Language Engineering, 2017, 23, 3-30.	2.1	70
16	Duplicates, redundancies and inconsistencies in the primary nucleotide databases: a descriptive study. Database: the Journal of Biological Databases and Curation, 2017, 2017, baw163.	1.4	36
17	Literature consistency of bioinformatics sequence databases is effective for assessing record quality. Database: the Journal of Biological Databases and Curation, 2017, 2017, .	1.4	8
18	Automated detection of records in biological sequence databases that are inconsistent with the literature. Journal of Biomedical Informatics, 2017, 71, 229-240.	2.5	12

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19	A Taxonomy of Query Auto Completion Modes. , 2017, , .		14
20	Learning Biological Sequence Types Using the Literature. , 2017, , .		2
21	Evaluation of CD-HIT for constructing non-redundant databases. , 2016, , .		10
22	A categorical analysis of coreference resolution errors in biomedical texts. Journal of Biomedical Informatics, 2016, 60, 309-318.	2.5	7
23	Coreference resolution improves extraction of Biological Expression Language statements from texts. Database: the Journal of Biological Databases and Curation, 2016, 2016, baw076.	1.4	9
24	Medical information retrieval: introduction to the special issue. Information Retrieval, 2016, 19, 1-5.	1.6	26
25	Supervised Learning for Detection of Duplicates in Genomic Sequence Databases. PLoS ONE, 2016, 11, e0159644.	1.1	10
26	High performance computing enabling exhaustive analysis of higher order single nucleotide polymorphism interaction in Genome Wide Association Studies. Health Information Science and Systems, 2015, 3, S3.	3.4	24
27	Bandage: interactive visualization of <i>de novo</i> genome assemblies. Bioinformatics, 2015, 31, 3350-3352.	1.8	1,671
28	Evaluation of a Machine Learning Duplicate Detection Method for Bioinformatics Databases. , 2015, , .		17
29	SRST2: Rapid genomic surveillance for public health and hospital microbiology labs. Genome Medicine, 2014, 6, 90.	3.6	953
30	Accurate and Robust Genomic Prediction of Celiac Disease Using Statistical Learning. PLoS Genetics, 2014, 10, e1004137.	1.5	95
31	Using Inter-file Similarity to Improve Intra-file Compression. , 2014, , .		3
32	Storing a Collection of Differentially Compressed Files Recursively. , 2014, , .		1
33	Supercomputing enabling exhaustive statistical analysis of genome wide association study data: Preliminary results. , 2012, 2012, 1258-61.		5
34	Redesigning the string hash table, burst trie, and BST to exploit cache. Journal of Experimental Algorithmics, 2010, 15, .	0.7	8
35	Against recall. ACM SIGIR Forum, 2009, 43, 3-8.	0.4	16
36	Rank-biased precision for measurement of retrieval effectiveness. ACM Transactions on Information Systems, 2008, 27, 1-27.	3.8	405

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37	SICO: A System for Detection of Near-Duplicate Images During Search. , 2007, , .		8
38	Does topic metadata help with Web search?. Journal of the Association for Information Science and Technology, 2007, 58, 613-628.	2.6	12
39	Efficient plagiarism detection for large code repositories. Software - Practice and Experience, 2007, 37, 151-175.	2.5	64
40	A pipelined architecture for distributed text query evaluation. Information Retrieval, 2007, 10, 205-231.	1.6	88
41	Compression techniques for fast external sorting. VLDB Journal, 2007, 16, 269-291.	2.7	9
42	Inverted files for text search engines. ACM Computing Surveys, 2006, 38, 6.	16.1	797
43	Searchable words on the Web. International Journal on Digital Libraries, 2005, 5, 99-105.	1.1	33
44	Fast on-line index construction by geometric partitioning. , 2005, , .		37
45	Chinese OOV translation and post-translation query expansion in chinese-english cross-lingual information retrieval. ACM Transactions on Asian Language Information Processing, 2005, 4, 57-77.	0.8	14
46	An architecture for effective music information retrieval. Journal of the Association for Information Science and Technology, 2004, 55, 1053-1057.	2.6	8
47	Methods for identifying versioned and plagiarized documents. Journal of the Association for Information Science and Technology, 2003, 54, 203-215.	2.6	220
48	Efficient single-pass index construction for text databases. Journal of the Association for Information Science and Technology, 2003, 54, 713-729.	2.6	73
49	Burst tries. ACM Transactions on Information Systems, 2002, 20, 192-223.	3.8	113
50	Self-adjusting trees in practice for large text collections. Software - Practice and Experience, 2001, 31, 925-939.	2.5	29
51	Compression techniques for Chinese text. Software - Practice and Experience, 1998, 28, 1299-1314.	2.5	2
52	Inverted files versus signature files for text indexing. ACM Transactions on Database Systems, 1998, 23, 453-490.	1.5	249
53	How reliable are the results of large-scale information retrieval experiments?. , 1998, , .		359
54	Passage retrieval revisited. ACM SIGIR Forum, 1997, 31, 178-185.	0.4	35

#	ARTICLE	IF	CITATIONS
55	A Formal Model for Representation and Querying of Structured Documents. Journal of Systems Integration, 1997, 7, 31-46.	0.1	1
56	Performance in Practice of String Hashing Functions. , 1997, , .		40
57	Filtered document retrieval with frequency-sorted indexes. Journal of the Association for Information Science and Technology, 1996, 47, 749-764.	1.2	66
58	Filtered document retrieval with frequency-sorted indexes. , 1996, 47, 749.		4
59	Filtered document retrieval with frequency-sorted indexes. , 1996, 47, 749.		63
60	Finding approximate matches in large lexicons. Software - Practice and Experience, 1995, 25, 331-345.	2.5	94
61	Adding compression to a full-text retrieval system. Software - Practice and Experience, 1995, 25, 891-903.	2.5	72
62	Rejoinder to letter from Stephen P. Harter. Journal of the Association for Information Science and Technology, 1993, 44, 213-213.	1.2	0
63	A model for word clustering. Journal of the Association for Information Science and Technology, 1992, 43, 616-627.	1.2	8