

James B Stewart

List of Publications by Year in descending order

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Version: 2024-02-01

52
papers

5,075
citations

109321

35
h-index

206112

48
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60
all docs

60
docs citations

60
times ranked

5645
citing authors

#	ARTICLE	IF	CITATIONS
1	The dynamics of mitochondrial DNA heteroplasmy: implications for human health and disease. <i>Nature Reviews Genetics</i> , 2015, 16, 530-542.	16.3	679
2	Incorporating Molecular Evolution into Phylogenetic Analysis, and a New Compilation of Conserved Polymerase Chain Reaction Primers for Animal Mitochondrial DNA. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2006, 37, 545-579.	8.3	496
3	Strong Purifying Selection in Transmission of Mammalian Mitochondrial DNA. <i>PLoS Biology</i> , 2008, 6, e10.	5.6	425
4	LRPPRC is necessary for polyadenylation and coordination of translation of mitochondrial mRNAs. <i>EMBO Journal</i> , 2012, 31, 443-456.	7.8	264
5	Germline mitochondrial DNA mutations aggravate ageing and can impair brain development. <i>Nature</i> , 2013, 501, 412-415.	27.8	231
6	Genome editing in mitochondria corrects a pathogenic mtDNA mutation in vivo. <i>Nature Medicine</i> , 2018, 24, 1691-1695.	30.7	215
7	MitoTALEN reduces mutant mtDNA load and restores tRNA ^{Ala} levels in a mouse model of heteroplasmic mtDNA mutation. <i>Nature Medicine</i> , 2018, 24, 1696-1700.	30.7	187
8	Ultra-Deep Sequencing of Mouse Mitochondrial DNA: Mutational Patterns and Their Origins. <i>PLoS Genetics</i> , 2011, 7, e1002028.	3.5	162
9	Purifying selection of mtDNA and its implications for understanding evolution and mitochondrial disease. <i>Nature Reviews Genetics</i> , 2008, 9, 657-662.	16.3	155
10	Extreme heterogeneity of human mitochondrial DNA from organelles to populations. <i>Nature Reviews Genetics</i> , 2021, 22, 106-118.	16.3	139
11	Variation in germline mtDNA heteroplasmy is determined prenatally but modified during subsequent transmission. <i>Nature Genetics</i> , 2012, 44, 1282-1285.	21.4	128
12	Hierarchical RNA Processing Is Required for Mitochondrial Ribosome Assembly. <i>Cell Reports</i> , 2016, 16, 1874-1890.	6.4	116
13	Simultaneous DNA and RNA Mapping of Somatic Mitochondrial Mutations across Diverse Human Cancers. <i>PLoS Genetics</i> , 2015, 11, e1005333.	3.5	102
14	A Phenotype-Driven Approach to Generate Mouse Models with Pathogenic mtDNA Mutations Causing Mitochondrial Disease. <i>Cell Reports</i> , 2016, 16, 2980-2990.	6.4	102
15	Characterization of mature mitochondrial transcripts in <i>Drosophila</i> , and the implications for the tRNA punctuation model in arthropods. <i>Gene</i> , 2009, 445, 49-57.	2.2	94
16	Insect mitochondrial genomics: the complete mitochondrial genome sequence of the meadow spittlebug <i>Philaenus spumarius</i> (Hemiptera: Auchenorrhyncha: Cercopoidae). <i>Genome</i> , 2005, 48, 46-54.	2.0	93
17	MTERF1 Binds mtDNA to Prevent Transcriptional Interference at the Light-Strand Promoter but Is Dispensable for rRNA Gene Transcription Regulation. <i>Cell Metabolism</i> , 2013, 17, 618-626.	16.2	93
18	No recombination of mtDNA after heteroplasmy for 50 generations in the mouse maternal germline. <i>Nucleic Acids Research</i> , 2014, 42, 1111-1116.	14.5	92

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19	Keeping mtDNA in Shape between Generations. <i>PLoS Genetics</i> , 2014, 10, e1004670.	3.5	90
20	Modulation of mtDNA copy number ameliorates the pathological consequences of a heteroplasmic mtDNA mutation in the mouse. <i>Science Advances</i> , 2019, 5, eaav9824.	10.3	86
21	Progressive loss of mitochondrial DNA in thymidine kinase 2-deficient mice. <i>Human Molecular Genetics</i> , 2008, 17, 2329-2335.	2.9	85
22	Increased Total mtDNA Copy Number Cures Male Infertility Despite Unaltered mtDNA Mutation Load. <i>Cell Metabolism</i> , 2017, 26, 429-436.e4.	16.2	84
23	Phylogenetic and genomic analysis of the complete mitochondrial DNA sequence of the spotted asparagus beetle <i>Crioceris duodecimpunctata</i> . <i>Molecular Phylogenetics and Evolution</i> , 2003, 26, 513-526.	2.7	70
24	The exonuclease activity of DNA polymerase β is required for ligation during mitochondrial DNA replication. <i>Nature Communications</i> , 2015, 6, 7303.	12.8	70
25	Mitochondrial DNA deletions are associated with non-B DNA conformations. <i>Nucleic Acids Research</i> , 2012, 40, 7606-7621.	14.5	64
26	The Challenges of Mitochondrial Replacement. <i>PLoS Genetics</i> , 2014, 10, e1004315.	3.5	61
27	<i>In vivo</i> mutagenesis reveals that OriL is essential for mitochondrial DNA replication. <i>EMBO Reports</i> , 2012, 13, 1130-1137.	4.5	59
28	Base-excision repair deficiency alone or combined with increased oxidative stress does not increase mtDNA point mutations in mice. <i>Nucleic Acids Research</i> , 2018, 46, 6642-6669.	14.5	58
29	Mice lacking the mitochondrial exonuclease MGME1 accumulate mtDNA deletions without developing progeria. <i>Nature Communications</i> , 2018, 9, 1202.	12.8	57
30	Insect mitochondrial genomics 3: the complete mitochondrial genome sequences of representatives from two neuropteroid orders: a dobsonfly (order Megaloptera) and a giant lacewing and an owlfly (order Neuroptera). <i>Genome</i> , 2009, 52, 31-38.	2.0	56
31	Insect mitochondrial genomics 2: the complete mitochondrial genome sequence of a giant stonefly, <i>Pteronarcys princeps</i> , asymmetric directional mutation bias, and conserved plecopteran A+T-region elements. <i>Genome</i> , 2006, 49, 815-824.	2.0	55
32	The Bicoid Stability Factor Controls Polyadenylation and Expression of Specific Mitochondrial mRNAs in <i>Drosophila melanogaster</i> . <i>PLoS Genetics</i> , 2011, 7, e1002324.	3.5	55
33	Mitochondrial DNA: Radically free of free-radical driven mutations. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2015, 1847, 1354-1361.	1.0	52
34	Complementation between polymerase- and exonuclease-deficient mitochondrial DNA polymerase mutants in genomically engineered flies. <i>Nature Communications</i> , 2015, 6, 8808.	12.8	48
35	Mitochondrial targeted meganuclease as a platform to eliminate mutant mtDNA in vivo. <i>Nature Communications</i> , 2021, 12, 3210.	12.8	42
36	Accurate mapping of mitochondrial DNA deletions and duplications using deep sequencing. <i>PLoS Genetics</i> , 2020, 16, e1009242.	3.5	41

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37	Similar patterns of clonally expanded somatic mtDNA mutations in the colon of heterozygous mtDNA mutator mice and ageing humans. <i>Mechanisms of Ageing and Development</i> , 2014, 139, 22-30.	4.6	33
38	Mitochondrial stress response triggered by defects in protein synthesis quality control. <i>Life Science Alliance</i> , 2019, 2, e201800219.	2.8	26
39	Mitochondrial DNA heteroplasmy is modulated during oocyte development propagating mutation transmission. <i>Science Advances</i> , 2021, 7, eabi5657.	10.3	22
40	Current progress with mammalian models of mitochondrial DNA disease. <i>Journal of Inherited Metabolic Disease</i> , 2021, 44, 325-342.	3.6	19
41	A novel histochemistry assay to assess and quantify focal cytochrome <i>c</i> oxidase deficiency. <i>Journal of Pathology</i> , 2018, 245, 311-323.	4.5	17
42	Large dataset of octocoral mitochondrial genomes provides new insights into mt-mutS evolution and function. <i>DNA Repair</i> , 2022, 110, 103273.	2.8	16
43	Tissue-specific modulation of mitochondrial DNA segregation by a defect in mitochondrial division. <i>Human Molecular Genetics</i> , 2016, 25, 706-714.	2.9	11
44	Delivery of mtZFNs into Early Mouse Embryos. <i>Methods in Molecular Biology</i> , 2018, 1867, 215-228.	0.9	6
45	Addressing RNA Integrity to Determine the Impact of Mitochondrial DNA Mutations on Brain Mitochondrial Function with Age. <i>PLoS ONE</i> , 2014, 9, e96940.	2.5	5
46	High-Throughput Detection of mtDNA Mutations Leading to tRNA Processing Errors. <i>Methods in Molecular Biology</i> , 2021, 2192, 117-132.	0.9	4
47	Characterization of the sea urchin mitochondrial transcription factor A reveals unusual features. <i>Mitochondrion</i> , 2014, 14, 34-41.	3.4	2
48	MitoTALEN reduces mutant mtDNA load and restores tRNA ^A levels in a mouse model of heteroplasmic mtDNA mutation. , 0, .		1
49	Accurate mapping of mitochondrial DNA deletions and duplications using deep sequencing. , 2020, 16, e1009242.		0
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