Brett R South

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10883086/publications.pdf

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21 papers	1,825 citations	623734 14 h-index	752698 20 g-index
21 all docs	21 docs citations	21 times ranked	1700 citing authors

#	Article	IF	Citations
1	Normalizing acronyms and abbreviations to aid patient understanding of clinical texts: ShARe/CLEF eHealth Challenge 2013, Task 2. Journal of Biomedical Semantics, 2016, 7, 43.	1.6	16
2	Extracting a stroke phenotype risk factor from Veteran Health Administration clinical reports: an information content analysis. Journal of Biomedical Semantics, 2016, 7, 26.	1.6	38
3	Evaluating the state of the art in disorder recognition and normalization of the clinical narrative. Journal of the American Medical Informatics Association: JAMIA, 2015, 22, 143-154.	4.4	107
4	Text de-identification for privacy protection: A study of its impact on clinical text information content. Journal of Biomedical Informatics, 2014, 50, 142-150.	4.3	44
5	Evaluating the effects of machine pre-annotation and an interactive annotation interface on manual de-identification of clinical text. Journal of Biomedical Informatics, 2014, 50, 162-172.	4.3	40
6	BoB, a best-of-breed automated text de-identification system for VHA clinical documents. Journal of the American Medical Informatics Association: JAMIA, 2013, 20, 77-83.	4.4	60
7	Overview of the ShARe/CLEF eHealth Evaluation Lab 2013. Lecture Notes in Computer Science, 2013, , 212-231.	1.3	127
8	"Sitting on pins and needles": characterization of symptom descriptions in clinical notes". AMIA Summits on Translational Science Proceedings, 2013, 2013, 67-71.	0.4	10
9	Evaluating the state of the art in coreference resolution for electronic medical records. Journal of the American Medical Informatics Association: JAMIA, 2012, 19, 786-791.	4.4	113
10	Automated extraction of ejection fraction for quality measurement using regular expressions in Unstructured Information Management Architecture (UIMA) for heart failure. Journal of the American Medical Informatics Association: JAMIA, 2012, 19, 859-866.	4.4	93
11	Evaluating current automatic de-identification methods with Veteranâ€™s health administration clinical documents. BMC Medical Research Methodology, 2012, 12, 109.	3.1	38
12	The relationship between structural characteristics of 2010 challenge documents and ratings of document quality. AMIA ... Annual Symposium proceedings, 2012, 2012, 848-55.	0.2	0
13	2010 i2b2/VA challenge on concepts, assertions, and relations in clinical text. Journal of the American Medical Informatics Association: JAMIA, 2011, 18, 552-556.	4.4	813
14	Qualitative analysis of workflow modifications used to generate the reference standard for the 2010 i2b2/VA challenge. AMIA ... Annual Symposium proceedings, 2011, 2011, 1243-51.	0.2	2
15	Automatic de-identification of textual documents in the electronic health record: a review of recent research. BMC Medical Research Methodology, 2010, 10, 70.	3.1	217
16	Textractor: a hybrid system for medications and reason for their prescription extraction from clinical text documents. Journal of the American Medical Informatics Association: JAMIA, 2010, 17, 559-562.	4.4	38
17	Analysis of False Positive Errors of an Acute Respiratory Infection Text Classifier due to Contextual Features. Summit on Translational Bioinformatics, 2010, 2010, 56-60.	0.7	2
18	Developing a manually annotated clinical document corpus to identify phenotypic information for inflammatory bowel disease. BMC Bioinformatics, 2009, 10, S12.	2.6	37

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19	Developing a manually annotated clinical document corpus to identify phenotypic information for inflammatory bowel disease. Summit on Translational Bioinformatics, 2009, 2009, 1-32.	0.7	2
20	Optimizing A syndromic surveillance text classifier for influenza-like illness: Does document source matter?. AMIA ... Annual Symposium proceedings, 2008, , 692-6.	0.2	17
21	Application of Natural Language Processing to VA Electronic Health Records to Identify Phenotypic Characteristics for Clinical and Research Purposes. Summit on Translational Bioinformatics, 2008, 2008, 36-40.	0.7	11