

# Howard T Jacobs, Howy Jacobs

## List of Publications by Year in descending order

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250  
papers

13,915  
citations

36303

51  
h-index

22832

112  
g-index

265  
all docs

265  
docs citations

265  
times ranked

13293  
citing authors

#	ARTICLE	IF	CITATIONS
1	Premature ageing in mice expressing defective mitochondrial DNA polymerase. <i>Nature</i> , 2004, 429, 417-423.	27.8	2,318
2	Succinate Dehydrogenase Supports Metabolic Repurposing of Mitochondria to Drive Inflammatory Macrophages. <i>Cell</i> , 2016, 167, 457-470.e13.	28.9	1,396
3	Human mitochondrial DNA deletions associated with mutations in the gene encoding Twinkle, a phage T7 gene 4-like protein localized in mitochondria. <i>Nature Genetics</i> , 2001, 28, 223-231.	21.4	803
4	Somatic mtDNA mutations cause aging phenotypes without affecting reactive oxygen species production. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 17993-17998.	7.1	491
5	Coupled Leading- and Lagging-Strand Synthesis of Mammalian Mitochondrial DNA. <i>Cell</i> , 2000, 100, 515-524.	28.9	372
6	Nucleotide sequence and gene organization of sea urchin mitochondrial DNA. <i>Journal of Molecular Biology</i> , 1988, 202, 185-217.	4.2	351
7	Mitochondria are physiologically maintained at close to 50 Å°C. <i>PLoS Biology</i> , 2018, 16, e2003992.	5.6	295
8	A novel mitochondrial point mutation in a maternal pedigree with sensorineural deafness. <i>Human Mutation</i> , 1994, 3, 243-247.	2.5	252
9	Biased Incorporation of Ribonucleotides on the Mitochondrial L-Strand Accounts for Apparent Strand-Asymmetric DNA Replication. <i>Cell</i> , 2002, 111, 495-505.	28.9	238
10	Mitochondrial medicine: A metabolic perspective on the pathology of oxidative phosphorylation disorders. <i>Cell Metabolism</i> , 2006, 3, 9-13.	16.2	210
11	Replication of vertebrate mitochondrial DNA entails transient ribonucleotide incorporation throughout the lagging strand. <i>EMBO Journal</i> , 2006, 25, 5358-5371.	7.8	205
12	Different cellular backgrounds confer a marked advantage to either mutant or wild-type mitochondrial genomes.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1995, 92, 6562-6566.	7.1	198
13	Mammalian Mitochondrial DNA Replicates Bidirectionally from an Initiation Zone. <i>Journal of Biological Chemistry</i> , 2003, 278, 50961-50969.	3.4	174
14	Mutations at the mitochondrial DNA polymerase (POLG) locus associated with male infertility. <i>Nature Genetics</i> , 2001, 29, 261-262.	21.4	173
15	Human mitochondrial transcription factor A induces a U-turn structure in the light strand promoter. <i>Nature Structural and Molecular Biology</i> , 2011, 18, 1281-1289.	8.2	168
16	In Vivo Functional Analysis of the Human Mitochondrial DNA Polymerase POLG Expressed in Cultured Human Cells. <i>Journal of Biological Chemistry</i> , 2000, 275, 24818-24828.	3.4	166
17	Expression of the <i>Ciona intestinalis</i> Alternative Oxidase (AOX) in <i>Drosophila</i> Complements Defects in Mitochondrial Oxidative Phosphorylation. <i>Cell Metabolism</i> , 2009, 9, 449-460.	16.2	156
18	Stabilization of Hypoxia-inducible Factor-1 $\alpha$ Protein in Hypoxia Occurs Independently of Mitochondrial Reactive Oxygen Species Production. <i>Journal of Biological Chemistry</i> , 2010, 285, 31277-31284.	3.4	154

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19	Alterations to the expression level of mitochondrial transcription factor A, TFAM, modify the mode of mitochondrial DNA replication in cultured human cells. <i>Nucleic Acids Research</i> , 2006, 34, 5815-5828.	14.5	151
20	A Bidirectional Origin of Replication Maps to the Major Noncoding Region of Human Mitochondrial DNA. <i>Molecular Cell</i> , 2005, 18, 651-662.	9.7	148
21	Eukaryotic gene expression: Very short repeats and coordinate induction of genes. <i>Nature</i> , 1983, 301, 468-470.	27.8	135
22	Expression of the yeast NADH dehydrogenase Ndi1 in <i>Drosophila</i> confers increased lifespan independently of dietary restriction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 9105-9110.	7.1	132
23	Mitochondrial DNA sequences in the nuclear genome of <i>Strongylocentrotus pupuratus</i> . <i>Journal of Molecular Biology</i> , 1983, 165, 609-632.	4.2	115
24	Mitochondrial DNA mutations in patients with postlingual, nonsyndromic hearing impairment. <i>European Journal of Human Genetics</i> , 2005, 13, 26-33.	2.8	110
25	Allotopic expression of a mitochondrial alternative oxidase confers cyanide resistance to human cell respiration. <i>EMBO Reports</i> , 2006, 7, 341-345.	4.5	110
26	Human Heart Mitochondrial DNA Is Organized in Complex Catenated Networks Containing Abundant Four-way Junctions and Replication Forks. <i>Journal of Biological Chemistry</i> , 2009, 284, 21446-21457.	3.4	110
27	Mammalian Mitochondrial DNA Replication Intermediates Are Essentially Duplex but Contain Extensive Tracts of RNA/DNA Hybrid. <i>Journal of Molecular Biology</i> , 2010, 397, 1144-1155.	4.2	110
28	Nuclear genes and mitochondrial translation: a new class of genetic disease. <i>Trends in Genetics</i> , 2005, 21, 312-314.	6.7	109
29	Mitochondrial electron transport chain is necessary for NLRP3 inflammasome activation. <i>Nature Immunology</i> , 2022, 23, 692-704.	14.5	107
30	The mitochondrial theory of aging: dead or alive?. <i>Aging Cell</i> , 2003, 2, 11-17.	6.7	106
31	A tRNA suppressor mutation in human mitochondria. <i>Nature Genetics</i> , 1998, 18, 350-353.	21.4	103
32	A Mitochondrial Ribosomal and RNA Decay Pathway Blocks Cell Proliferation. <i>Current Biology</i> , 2013, 23, 535-541.	3.9	103
33	Mitochondrial ROS production correlates with, but does not directly regulate lifespan in <i>Drosophila</i> . <i>Aging</i> , 2010, 2, 200-223.	3.1	101
34	Alternative Oxidase Expression in the Mouse Enables Bypassing Cytochrome c Oxidase Blockade and Limits Mitochondrial ROS Overproduction. <i>PLoS Genetics</i> , 2013, 9, e1003182.	3.5	96
35	No Influence of Indy on Lifespan in <i>Drosophila</i> after Correction for Genetic and Cytoplasmic Background Effects. <i>PLoS Genetics</i> , 2007, 3, e95.	3.5	95
36	The mitochondrial transcription termination factor mTERF modulates replication pausing in human mitochondrial DNA. <i>Nucleic Acids Research</i> , 2007, 35, 6458-6474.	14.5	95

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37	Mitochondrial DNA replication proceeds via a "bootlace"™ mechanism involving the incorporation of processed transcripts. <i>Nucleic Acids Research</i> , 2013, 41, 5837-5850.	14.5	93
38	Prominent mitochondrial DNA recombination intermediates in human heart muscle. <i>EMBO Reports</i> , 2001, 2, 1007-1012.	4.5	92
39	Behaviour of a population of partially duplicated mitochondrial DNA molecules in cell culture: segregation, maintenance and recombination dependent upon nuclear background. <i>Human Molecular Genetics</i> , 1997, 6, 1251-1260.	2.9	91
40	Expression of the alternative oxidase complements cytochrome <i>c</i> oxidase deficiency in human cells. <i>EMBO Molecular Medicine</i> , 2009, 1, 30-36.	6.9	89
41	No sex please, we're mitochondria: a hypothesis on the somatic unit of inheritance of mammalian mtDNA. <i>BioEssays</i> , 2000, 22, 564-572.	2.5	88
42	<i>c</i> technical knockout, a <i>Drosophila</i> Model of Mitochondrial Deafness. <i>Genetics</i> , 2001, 159, 241-254.	2.9	88
43	Disorders of mitochondrial protein synthesis. <i>Human Molecular Genetics</i> , 2003, 12, R293-R301.	2.9	84
44	Primer retention owing to the absence of RNase H1 is catastrophic for mitochondrial DNA replication. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 9334-9339.	7.1	75
45	Perturbed Redox Signaling Exacerbates a Mitochondrial Myopathy. <i>Cell Metabolism</i> , 2018, 28, 764-775.e5.	16.2	70
46	Wobble modification deficiency in mutant tRNAs in patients with mitochondrial diseases. <i>FEBS Letters</i> , 2005, 579, 2948-2952.	2.8	68
47	Gene Expression in a <i>Drosophila</i> Model of Mitochondrial Disease. <i>PLoS ONE</i> , 2010, 5, e8549.	2.5	64
48	Response: The mitochondrial DNA replication bubble has not burst. <i>Trends in Biochemical Sciences</i> , 2003, 28, 355-356.	7.5	62
49	Conserved tRNA gene cluster in starfish mitochondrial DNA. <i>Current Genetics</i> , 1989, 15, 193-206.	1.7	57
50	Mitochondrial DNA "all things bad?". <i>Trends in Genetics</i> , 1999, 15, 91-93.	6.7	57
51	Expression of alternative oxidase in <i>Drosophila</i> ameliorates diverse phenotypes due to cytochrome oxidase deficiency. <i>Human Molecular Genetics</i> , 2014, 23, 2078-2093.	2.9	57
52	Genotypic Stability, Segregation and Selection in Heteroplasmic Human Cell Lines Containing np 3243 Mutant mtDNA. <i>Genetics</i> , 2000, 154, 363-380.	2.9	57
53	Alternative oxidase-mediated respiration prevents lethal mitochondrial cardiomyopathy. <i>EMBO Molecular Medicine</i> , 2019, 11, .	6.9	53
54	Alternative oxidase rescues mitochondria-mediated dopaminergic cell loss in <i>Drosophila</i> . <i>Human Molecular Genetics</i> , 2012, 21, 2698-2712.	2.9	51

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55	Molecular Phenotype of a Human Lymphoblastoid Cell-line Homoplasmic for the np 7445 Deafness-associated Mitochondrial Mutation. <i>Human Molecular Genetics</i> , 1997, 6, 443-449.	2.9	49
56	Engineering the alternative oxidase gene to better understand and counteract mitochondrial defects: state of the art and perspectives. <i>British Journal of Pharmacology</i> , 2014, 171, 2243-2249.	5.4	49
57	The 7472insC Mitochondrial DNA Mutation Impairs the Synthesis and Extent of Aminoacylation of tRNASer(UCN) but Not Its Structure or Rate of Turnover. <i>Journal of Biological Chemistry</i> , 2002, 277, 22240-22250.	3.4	47
58	Sea urchin egg mitochondrial DNA contains a short displacement loop (D-loop) in the replication origin region. <i>Nucleic Acids Research</i> , 1989, 17, 8949-8965.	14.5	46
59	The human homologue of the yeast mitochondrial AAA metalloprotease Yme1p complements a yeast yme1 disruptant. <i>FEBS Letters</i> , 2000, 478, 267-270.	2.8	46
60	Novel Repetitive Structures, Deviant Protein-Encoding Sequences and Unidentified ORFs in the Mitochondrial Genome of the Brachiopod <i>Lingula anatina</i> . <i>Journal of Molecular Evolution</i> , 2005, 61, 36-53.	1.8	46
61	Human mitochondrial mTERF wraps around DNA through a left-handed superhelical tandem repeat. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 891-893.	8.2	46
62	Glucose Modulates Respiratory Complex I Activity in Response to Acute Mitochondrial Dysfunction. <i>Journal of Biological Chemistry</i> , 2012, 287, 38729-38740.	3.4	46
63	Broad AOX expression in a genetically tractable mouse model does not disturb normal physiology. <i>DMM Disease Models and Mechanisms</i> , 2017, 10, 163-171.	2.4	46
64	Molecular Phenotype of the np 7472 Deafness-Associated Mitochondrial Mutation in Osteosarcoma Cell Cybrids. <i>Human Molecular Genetics</i> , 1999, 8, 2275-2283.	2.9	44
65	Developmental and Pathological Changes in the Human Cardiac Muscle Mitochondrial DNA Organization, Replication and Copy Number. <i>PLoS ONE</i> , 2010, 5, e10426.	2.5	43
66	Metazoan nuclear genes for mitoribosomal protein S12. <i>Gene</i> , 1997, 204, 55-62.	2.2	42
67	Complete nucleotide sequences of the nuclear pseudogenes for cytochrome oxidase subunit I and the large mitochondrial ribosomal RNA in the sea urchin <i>Strongylocentrotus purpuratus</i> . <i>Journal of Molecular Biology</i> , 1986, 187, 509-527.	4.2	41
68	Mitochondrial Deafness. <i>Annals of Medicine</i> , 1997, 29, 483-491.	3.8	41
69	Respiratory chain alternative enzymes as tools to better understand and counteract respiratory chain deficiencies in human cells and animals. <i>Physiologia Plantarum</i> , 2009, 137, 362-370.	5.2	41
70	Familial mitochondrial DNA depletion in liver: haplotype analysis of candidate genes. <i>Human Genetics</i> , 1998, 102, 327-331.	3.8	40
71	Bypassing mitochondrial complex III using alternative oxidase inhibits acute pulmonary oxygen sensing. <i>Science Advances</i> , 2020, 6, eaba0694.	10.3	39
72	Analysis of the trinucleotide CAG repeat from the human mitochondrial DNA polymerase gene in healthy and diseased individuals. <i>European Journal of Human Genetics</i> , 1999, 7, 140-146.	2.8	38

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73	Mitochondrial genotype modulates mtDNA copy number and organismal phenotype in <i>Drosophila</i> . <i>Mitochondrion</i> , 2017, 34, 75-83.	3.4	38
74	A Rolling Circle Replication Mechanism Produces Multimeric Lariats of Mitochondrial DNA in <i>Caenorhabditis elegans</i> . <i>PLoS Genetics</i> , 2015, 11, e1004985.	3.5	37
75	Alternative Oxidase Attenuates Cigarette Smoke-induced Lung Dysfunction and Tissue Damage. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2019, 60, 515-522.	2.9	37
76	Acquisition of the wobble modification in mitochondrial tRNA <sup>Leu</sup> (CUN) bearing the G12300A mutation suppresses the MELAS molecular defect. <i>Human Molecular Genetics</i> , 2006, 15, 897-904.	2.9	36
77	Overexpression of MTERFD1 or MTERFD3 impairs the completion of mitochondrial DNA replication. <i>Molecular Biology Reports</i> , 2011, 38, 1321-1328.	2.3	36
78	Modelling in <i>Escherichia coli</i> of mutations in mitoribosomal protein S12: novel mutant phenotypes of rpsL. <i>Molecular Microbiology</i> , 1999, 31, 1735-1746.	2.5	33
79	Comparative Genetics of Functional Trinucleotide Tandem Repeats in Humans and Apes. <i>Journal of Molecular Evolution</i> , 2004, 59, 329-339.	1.8	33
80	Screen for mitochondrial DNA copy number maintenance genes reveals essential role for ATP synthase. <i>Molecular Systems Biology</i> , 2014, 10, 734.	7.2	33
81	The mitochondrial inner membrane AAA metalloprotease family in metazoans. <i>FEBS Letters</i> , 2000, 481, 91-95.	2.8	32
82	Mitochondrial transcription termination factor 1 directs polar replication fork pausing. <i>Nucleic Acids Research</i> , 2016, 44, 5732-5742.	14.5	32
83	The alternative oxidase, a tool for compensating cytochrome <i>c</i> oxidase deficiency in human cells. <i>Physiologia Plantarum</i> , 2009, 137, 427-434.	5.2	31
84	Unique features of DNA replication in mitochondria: A functional and evolutionary perspective. <i>BioEssays</i> , 2014, 36, 1024-1031.	2.5	31
85	Expression of the alternative oxidase mitigates beta-amyloid production and toxicity in model systems. <i>Free Radical Biology and Medicine</i> , 2016, 96, 57-66.	2.9	31
86	Systematic Segregation to Mutant Mitochondrial DNA and Accompanying Loss of Mitochondrial DNA in Human NT2 Teratocarcinoma Cybrids. <i>Genetics</i> , 2005, 170, 1879-1885.	2.9	29
87	Analysis of Replication Intermediates Indicates That <i>Drosophila melanogaster</i> Mitochondrial DNA Replicates by a Strand-Coupled Theta Mechanism. <i>PLoS ONE</i> , 2013, 8, e53249.	2.5	29
88	Post-transcriptional regulation of messenger abundance in rat liver and hepatoma. <i>Nucleic Acids Research</i> , 1980, 8, 3087-3104.	14.5	27
89	The bidirectional promoter of two genes for the mitochondrial translational apparatus in mouse is regulated by an array of CCAAT boxes interacting with the transcription factor NF-Y. <i>Nucleic Acids Research</i> , 2007, 35, 664-677.	14.5	27
90	Phenotypic rescue of a <i>Drosophila</i> model of mitochondrial ANT1 disease. <i>DMM Disease Models and Mechanisms</i> , 2014, 7, 635-48.	2.4	27

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91	Mitochondrial Dysfunction Plus High-Sugar Diet Provokes a Metabolic Crisis That Inhibits Growth. PLoS ONE, 2016, 11, e0145836.	2.5	27
92	Mitochondrial disease in flies. Biochimica Et Biophysica Acta - Bioenergetics, 2004, 1659, 190-196.	1.0	26
93	Two distinct, sequence-specific DNA-binding proteins interact independently with the major replication pause region of sea urchin mtDNA. Nucleic Acids Research, 1993, 21, 2801-2808.	14.5	25
94	The relationship between somatic mtDNA rearrangements, human heart disease and aging. Human Molecular Genetics, 2002, 11, 317-324.	2.9	25
95	dj-1 <sup>2</sup> regulates oxidative stress, insulin-like signaling and development in Drosophila melanogaster. Cell Cycle, 2012, 11, 3876-3886.	2.6	25
96	Expression of the Gene for Mitoribosomal Protein S12 Is Controlled in Human Cells at the Levels of Transcription, RNA Splicing, and Translation. Journal of Biological Chemistry, 1999, 274, 31853-31862.	3.4	24
97	Isolation and Purification of Rat Hepatoma Nuclei Active in the Transport of Messenger RNA in vitro. FEBS Journal, 1982, 121, 597-607.	0.2	23
98	The 7472insC mtDNA mutation impairs 5â€² and 3â€² processing of tRNASer(UCN). Biochemical and Biophysical Research Communications, 2004, 322, 803-813.	2.1	23
99	DNA conformationâ€¦dependent activities of human mitochondrial RNA polymerase. Genes To Cells, 2009, 14, 1029-1042.	1.2	23
100	A Cytoplasmic Suppressor of a Nuclear Mutation Affecting Mitochondrial Functions in <i>Drosophila</i> . Genetics, 2012, 192, 483-493.	2.9	23
101	Making mitochondrial mutants. Trends in Genetics, 2001, 17, 653-660.	6.7	22
102	A prevalent POLG CAG microsatellite length allele in humans and African great apes. Mammalian Genome, 2004, 15, 492-502.	2.2	22
103	The Alternative Oxidase AOX Does Not Rescue the Phenotype of <i>tko25t</i> Mutant Flies. G3: Genes, Genomes, Genetics, 2014, 4, 2013-2021.	1.8	22
104	Structural similarities between a mitochondrially encoded polypeptide and a family of prokaryotic respiratory toxins involved in plasmid maintenance suggest a novel mechanism for the evolutionary maintenance of mitochondrial DNA. Journal of Molecular Evolution, 1991, 32, 333-339.	1.8	21
105	Alternative respiratory chain enzymes: Therapeutic potential and possible pitfalls. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2019, 1865, 854-866.	3.8	21
106	Chromosomal Locations of Three Human Nuclear Genes (RPSM12, TUFM, and AFG3L1) Specifying Putative Components of the Mitochondrial Gene Expression Apparatus. Genomics, 1998, 48, 384-388.	2.9	20
107	The Dominant Cold-Sensitive Out-Cold Mutants of Drosophila melanogaster Have Novel Missense Mutations in the Voltage-Gated Sodium Channel Gene paralytic. Genetics, 2008, 180, 873-884.	2.9	20
108	Transcripts of three mitochondrial genes in the RNA of sea urchin eggs and embryos. Developmental Biology, 1983, 97, 500-505.	2.0	19



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109	Gene dosage and selective expression modify phenotype in a <i>Drosophila</i> model of human mitochondrial disease. <i>Mitochondrion</i> , 2003, 3, 83-96.	3.4	19
110	Therapies in inborn errors of oxidative metabolism. <i>Trends in Endocrinology and Metabolism</i> , 2012, 23, 488-495.	7.1	19
111	Characterization of a high-affinity binding site for a DNA-binding protein from sea urchin embryo mitochondria. <i>Nucleic Acids Research</i> , 1993, 21, 811-816.	14.5	18
112	Effects on mitochondrial transcription of manipulating mTERF protein levels in cultured human HEK293 cells. <i>BMC Molecular Biology</i> , 2010, 11, 72.	3.0	17
113	Mitochondrial Transcription Terminator Family Members mTTF and mTerf5 Have Opposing Roles in Coordination of mtDNA Synthesis. <i>PLoS Genetics</i> , 2013, 9, e1003800.	3.5	17
114	Long-Extension PCR to Detect Deleted Mitochondrial DNA Molecules Is Compromized by Technical Artefacts. <i>Biochemical and Biophysical Research Communications</i> , 1999, 254, 507-514.	2.1	16
115	<i>Ciona intestinalis</i> NADH dehydrogenase NDX confers stress-resistance and extended lifespan on <i>Drosophila</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2014, 1837, 1861-1869.	1.0	16
116	Phenotypic suppression of the <i>Drosophila</i> mitochondrial disease-like mutant <i>tko25t</i> by duplication of the mutant gene in its natural chromosomal context. <i>Mitochondrion</i> , 2009, 9, 353-363.	3.4	15
117	Diiron centre mutations in <i>Ciona intestinalis</i> alternative oxidase abolish enzymatic activity and prevent rescue of cytochrome oxidase deficiency in flies. <i>Scientific Reports</i> , 2016, 5, 18295.	3.3	15
118	Respiratory chain signalling is essential for adaptive remodelling following cardiac ischaemia. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 3534-3548.	3.6	15
119	Immunochemical Characterization of a Novel Mitochondrially Located Protein Encoded by a Nuclear Gene within the DFNB8/10 Critical Region on 21q22.3. <i>Biochemical and Biophysical Research Communications</i> , 1997, 238, 806-810.	2.1	14
120	Expression of the nuclear gene encoding mitochondrial ATP synthase subunit alpha in early development of <i>Drosophila</i> and sea urchin. <i>Molecular Biology Reports</i> , 1998, 25, 87-94.	2.3	14
121	Expression of <i>Ciona intestinalis</i> AOX causes male reproductive defects in <i>Drosophila melanogaster</i> . <i>BMC Developmental Biology</i> , 2017, 17, 9.	2.1	14
122	Polyadenylation and degradation of structurally abnormal mitochondrial tRNAs in human cells. <i>Nucleic Acids Research</i> , 2018, 46, 5209-5226.	14.5	14
123	Developmental arrest in <i>Drosophila melanogaster</i> caused by mitochondrial DNA replication defects cannot be rescued by the alternative oxidase. <i>Scientific Reports</i> , 2018, 8, 10882.	3.3	14
124	RNase H1 promotes replication fork progression through oppositely transcribed regions of <i>Drosophila</i> mitochondrial DNA. <i>Journal of Biological Chemistry</i> , 2019, 294, 4331-4344.	3.4	14
125	Lethal Interaction of Nuclear and Mitochondrial Genotypes in <i>Drosophila melanogaster</i> . G3: Genes, Genomes, Genetics, 2019, 9, 2225-2234.	1.8	13
126	Alternative NADH dehydrogenase extends lifespan and increases resistance to xenobiotics in <i>Drosophila</i> . <i>Biogerontology</i> , 2020, 21, 155-171.	3.9	13



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127	Regulation of growth in <i>Drosophila melanogaster</i> : the roles of mitochondrial metabolism. <i>Journal of Biochemistry</i> , 2020, 167, 267-277.	1.7	13
128	NF-Y influences directionality of transcription from the bidirectional <i>Mrps12/Sarsm</i> promoter in both mouse and human cells. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2009, 1789, 432-442.	1.9	12
129	Ligand-Bound GeneSwitch Causes Developmental Aberrations in <i>Drosophila</i> that Are Alleviated by the Alternative Oxidase. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 2839-2846.	1.8	12
130	What physiological role(s) does the alternative oxidase perform in animals?. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2022, 1863, 148556.	1.0	12
131	The dangers of extended PCR in the clinic. <i>Nature Medicine</i> , 1999, 5, 965-966.	30.7	11
132	In vivo mitochondrial DNA-protein interactions in sea urchin eggs and embryos. <i>Current Genetics</i> , 1999, 34, 449-458.	1.7	11
133	Novel coding-region polymorphisms in mitochondrial seryl-tRNA synthetase ( <i>SARSM</i> ) and mitoribosomal protein <i>S12</i> ( <i>RPMS12</i> ) genes in <i>DFNA4</i> autosomal dominant deafness families. <i>Human Mutation</i> , 2001, 17, 433-434.	2.5	11
134	Modulation of <i>Mrps12/Sarsm</i> promoter activity in response to mitochondrial stress. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2008, 1783, 2352-2362.	4.1	11
135	Postdockin' in the free world. <i>EMBO Reports</i> , 2010, 11, 1-1.	4.5	11
136	Genome wide study of NF-Y type CCAAT boxes in unidirectional and bidirectional promoters in human and mouse. <i>Journal of Theoretical Biology</i> , 2011, 281, 74-83.	1.7	11
137	$\hat{\Gamma}^2$ carbonic anhydrase is required for female fertility in <i>Drosophila melanogaster</i> . <i>Frontiers in Zoology</i> , 2015, 12, 19.	2.0	11
138	Expression of the Alternative Oxidase Influences Jun N-Terminal Kinase Signaling and Cell Migration. <i>Molecular and Cellular Biology</i> , 2018, 38, .	2.3	11
139	Hyperoxia but not AOX expression mitigates pathological cardiac remodeling in a mouse model of inflammatory cardiomyopathy. <i>Scientific Reports</i> , 2019, 9, 12741.	3.3	11
140	Apparent functional independence of the mitochondrial and nuclear transcription systems in cultured human cells. <i>Molecular Genetics and Genomics</i> , 1994, 245, 760-768.	2.4	10
141	The study of cell-death proteins in the outer mitochondrial membrane by chemical cross-linking. <i>Biochemical Journal</i> , 1997, 325, 321-324.	3.7	10
142	Structural rearrangements in the mitochondrial genome of <i>Drosophila melanogaster</i> induced by elevated levels of the replicative DNA helicase. <i>Nucleic Acids Research</i> , 2018, 46, 3034-3046.	14.5	10
143	Post-transcriptional control of messenger abundance. <i>Molecular Biology Reports</i> , 1981, 7, 159-162.	2.3	8
144	Do ribosomes regulate mitochondrial RNA synthesis?. <i>BioEssays</i> , 1989, 11, 27-34.	2.5	8

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145	Phenotypic effects of dietary stress in combination with a respiratory chain bypass in mice. <i>Physiological Reports</i> , 2019, 7, e14159.	1.7	8
146	The alternative oxidase (AOX) increases sulphide tolerance in the highly invasive marine invertebrate <i>Ciona intestinalis</i> . <i>Journal of Experimental Biology</i> , 2021, 224, .	1.7	8
147	Cyanide resistant respiration and the alternative oxidase pathway: A journey from plants to mammals. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2022, 1863, 148567.	1.0	8
148	The V368i mutation in Twinkle does not segregate with adPEO. <i>Annals of Neurology</i> , 2003, 53, 278-278.	5.3	7
149	From and to a very grey area. <i>EMBO Reports</i> , 2011, 12, 479-479.	4.5	7
150	Alternative oxidase confers nutritional limitation on <i>Drosophila</i> development. <i>Journal of Experimental Zoology Part A: Ecological and Integrative Physiology</i> , 2019, 331, 341-356.	1.9	7
151	Manipulating mtDNA in vivo reprograms metabolism via novel response mechanisms. <i>PLoS Genetics</i> , 2019, 15, e1008410.	3.5	7
152	The selfish organelle. <i>Trends in Genetics</i> , 1987, 3, 337-341.	6.7	6
153	Nuclear expression of mitochondrial genes implicated in human encephalomyopathies. <i>Biochemical Society Transactions</i> , 1994, 22, 413S-413S.	3.4	6
154	Rebuttal to Pak et al.: New data, old chestnuts. <i>Aging Cell</i> , 2003, 2, 19-20.	6.7	6
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