

Laura K Wiley

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

1,547
citations

1684129

5
h-index

1720014

7
g-index

20
all docs

20
docs citations

20
times ranked

3094
citing authors

#	ARTICLE	IF	CITATIONS
1	Opportunity for Genotype-Guided Prescribing Among Adult Patients in 11 US Health Systems. <i>Clinical Pharmacology and Therapeutics</i> , 2021, 110, 179-188.	4.7	35
2	Assessing the utility and accuracy of ICD10-CM non-traumatic subarachnoid hemorrhage codes for intracranial aneurysm research. <i>Learning Health Systems</i> , 2021, 5, e10257.	2.0	0
3	Developing and Deploying a Scalable Computing Platform to Support MOOC Education in Clinical Data Science. <i>AMIA Summits on Translational Science Proceedings</i> , 2021, 2021, 430-437.	0.4	0
4	Recommendations for patient similarity classes: results of the AMIA 2019 workshop on defining patient similarity. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2020, 27, 1808-1812.	4.4	15
5	Predictors of 90-Day Readmission Rate After Unruptured Intracranial Aneurysm Repair. <i>World Neurosurgery</i> , 2020, 141, e728-e735.	1.3	1
6	The diversity and disparity in biomedical informatics (DDBI) workshop. , 2018, , .		1
7	Opportunities and obstacles for deep learning in biology and medicine. <i>Journal of the Royal Society Interface</i> , 2018, 15, 20170387.	3.4	1,282
8	STRATEGIES FOR EQUITABLE PHARMACOGENOMIC-GUIDED WARFARIN DOSING AMONG EUROPEAN AND AFRICAN AMERICAN INDIVIDUALS IN A CLINICAL POPULATION. , 2017, 22, 545-556.		6
9	Use of Clinical Decision Support to Tailor Drug Therapy Based on Genomics. <i>Computers in Health Care</i> , 2016, , 209-226.	0.3	0
10	Phenotyping Adverse Drug Reactions: Statin-Related Myotoxicity. <i>AMIA Summits on Translational Science Proceedings</i> , 2015, 2015, 466-70.	0.4	5
11	ICD-9 tobacco use codes are effective identifiers of smoking status. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2013, 20, 652-658.	4.4	127
12	Rapid storage and retrieval of genomic intervals from a relational database system using nested containment lists. <i>Database: the Journal of Biological Databases and Curation</i> , 2013, 2013, bat056.	3.0	1