## Theresa L Walunas

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

8,197 41 50 21 h-index g-index citations papers 8,924 5.16 50 9.2 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
41	Remdesivir for Severe Coronavirus Disease 2019 (COVID-19) Versus a Cohort Receiving Standard of Care. <i>Clinical Infectious Diseases</i> , <b>2021</b> , 73, e4166-e4174	11.6	72
40	Remdesivir Versus Standard-of-Care for Severe Coronavirus Disease 2019 Infection: An Analysis of 28-Day Mortality. <i>Open Forum Infectious Diseases</i> , <b>2021</b> , 8, ofab278	1	8
39	Pleiotropy of systemic lupus erythematosus risk alleles and cardiometabolic disorders: A phenome-wide association study and inverse-variance weighted meta-analysis. <i>Lupus</i> , <b>2021</b> , 30, 1264-12	272	1
38	Neptune: an environment for the delivery of genomic medicine. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 1838-18	<b>48.</b> 1	1
37	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> , <b>2021</b> , 16, 33	8.4	6
36	Development of preclinical and clinical models for immune-related adverse events following checkpoint immunotherapy: a perspective from SITC and AACR <b>2021</b> , 9,		4
35	Identifying Practice Facilitation Delays and Barriers in Primary Care Quality Improvement. <i>Journal of the American Board of Family Medicine</i> , <b>2020</b> , 33, 655-664	1.6	12
34	Lung Cancer Survival in Patients With Autoimmune Disease. <i>JAMA Network Open</i> , <b>2020</b> , 3, e2029917	10.4	5
33	A Polygenic and Phenotypic Risk Prediction for Polycystic Ovary Syndrome Evaluated by Phenome-Wide Association Studies. <i>Journal of Clinical Endocrinology and Metabolism</i> , <b>2020</b> , 105,	5.6	17
32	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> , <b>2020</b> , 46, 11-17	1.4	3
31	Effects of 2 Forms of Practice Facilitation on Cardiovascular Prevention in Primary Care: A Practice-randomized, Comparative Effectiveness Trial. <i>Medical Care</i> , <b>2020</b> , 58, 344-351	3.1	6
30	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> , <b>2020</b> , 42, e32-e38	1	1
29	Harmonizing Clinical Sequencing and Interpretation for the eMERGE III Network. <i>American Journal of Human Genetics</i> , <b>2019</b> , 105, 588-605	11	63
28	Making work visible for electronic phenotype implementation: Lessons learned from the eMERGE network. <i>Journal of Biomedical Informatics</i> , <b>2019</b> , 99, 103293	10.2	11
27	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> , <b>2019</b> , 2, 423-428	2.9	5
26	Qualitative evaluation of a cardiovascular quality improvement programmereveals sizable data inaccuracies in small primary care practices. <i>BMJ Open Quality</i> , <b>2019</b> , 8, e000702	1.9	О
25	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> , <b>2019</b> , 264, 1466	-1457	

## (2000-2018)

24	Engaging Primary Care Practices in Studies of Improvement: Did You Budget Enough for Practice Recruitment?. <i>Annals of Family Medicine</i> , <b>2018</b> , 16, S72-S79	2.9	11	
23	Design of healthy hearts in the heartland (H3): A practice-randomized, comparative effectiveness study. <i>Contemporary Clinical Trials</i> , <b>2018</b> , 71, 47-54	2.3	13	
22	Practice FacilitatorsTand LeadersTPerspectives on a Facilitated Quality Improvement Program. <i>Annals of Family Medicine</i> , <b>2018</b> , 16, S65-S71	2.9	8	
21	Disease Outcomes and Care Fragmentation Among Patients With Systemic Lupus Erythematosus. <i>Arthritis Care and Research</i> , <b>2017</b> , 69, 1369-1376	4.7	18	
20	Payment Reform Needed to Address Health Disparities of Undiagnosed Diabetic Retinopathy in the City of Chicago. <i>Ophthalmology and Therapy</i> , <b>2017</b> , 6, 123-131	5	6	
19	Design and implementation of a privacy preserving electronic health record linkage tool in Chicago.  Journal of the American Medical Informatics Association: JAMIA, 2015, 22, 1072-80	8.6	75	
18	The genetic basis of laboratory adaptation in Caulobacter crescentus. <i>Journal of Bacteriology</i> , <b>2010</b> , 192, 3678-88	3.5	124	
17	Living with genome instability: the adaptation of phytoplasmas to diverse environments of their insect and plant hosts. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 3682-96	3.5	284	
16	Identification of open reading frames unique to a select agent: Ralstonia solanacearum race 3 biovar 2. <i>Molecular Plant-Microbe Interactions</i> , <b>2006</b> , 19, 69-79	3.6	98	
15	The complete genome sequence of Lactobacillus bulgaricus reveals extensive and ongoing reductive evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2006</b> , 103, 9274-9	11.5	318	
14	Comparative genome analysis of Bacillus cereus group genomes with Bacillus subtilis. <i>FEMS Microbiology Letters</i> , <b>2005</b> , 250, 175-84	2.9	64	
13	The ERGO genome analysis and discovery system. <i>Nucleic Acids Research</i> , <b>2003</b> , 31, 164-71	20.1	173	
12	Genome sequence of Bacillus cereus and comparative analysis with Bacillus anthracis. <i>Nature</i> , <b>2003</b> , 423, 87-91	50.4	670	
11	Genome analysis of F. nucleatum sub spp vincentii and its comparison with the genome of F. nucleatum ATCC 25586. <i>Genome Research</i> , <b>2003</b> , 13, 1180-9	9.7	67	
10	Whole-genome comparative analysis of three phytopathogenic Xylella fastidiosa strains. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2002</b> , 99, 12403-8	11.5	81	
9	Draft sequencing and comparative genomics of Xylella fastidiosa strains reveal novel biological insights. <i>Genome Research</i> , <b>2002</b> , 12, 1556-63	9.7	59	
8	Genome sequence and analysis of the oral bacterium Fusobacterium nucleatum strain ATCC 25586. <i>Journal of Bacteriology</i> , <b>2002</b> , 184, 2005-18	3.5	268	
7	Cutting edge: the Ets1 transcription factor is required for the development of NK T cells in mice. <i>Journal of Immunology</i> , <b>2000</b> , 164, 2857-60	5.3	82	

6	The Ets-1 transcription factor is required for the development of natural killer cells in mice. <i>Immunity</i> , <b>1998</b> , 9, 555-63	32.3	298
5	CTLA-4 ligation blocks CD28-dependent T cell activation. <i>Journal of Experimental Medicine</i> , <b>1996</b> , 183, 2541-50	16.6	657
4	CD28/B7 system of T cell costimulation. <i>Annual Review of Immunology</i> , <b>1996</b> , 14, 233-58	34.7	2245
3	CTLA-4: a negative regulator of autoimmune disease. <i>Journal of Experimental Medicine</i> , <b>1996</b> , 184, 783-	-816.6	339
2	Absence of B7-dependent responses in CD28-deficient mice. <i>Immunity</i> , <b>1994</b> , 1, 501-8	32.3	326