

# Karen R Siklosi

## List of Publications by Year in descending order

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Version: 2024-02-01

40  
papers

2,178  
citations

430754

18  
h-index

345118

36  
g-index

41  
all docs

41  
docs citations

41  
times ranked

2942  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparing encounter-based and annualized chronic pseudomonas infection definitions in cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2022, 21, 40-44.	0.3	3
2	Complete CFTR gene sequencing in 5,058 individuals with cystic fibrosis informs variant-specific treatment. <i>Journal of Cystic Fibrosis</i> , 2022, 21, 463-470.	0.3	13
3	Accurate assignment of disease liability to genetic variants using only population data. <i>Genetics in Medicine</i> , 2022, 24, 87-99.	1.1	4
4	Genetic counseling access for parents of newborns who screen positive for cystic fibrosis: Consensus guidelines. <i>Pediatric Pulmonology</i> , 2022, 57, 894-902.	1.0	6
5	<i>C</i> <i>FTR</i> variants are associated with chronic bronchitis in smokers. <i>European Respiratory Journal</i> , 2022, 60, 2101994.	3.1	6
6	Things come in threes: A new complex allele and a novel deletion within the <i>CFTR</i> gene complicate an accurate diagnosis of cystic fibrosis. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2022, 10, e1926.	0.6	2
7	CFTR bearing variant p.Phe312del exhibits function inconsistent with phenotype and negligible response to ivacaftor. <i>JCI Insight</i> , 2022, 7, .	2.3	3
8	DNA sequencing analysis of cystic fibrosis transmembrane conductance regulator gene identifies cystic fibrosis-associated variants in the Severe Asthma Research Program. <i>Pediatric Pulmonology</i> , 2022, 57, 1782-1788.	1.0	3
9	Caution advised in the use of CFTR modulator treatment for individuals harboring specific CFTR variants. <i>Journal of Cystic Fibrosis</i> , 2022, 21, 856-860.	0.3	8
10	The future of cystic fibrosis care: a global perspective. <i>Lancet Respiratory Medicine</i> , the, 2020, 8, 65-124.	5.2	573
11	Genetic Modifiers of Cystic Fibrosis-Related Diabetes Have Extensive Overlap With Type 2 Diabetes and Related Traits. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, 1401-1415.	1.8	34
12	Predictive effects of low birth weight and small for gestational age status on respiratory and nutritional outcomes in cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2020, 19, 888-895.	0.3	2
13	Cystic fibrosis transmembrane conductance regulator function, not TAS2R38 gene haplotypes, predict sinus surgery in children and young adults with cystic fibrosis. <i>International Forum of Allergy and Rhinology</i> , 2020, 10, 748-754.	1.5	7
14	Evaluation of both exonic and intronic variants for effects on RNA splicing allows for accurate assessment of the effectiveness of precision therapies. <i>PLoS Genetics</i> , 2020, 16, e1009100.	1.5	23
15	Title is missing!. , 2020, 16, e1009100.		0
16	Title is missing!. , 2020, 16, e1009100.		0
17	Title is missing!. , 2020, 16, e1009100.		0
18	Title is missing!. , 2020, 16, e1009100.		0

#	ARTICLE	IF	CITATIONS
19	Practice variation of genetic counselor engagement in the cystic fibrosis newborn screenâ€positive diagnostic resolution process. <i>Journal of Genetic Counseling</i> , 2019, 28, 1178-1188.	0.9	8
20	Correlating Cystic Fibrosis Transmembrane Conductance Regulator Function with Clinical Features to Inform Precision Treatment of Cystic Fibrosis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2019, 199, 1116-1126.	2.5	76
21	The increasing challenge of genetic counseling for cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2019, 18, 167-174.	0.3	17
22	Capitalizing on the heterogeneous effects of CFTR nonsense and frameshift variants to inform therapeutic strategy for cystic fibrosis. <i>PLoS Genetics</i> , 2018, 14, e1007723.	1.5	44
23	Functional Assays Are Essential for Interpretation of Missense Variants Associated with Variable Expressivity. <i>American Journal of Human Genetics</i> , 2018, 102, 1062-1077.	2.6	69
24	Residual function of cystic fibrosis mutants predicts response to small molecule CFTR modulators. <i>JCI Insight</i> , 2018, 3, .	2.3	86
25	Applying Cystic Fibrosis Transmembrane Conductance Regulator Genetics and CFTR2 Data to Facilitate Diagnoses. <i>Journal of Pediatrics</i> , 2017, 181, S27-S32.e1.	0.9	58
26	Ethnicity impacts the cystic fibrosis diagnosis: A note of caution. <i>Journal of Cystic Fibrosis</i> , 2017, 16, 488-491.	0.3	34
27	Diagnosis and Treatment of Cystic Fibrosis: A (Not-so) Simple Recessive Condition. <i>Current Genetic Medicine Reports</i> , 2017, 5, 91-99.	1.9	1
28	Systematic Computational Identification of Variants That Activate Exonic and Intronic Cryptic Splice Sites. <i>American Journal of Human Genetics</i> , 2017, 100, 751-765.	2.6	68
29	Benign and Deleterious Cystic Fibrosis Transmembrane Conductance Regulator Mutations Identified by Sequencing in Positive Cystic Fibrosis Newborn Screen Children from California. <i>PLoS ONE</i> , 2016, 11, e0155624.	1.1	27
30	Deep resequencing of CFTR in 762 F508del homozygotes reveals clusters of non-coding variants associated with cystic fibrosis disease traits. <i>Human Genome Variation</i> , 2016, 3, 16038.	0.4	34
31	Respiratory pathogens mediate the association between lung function and temperature in cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2016, 15, 794-801.	0.3	14
32	Molecular Genetics of Cystic Fibrosis Transmembrane Conductance Regulator. <i>Pediatric Clinics of North America</i> , 2016, 63, 585-598.	0.9	38
33	Bias in CFTR screening panels. <i>Genetics in Medicine</i> , 2016, 18, 209-209.	1.1	3
34	Loss of carbonic anhydrase XII function in individuals with elevated sweat chloride concentration and pulmonary airway disease. <i>Human Molecular Genetics</i> , 2016, 25, 1923-1933.	1.4	32
35	Benign outcome among positive cystic fibrosis newborn screen children with non-CF-causing variants. <i>Journal of Cystic Fibrosis</i> , 2015, 14, 714-719.	0.3	35
36	Missense variants in CFTR nucleotide-binding domains predict quantitative phenotypes associated with cystic fibrosis disease severity. <i>Human Molecular Genetics</i> , 2015, 24, 1908-1917.	1.4	11

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37	Genome-wide association meta-analysis identifies five modifier loci of lung disease severity in cystic fibrosis. <i>Nature Communications</i> , 2015, 6, 8382.	5.8	242
38	Experimental Assessment of Splicing Variants Using Expression Minigenes and Comparison with In Silico Predictions. <i>Human Mutation</i> , 2014, 35, 1249-1259.	1.1	56
39	Defining the disease liability of variants in the cystic fibrosis transmembrane conductance regulator gene. <i>Nature Genetics</i> , 2013, 45, 1160-1167.	9.4	513
40	Development, validation, and implementation of a questionnaire assessing disease knowledge and understanding in adult cystic fibrosis patients. <i>Journal of Cystic Fibrosis</i> , 2010, 9, 400-405.	0.3	24