

# Batsheva Kerem

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/345534/publications.pdf>

Version: 2024-02-01

46  
papers

4,894  
citations

186265

28  
h-index

223800

46  
g-index

50  
all docs

50  
docs citations

50  
times ranked

5814  
citing authors

#	ARTICLE	IF	CITATIONS
1	Nucleotide Deficiency Promotes Genomic Instability in Early Stages of Cancer Development. <i>Cell</i> , 2011, 145, 435-446.	28.9	685
2	Gentamicin-Induced Correction of CFTR Function in Patients with Cystic Fibrosis and CFTR Stop Mutations. <i>New England Journal of Medicine</i> , 2003, 349, 1433-1441.	27.0	473
3	Effectiveness of PTC124 treatment of cystic fibrosis caused by nonsense mutations: a prospective phase II trial. <i>Lancet</i> , 2008, 372, 719-727.	13.7	364
4	A role for common fragile site induction in amplification of human oncogenes. <i>Cancer Cell</i> , 2002, 1, 89-97.	16.8	267
5	Nonsense-mediated mRNA decay affects nonsense transcript levels and governs response of cystic fibrosis patients to gentamicin. <i>Journal of Clinical Investigation</i> , 2007, 117, 683-692.	8.2	252
6	The presence of extra chromosomes leads to genomic instability. <i>Nature Communications</i> , 2016, 7, 10754.	12.8	235
7	Molecular Basis for Expression of Common and Rare Fragile Sites. <i>Molecular and Cellular Biology</i> , 2003, 23, 7143-7151.	2.3	211
8	A Pilot Study of the Effect of Gentamicin on Nasal Potential Difference Measurements in Cystic Fibrosis Patients Carrying Stop Mutations. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2000, 161, 860-865.	5.6	202
9	Splicing regulation as a potential genetic modifier. <i>Trends in Genetics</i> , 2002, 18, 123-127.	6.7	197
10	Failure of Origin Activation in Response to Fork Stalling Leads to Chromosomal Instability at Fragile Sites. <i>Molecular Cell</i> , 2011, 43, 122-131.	9.7	157
11	The molecular basis of common and rare fragile sites. <i>Cancer Letters</i> , 2006, 232, 13-26.	7.2	131
12	Homologous recombination and nonhomologous end-joining repair pathways regulate fragile site stability. <i>Genes and Development</i> , 2005, 19, 2715-2726.	5.9	114
13	Replication Delay along FRA7H, a Common Fragile Site on Human Chromosome 7, Leads to Chromosomal Instability. <i>Molecular and Cellular Biology</i> , 2000, 20, 4420-4427.	2.3	110
14	Clinical and Genetic Risk Factors for Cystic Fibrosis-related Liver Disease. <i>Pediatrics</i> , 1999, 103, 52-57.	2.1	108
15	Genomic Instability in Human Pluripotent Stem Cells Arises from Replicative Stress and Chromosome Condensation Defects. <i>Cell Stem Cell</i> , 2016, 18, 253-261.	11.1	106
16	Genotype-phenotype correlations in cystic fibrosis. <i>Am J Hum Genet</i> , 1996, 22, 387-395.		102
17	The efficiency of nonsense-mediated mRNA decay is an inherent character and varies among different cells. <i>European Journal of Human Genetics</i> , 2007, 15, 1156-1162.	2.8	102
18	The Molecular Basis for Disease Variability in Cystic Fibrosis. <i>European Journal of Human Genetics</i> , 1996, 4, 65-73.	2.8	101

#	ARTICLE	IF	CITATIONS
19	The Molecular Basis of Disease Variability among Cystic Fibrosis Patients Carrying the 3849+10 kb CâtT Mutation. <i>Genomics</i> , 1998, 53, 276-283.	2.9	87
20	The splicing machinery is a genetic modifier of disease severity. <i>Trends in Genetics</i> , 2005, 21, 480-483.	6.7	72
21	The complex basis underlying common fragile site instability in cancer. <i>Trends in Genetics</i> , 2012, 28, 295-302.	6.7	72
22	Oncogenes create a unique landscape of fragile sites. <i>Nature Communications</i> , 2015, 6, 7094.	12.8	72
23	Restoration of the cystic fibrosis transmembrane conductance regulator function by splicing modulation. <i>EMBO Reports</i> , 2004, 5, 1071-1077.	4.5	65
24	Identification of Dormancy-Associated MicroRNAs for the Design of Osteosarcoma-Targeted Dendritic Polyglycerol Nanopolyplexes. <i>ACS Nano</i> , 2016, 10, 2028-2045.	14.6	64
25	Highly variable incidence of cystic fibrosis and different mutation distribution among different Jewish ethnic groups in Israel. <i>Human Genetics</i> , 1995, 96, 193-197.	3.8	63
26	The complex nature of fragile site plasticity and its importance in cancer. <i>Current Opinion in Cell Biology</i> , 2016, 40, 131-136.	5.4	56
27	3D genome organization contributes to genome instability at fragile sites. <i>Nature Communications</i> , 2020, 11, 3613.	12.8	46
28	New approaches to genetic therapies for cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2020, 19, S54-S59.	0.7	46
29	The unfolded protein response affects readthrough of premature termination codons. <i>EMBO Molecular Medicine</i> , 2014, 6, 685-701.	6.9	31
30	The suppression of premature termination codons and the repair of splicing mutations in CFTR. <i>Current Opinion in Pharmacology</i> , 2017, 34, 125-131.	3.5	30
31	Antisense oligonucleotide-based drug development for Cystic Fibrosis patients carrying the 3849+10Åkb C-to-T splicing mutation. <i>Journal of Cystic Fibrosis</i> , 2021, 20, 865-875.	0.7	30
32	Folate levels modulate oncogeneâ€”induced replication stress and tumorigenicity. <i>EMBO Molecular Medicine</i> , 2015, 7, 1138-1152.	6.9	28
33	AT-dinucleotide rich sequences drive fragile site formation. <i>Nucleic Acids Research</i> , 2019, 47, 9685-9695.	14.5	28
34	Interplay between genetic and epigenetic factors governs common fragile site instability in cancer. <i>Cellular and Molecular Life Sciences</i> , 2014, 71, 4495-4506.	5.4	26
35	FRA18C: a new aphidicolin-inducible fragile site on chromosome 18q22, possibly associated with in vivo chromosome breakage. <i>Journal of Medical Genetics</i> , 2007, 44, 347-352.	3.2	24
36	Transcriptional Dynamics in Colorectal Carcinogenesis: New Insights into the Role of c-Myc and miR17 in Benign to Cancer Transformation. <i>Cancer Research</i> , 2014, 74, 5532-5540.	0.9	19

#	ARTICLE	IF	CITATIONS
37	DNA replication stress drives fragile site instability. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2018, 808, 56-61.	1.0	19
38	Impaired Replication Stress Response in Cells from Immunodeficiency Patients Carrying Cernunnos/XLF Mutations. <i>PLoS ONE</i> , 2009, 4, e4516.	2.5	19
39	Oncogene-Induced Replication Stress Drives Genome Instability and Tumorigenesis. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1339.	4.1	18
40	Antisense oligonucleotide splicing modulation as a novel Cystic Fibrosis therapeutic approach for the W1282X nonsense mutation. <i>Journal of Cystic Fibrosis</i> , 2022, 21, 630-636.	0.7	18
41	Genomic instability in fragile sitesâ€”still adding the pieces. <i>Genes Chromosomes and Cancer</i> , 2019, 58, 295-304.	2.8	15
42	Screening of CFTR mutations in an isolated population: identification of carriers and patients. <i>European Journal of Human Genetics</i> , 1998, 6, 181-184.	2.8	11
43	Genotypeâ€”phenotype correlations in cystic fibrosis. <i>Pediatric Pulmonology</i> , 1996, 22, 387-395.	2.0	4
44	Continuous chromosomal instability in human pluripotent stem cells â€” the role of DNA replication. <i>Molecular and Cellular Oncology</i> , 2016, 3, e1183743.	0.7	3
45	Take it personally: how personal we reach when we are so different from each other?. <i>Journal of Cystic Fibrosis</i> , 2019, 18, 6-7.	0.7	3
46	To break or not to break â€” context matters. <i>Molecular and Cellular Oncology</i> , 2016, 3, e1072657.	0.7	1