

Theresa L Walunas

List of Publications by Year in descending order

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Version: 2024-02-01

46
papers

9,558
citations

236912

25
h-index

265191

42
g-index

50
all docs

50
docs citations

50
times ranked

9815
citing authors

#	ARTICLE	IF	CITATIONS
1	CD28/B7 SYSTEM OF T CELL COSTIMULATION. Annual Review of Immunology, 1996, 14, 233-258.	21.8	2,466
2	CTLA-4 can function as a negative regulator of T cell activation. Immunity, 1994, 1, 405-413.	14.3	1,949
3	Genome sequence of <i>Bacillus cereus</i> and comparative analysis with <i>Bacillus anthracis</i> . Nature, 2003, 423, 87-91.	27.8	740
4	CTLA-4 ligation blocks CD28-dependent T cell activation.. Journal of Experimental Medicine, 1996, 183, 2541-2550.	8.5	732
5	The complete genome sequence of <i>Lactobacillus bulgaricus</i> reveals extensive and ongoing reductive evolution. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 9274-9279.	7.1	382
6	CTLA-4: a negative regulator of autoimmune disease.. Journal of Experimental Medicine, 1996, 184, 783-788.	8.5	369
7	Absence of B7-dependent responses in CD28-deficient mice. Immunity, 1994, 1, 501-508.	14.3	359
8	Living with Genome Instability: the Adaptation of Phytoplasmas to Diverse Environments of Their Insect and Plant Hosts. Journal of Bacteriology, 2006, 188, 3682-3696.	2.2	356
9	The Ets-1 Transcription Factor Is Required for the Development of Natural Killer Cells in Mice. Immunity, 1998, 9, 555-563.	14.3	338
10	Genome Sequence and Analysis of the Oral Bacterium <i>Fusobacterium nucleatum</i> Strain ATCC 25586. Journal of Bacteriology, 2002, 184, 2005-2018.	2.2	311
11	The ERGOTM genome analysis and discovery system. Nucleic Acids Research, 2003, 31, 164-171.	14.5	207
12	The Genetic Basis of Laboratory Adaptation in <i>Caulobacter crescentus</i> . Journal of Bacteriology, 2010, 192, 3678-3688.	2.2	166
13	Remdesivir for Severe Coronavirus Disease 2019 (COVID-19) Versus a Cohort Receiving Standard of Care. Clinical Infectious Diseases, 2021, 73, e4166-e4174.	5.8	135
14	Identification of Open Reading Frames Unique to a Select Agent: <i>Ralstonia solanacearum</i> Race 3 Biovar 2. Molecular Plant-Microbe Interactions, 2006, 19, 69-79.	2.6	121
15	Design and implementation of a privacy preserving electronic health record linkage tool in Chicago. Journal of the American Medical Informatics Association: JAMIA, 2015, 22, 1072-1080.	4.4	101
16	Harmonizing Clinical Sequencing and Interpretation for the eMERGE III Network. American Journal of Human Genetics, 2019, 105, 588-605.	6.2	99
17	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.	7.1	94
18	Cutting Edge: The Ets1 Transcription Factor Is Required for the Development of NK T Cells in Mice. Journal of Immunology, 2000, 164, 2857-2860.	0.8	86

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19	Comparative genome analysis of <i>Bacillus cereus</i> group genomes with <i>Bacillus subtilis</i> . <i>FEMS Microbiology Letters</i> , 2005, 250, 175-184.	1.8	73
20	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.	5.5	72
21	Draft Sequencing and Comparative Genomics of <i>Xylella fastidiosa</i> Strains Reveal Novel Biological Insights. <i>Genome Research</i> , 2002, 12, 1556-1563.	5.5	70
22	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.	3.6	40
23	Disease Outcomes and Care Fragmentation Among Patients With Systemic Lupus Erythematosus. <i>Arthritis Care and Research</i> , 2017, 69, 1369-1376.	3.4	33
24	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.	1.5	32
25	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.	0.9	31
26	Making work visible for electronic phenotype implementation: Lessons learned from the eMERGE network. <i>Journal of Biomedical Informatics</i> , 2019, 99, 103293.	4.3	27
27	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.	6.9	21
28	Design of healthy hearts in the heartland (H3): A practice-randomized, comparative effectiveness study. <i>Contemporary Clinical Trials</i> , 2018, 71, 47-54.	1.8	20
29	Lung Cancer Survival in Patients With Autoimmune Disease. <i>JAMA Network Open</i> , 2020, 3, e2029917.	5.9	16
30	Practice Facilitators™ and Leaders™ Perspectives on a Facilitated Quality Improvement Program. <i>Annals of Family Medicine</i> , 2018, 16, S65-S71.	1.9	15
31	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.	1.9	15
32	Development of preclinical and clinical models for immune-related adverse events following checkpoint immunotherapy: a perspective from SITC and AACR. , 2021, 9, e002627.		15
33	Effects of 2 Forms of Practice Facilitation on Cardiovascular Prevention in Primary Care. <i>Medical Care</i> , 2020, 58, 344-351.	2.4	14
34	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.	2.3	10
35	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.	2.0	8
36	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.	2.0	8

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37	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.	0.7	6
38	A Taxonomy for External Support for Practice Transformation. <i>Journal of the American Board of Family Medicine</i> , 2021, 34, 32-39.	1.5	6
39	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.	2.7	6
40	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.	0.7	3
41	Neptune: an environment for the delivery of genomic medicine. <i>Genetics in Medicine</i> , 2021, 23, 1838-1846.	2.4	3
42	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.	1.6	2
43	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.	1.1	1
44	BD-09â€¦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
45	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.	0.3	0
46	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.	2.0	0