

Theresa L Walunas

List of Publications by Year in descending order

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46
papers

9,558
citations

270111

25
h-index

299063

42
g-index

50
all docs

50
docs citations

50
times ranked

10789
citing authors

#	ARTICLE	IF	CITATIONS
1	Identifying Contextual Factors and Strategies for Practice Facilitation in Primary Care Quality Improvement Using an Informatics-Driven Model: Framework Development and Mixed Methods Case Study. <i>JMIR Human Factors</i> , 2022, 9, e32174.	1.0	8
2	Metadata Correction: Identifying Contextual Factors and Strategies for Practice Facilitation in Primary Care Quality Improvement Using an Informatics-Driven Model: Framework Development and Mixed Methods Case Study. <i>JMIR Human Factors</i> , 2022, 9, e40674.	1.0	0
3	Remdesivir for Severe Coronavirus Disease 2019 (COVID-19) Versus a Cohort Receiving Standard of Care. <i>Clinical Infectious Diseases</i> , 2021, 73, e4166-e4174.	2.9	135
4	A Taxonomy for External Support for Practice Transformation. <i>Journal of the American Board of Family Medicine</i> , 2021, 34, 32-39.	0.8	6
5	Does coaching matter? Examining the impact of specific practice facilitation strategies on implementation of quality improvement interventions in the Healthy Hearts in the Heartland study. <i>Implementation Science</i> , 2021, 16, 33.	2.5	21
6	Evaluation of structured data from electronic health records to identify clinical classification criteria attributes for systemic lupus erythematosus. <i>Lupus Science and Medicine</i> , 2021, 8, e000488.	1.1	6
7	Remdesivir Versus Standard-of-Care for Severe Coronavirus Disease 2019 Infection: An Analysis of 28-Day Mortality. <i>Open Forum Infectious Diseases</i> , 2021, 8, ofab278.	0.4	31
8	Pleiotropy of systemic lupus erythematosus risk alleles and cardiometabolic disorders: A phenome-wide association study and inverse-variance weighted meta-analysis. <i>Lupus</i> , 2021, 30, 1264-1272.	0.8	2
9	Neptune: an environment for the delivery of genomic medicine. <i>Genetics in Medicine</i> , 2021, 23, 1838-1846.	1.1	3
10	Development of preclinical and clinical models for immune-related adverse events following checkpoint immunotherapy: a perspective from SITC and AACR. , 2021, 9, e002627.		15
11	A Polygenic and Phenotypic Risk Prediction for Polycystic Ovary Syndrome Evaluated by Phenome-Wide Association Studies. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, 1918-1936.	1.8	40
12	Using Electronic Health Records to Measure Quality Improvement Efforts: Findings from a Large Practice Facilitation Initiative. <i>Joint Commission Journal on Quality and Patient Safety</i> , 2020, 46, 11-17.	0.4	6
13	Effects of 2 Forms of Practice Facilitation on Cardiovascular Prevention in Primary Care. <i>Medical Care</i> , 2020, 58, 344-351.	1.1	14
14	Contrasting Perspectives of Practice Leaders and Practice Facilitators May Be Common in Quality Improvement Initiatives. <i>Journal for Healthcare Quality: Official Publication of the National Association for Healthcare Quality</i> , 2020, 42, e32-e38.	0.3	3
15	Identifying Practice Facilitation Delays and Barriers in Primary Care Quality Improvement. <i>Journal of the American Board of Family Medicine</i> , 2020, 33, 655-664.	0.8	32
16	Lung Cancer Survival in Patients With Autoimmune Disease. <i>JAMA Network Open</i> , 2020, 3, e2029917.	2.8	16
17	Making work visible for electronic phenotype implementation: Lessons learned from the eMERGE network. <i>Journal of Biomedical Informatics</i> , 2019, 99, 103293.	2.5	27
18	Challenges to electronic clinical quality measurement using third-party platforms in primary care practices: the healthy hearts in the heartland experience. <i>JAMIA Open</i> , 2019, 2, 423-428.	1.0	8

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19	Harmonizing Clinical Sequencing and Interpretation for the eMERGE III Network. <i>American Journal of Human Genetics</i> , 2019, 105, 588-605.	2.6	99
20	Qualitative evaluation of a cardiovascular quality improvement program reveals sizable data inaccuracies in small primary care practices. <i>BMJ Open Quality</i> , 2019, 8, e000702.	0.4	1
21	Assessing the Concordance of Clinical Classification Criteria for Lupus Between Electronic Health Records and a Physician Curated Registry. <i>Studies in Health Technology and Informatics</i> , 2019, 264, 1466-1467.	0.2	0
22	BD-09...Preliminary report: rule-based algorithms using systemic lupus international collaborating clinics (SLICC) classification criteria to identify patients with systemic lupus erythematosus (SLE) from electronic health record (EHR) data. , 2018, , .		0
23	Practice Facilitators™ and Leaders™ Perspectives on a Facilitated Quality Improvement Program. <i>Annals of Family Medicine</i> , 2018, 16, S65-S71.	0.9	15
24	Engaging Primary Care Practices in Studies of Improvement: Did You Budget Enough for Practice Recruitment?. <i>Annals of Family Medicine</i> , 2018, 16, S72-S79.	0.9	15
25	Design of healthy hearts in the heartland (H3): A practice-randomized, comparative effectiveness study. <i>Contemporary Clinical Trials</i> , 2018, 71, 47-54.	0.8	20
26	Disease Outcomes and Care Fragmentation Among Patients With Systemic Lupus Erythematosus. <i>Arthritis Care and Research</i> , 2017, 69, 1369-1376.	1.5	33
27	Payment Reform Needed to Address Health Disparities of Undiagnosed Diabetic Retinopathy in the City of Chicago. <i>Ophthalmology and Therapy</i> , 2017, 6, 123-131.	1.0	10
28	Design and implementation of a privacy preserving electronic health record linkage tool in Chicago. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2015, 22, 1072-1080.	2.2	101
29	The Genetic Basis of Laboratory Adaptation in <i>Caulobacter crescentus</i> . <i>Journal of Bacteriology</i> , 2010, 192, 3678-3688.	1.0	166
30	Living with Genome Instability: the Adaptation of Phytoplasmata to Diverse Environments of Their Insect and Plant Hosts. <i>Journal of Bacteriology</i> , 2006, 188, 3682-3696.	1.0	356
31	Identification of Open Reading Frames Unique to a Select Agent: <i>Ralstonia solanacearum</i> Race 3 Biovar 2. <i>Molecular Plant-Microbe Interactions</i> , 2006, 19, 69-79.	1.4	121
32	The complete genome sequence of <i>Lactobacillus bulgaricus</i> reveals extensive and ongoing reductive evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 9274-9279.	3.3	382
33	Comparative genome analysis of <i>Bacillus cereus</i> group genomes with <i>Bacillus subtilis</i> . <i>FEMS Microbiology Letters</i> , 2005, 250, 175-184.	0.7	73
34	Genome sequence of <i>Bacillus cereus</i> and comparative analysis with <i>Bacillus anthracis</i> . <i>Nature</i> , 2003, 423, 87-91.	13.7	740
35	Genome Analysis of <i>F. nucleatum</i> sub spp <i>vincentii</i> and Its Comparison With the Genome of <i>F. nucleatum</i> ATCC 25586. <i>Genome Research</i> , 2003, 13, 1180-1189.	2.4	72
36	The ERGOTM genome analysis and discovery system. <i>Nucleic Acids Research</i> , 2003, 31, 164-171.	6.5	207

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37	Whole-genome comparative analysis of three phytopathogenic <i>Xylella fastidiosa</i> strains. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 12403-12408.	3.3	94
38	Draft Sequencing and Comparative Genomics of <i>Xylella fastidiosa</i> Strains Reveal Novel Biological Insights. Genome Research, 2002, 12, 1556-1563.	2.4	70
39	Genome Sequence and Analysis of the Oral Bacterium <i>Fusobacterium nucleatum</i> Strain ATCC 25586. Journal of Bacteriology, 2002, 184, 2005-2018.	1.0	311
40	Cutting Edge: The Ets1 Transcription Factor Is Required for the Development of NK T Cells in Mice. Journal of Immunology, 2000, 164, 2857-2860.	0.4	86
41	The Ets-1 Transcription Factor Is Required for the Development of Natural Killer Cells in Mice. Immunity, 1998, 9, 555-563.	6.6	338
42	CTLA-4 ligation blocks CD28-dependent T cell activation.. Journal of Experimental Medicine, 1996, 183, 2541-2550.	4.2	732
43	CD28/B7 SYSTEM OF T CELL COSTIMULATION. Annual Review of Immunology, 1996, 14, 233-258.	9.5	2,466
44	CTLA-4: a negative regulator of autoimmune disease.. Journal of Experimental Medicine, 1996, 184, 783-788.	4.2	369
45	Absence of B7-dependent responses in CD28-deficient mice. Immunity, 1994, 1, 501-508.	6.6	359
46	CTLA-4 can function as a negative regulator of T cell activation. Immunity, 1994, 1, 405-413.	6.6	1,949