Vilhelm A Bohr

List of Publications by Year
in descending order

392	36,297	2093	4750
papers	citations	100 h-index	169 g-index

429	429	429	28253
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Ageing as a risk factor for neurodegenerative disease. Nature Reviews Neurology, 2019, 15, 565-581.	4.9	1,578
2	Mitophagy inhibits amyloid-Î² and tau pathology and reverses cognitive deficits in models of Alzheimerâ€™s disease. Nature Neuroscience, 2019, 22, 401-412.	7.1	1,008
3	SIRT6 is a histone H3 lysine 9 deacetylase that modulates telomeric chromatin. Nature, 2008, 452, 492-496.	13.7	945
4	Nutrient-Sensitive Mitochondrial NAD+ Levels Dictate Cell Survival. Cell, 2007, 130, 1095-1107.	13.5	855
5	Senolytic therapy alleviates Aβ-associated oligodendrocyte progenitor cell senescence and cognitive deficits in an Alzheimerâ€™s disease model. Nature Neuroscience, 2019, 22, 719-728.	7.1	577
6	Defective Mitophagy in XPA via PARP-1 Hyperactivation and NAD+/SIRT1 Reduction. Cell, 2014, 157, 882-896.	13.5	554
7	Mitophagy and Alzheimerâ€™s Disease: Cellular and Molecular Mechanisms. Trends in Neurosciences, 2017, 40, 151-166.	4.2	553
8	The Bloom's and Werner's syndrome proteins are DNA structure-specific helicases. Nucleic Acids Research, 2001, 29, 2843-2849.	6.5	518
9	Base excision repair of oxidative DNA damage and association with cancer and aging. Carcinogenesis, 2008, 30, 2-10.	1.3	511
10	Human RecQ Helicases in DNA Repair, Recombination, and Replication. Annual Review of Biochemistry, 2014, 83, 519-552.	5.0	461
11	Repair of Oxidative Damage to Nuclear and Mitochondrial DNA in Mammalian Cells. Journal of Biological Chemistry, 1997, 272, 25409-25412.	1.6	427
12	NAD + Replenishment Improves Lifespan and Healthspan in Ataxia Telangiectasia Models via Mitophagy and DNA Repair. Cell Metabolism, 2016, 24, 566-581.	7.2	420
13	Werner's syndrome protein (WRN) migrates Holliday junctions and coâ€localizes with RPA upon replication arrest. EMBO Reports, 2000, 1, 80-84.	2.0	378
14	A research agenda for aging in China in the 21st century. Ageing Research Reviews, 2015, 24, 197-205.	5.0	374
15	Effects of Sex, Strain, and Energy Intake on Hallmarks of Aging in Mice. Cell Metabolism, 2016, 23, 1093-1112.	7.2	360
16	Repair of oxidative DNA damage in nuclear and mitochondrial DNA, and some changes with aging in mammalian cells1,2 1Guest Editor: Miral Dizdaroglu 2This article is part of a series of reviews on â€œOxidative DNA Damage and Repair.â€•The full list of papers may be found on the homepage of the journal.. Free Radical Biology and Medicine, 2002, 32, 804-812.	1.3	346
17	Telomere-binding Protein TRF2 Binds to and Stimulates the Werner and Bloom Syndrome Helicases. Journal of Biological Chemistry, 2002, 277, 41110-41119.	1.6	334
18	NAD + in Aging: Molecular Mechanisms and Translational Implications. Trends in Molecular Medicine, 2017, 23, 899-916.	3.5	333

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19	NAD ‹sup›+‹/sup› supplementation normalizes key Alzheimerâ€™s features and DNA damage responses in a new AD mouse model with introduced DNA repair deficiency. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E1876-E1885.	3.3	316
20	A High-Fat Diet and NAD + Activate Sirt1 to Rescue Premature Aging in Cockayne Syndrome. Cell Metabolism, 2014, 20, 840-855.	7.2	306
21	Nuclear DNA damage signalling to mitochondria in ageing. Nature Reviews Molecular Cell Biology, 2016, 17, 308-321.	16.1	294
22	Functional and Physical Interaction between WRN Helicase and Human Replication Protein A. Journal of Biological Chemistry, 1999, 274, 18341-18350.	1.6	287
23	Mitochondrial SIRT3 Mediates Adaptive Responses of Neurons to Exercise and Metabolic and Excitatory Challenges. Cell Metabolism, 2016, 23, 128-142.	7.2	286
24	DNA Damage, DNA Repair, Aging, and Neurodegeneration. Cold Spring Harbor Perspectives in Medicine, 2015, 5, a025130.	2.9	285
25	The mechanics of base excision repair, and its relationship to aging and disease. DNA Repair, 2007, 6, 544-559.	1.3	280
26	DNA repair deficiency in neurodegeneration. Progress in Neurobiology, 2011, 94, 166-200.	2.8	280
27	Ku complex interacts with and stimulates the Werner protein. Genes and Development, 2000, 14, 907-912.	2.7	276
28	Replication Protein A Physically Interacts with the Bloom's Syndrome Protein and Stimulates Its Helicase Activity. Journal of Biological Chemistry, 2000, 275, 23500-23508.	1.6	274
29	Mitophagy in neurodegeneration and aging. Neurochemistry International, 2017, 109, 202-209.	1.9	272
30	SIRT6 stabilizes DNA-dependent Protein Kinase at chromatin for DNA double-strand break repair. Aging, 2009, 1, 109-121.	1.4	270
31	Protecting the mitochondrial powerhouse. Trends in Cell Biology, 2015, 25, 158-170.	3.6	260
32	Repair of mitochondrial DNA after various types of DNA damage in Chinese hamster ovary cells. Carcinogenesis, 1992, 13, 1967-1973.	1.3	259
33	Defective DNA base excision repair in brain from individuals with Alzheimer's disease and amnestic mild cognitive impairment. Nucleic Acids Research, 2007, 35, 5545-5555.	6.5	253
34	Nicotinamide Improves Aspects of Healthspan, but Not Lifespan, in Mice. Cell Metabolism, 2018, 27, 667-676.e4.	7.2	242
35	Epigenetic inactivation of the premature aging Werner syndrome gene in human cancer. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 8822-8827.	3.3	240
36	Oxidative damage in telomeric DNA disrupts recognition by TRF1 and TRF2. Nucleic Acids Research, 2005, 33, 1230-1239.	6.5	237

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37	The Werner syndrome protein operates in base excision repair and cooperates with DNA polymerase Â. Nucleic Acids Research, 2006, 34, 745-754.	6.5	228
38	Rising from the RecQ-age: the role of human RecQ helicases in genome maintenance. Trends in Biochemical Sciences, 2008, 33, 609-620.	3.7	224
39	Repair Pathways for Processing of 8-Oxoguanine in DNA by Mammalian Cell Extracts. Journal of Biological Chemistry, 1998, 273, 33811-33816.	1.6	220
40	Role of DNA Polymerase Î² in the Excision Step of Long Patch Mammalian Base Excision Repair. Journal of Biological Chemistry, 1999, 274, 13741-13743.	1.6	202
41	Mitochondrial DNA repair pathways. Mutation Research DNA Repair, 1999, 434, 137-148.	3.8	200
42	Human Embryonic Stem Cells Have Enhanced Repair of Multiple Forms of DNA Damage. Stem Cells, 2008, 26, 2266-2274.	1.4	193
43	Removal of Oxidative DNA Damage via FEN1-Dependent Long-Patch Base Excision Repair in Human Cell Mitochondria. Molecular and Cellular Biology, 2008, 28, 4975-4987.	1.1	192
44	FEN1 Stimulation of DNA Polymerase Î² Mediates an Excision Step in Mammalian Long Patch Base Excision Repair. Journal of Biological Chemistry, 2000, 275, 4460-4466.	1.6	187
45	Human premature aging, DNA repair and RecQ helicases. Nucleic Acids Research, 2007, 35, 7527-7544.	6.5	186
46	Cockayne syndrome: Clinical features, model systems and pathways. Ageing Research Reviews, 2017, 33, 3-17.	5.0	184
47	Roles of Werner syndrome protein in protection of genome integrity. DNA Repair, 2010, 9, 331-344.	1.3	183
48	Mitochondrial DNA repair of oxidative damage in mammalian cells. Gene, 2002, 286, 127-134.	1.0	179
49	Cockayne syndrome group B protein prevents the accumulation of damaged mitochondria by promoting mitochondrial autophagy. Journal of Experimental Medicine, 2012, 209, 855-869.	4.2	177
50	NAD ⁺ supplementation reduces neuroinflammation and cell senescence in a transgenic mouse model of Alzheimerâ€™s disease via cGASâ€“STING. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	176
51	Repair of Formamidopyrimidines in DNA Involves Different Glycosylases. Journal of Biological Chemistry, 2005, 280, 40544-40551.	1.6	174
52	Novel DNA mismatch-repair activity involving YB-1 in human mitochondria. DNA Repair, 2009, 8, 704-719.	1.3	174
53	Increased Hypermutation at G and C Nucleotides in Immunoglobulin Variable Genes from Mice Deficient in the MSH2 Mismatch Repair Protein. Journal of Experimental Medicine, 1998, 187, 1745-1751.	4.2	170
54	Mitochondrial and nuclear DNA-repair capacity of various brain regions in mouse is altered in an age-dependent manner. Neurobiology of Aging, 2006, 27, 1129-1136.	1.5	168

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55	Gene specific DNA repair. Carcinogenesis, 1991, 12, 1983-1992.	1.3	167
56	NAD+ augmentation restores mitophagy and limits accelerated aging in Werner syndrome. Nature Communications, 2019, 10, 5284.	5.8	165
57	Werner syndrome and the function of the Werner protein; what they can teach us about the molecular aging process.. Carcinogenesis, 2003, 24, 791-802.	1.3	164
58	Human DNA polymerase Î² initiates DNA synthesis during long-patch repair of reduced AP sites in DNA. EMBO Journal, 2001, 20, 1477-1482.	3.5	159
59	DNA damage, mutation and fine structure DNA repair in aging. Mutation Research - DNAging, 1995, 338, 25-34.	3.3	157
60	POT1 Stimulates RecQ Helicases WRN and BLM to Unwind Telomeric DNA Substrates. Journal of Biological Chemistry, 2005, 280, 32069-32080.	1.6	157
61	The HRDC domain of BLM is required for the dissolution of double Holliday junctions. EMBO Journal, 2005, 24, 2679-2687.	3.5	150
62	The clinical characteristics of Werner syndrome: molecular and biochemical diagnosis. Human Genetics, 2008, 124, 369-377.	1.8	147
63	Mitochondrial DNA damage and repair in neurodegenerative disorders. DNA Repair, 2008, 7, 1110-1120.	1.3	146
64	Base excision repair in nuclear and mitochondrial DNA. Progress in Molecular Biology and Translational Science, 2001, 68, 285-297.	1.9	144
65	Cockayne Syndrome Group B Cellular and Biochemical Functions. American Journal of Human Genetics, 2003, 73, 1217-1239.	2.6	144
66	An Oxidative Damage-specific Endonuclease from Rat Liver Mitochondria. Journal of Biological Chemistry, 1997, 272, 27338-27344.	1.6	143
67	Werner Protein Is a Target of DNA-dependent Protein Kinase in Vivo and in Vitro, and Its Catalytic Activities Are Regulated by Phosphorylation. Journal of Biological Chemistry, 2002, 277, 18291-18302.	1.6	141
68	Central Role for the Werner Syndrome Protein/Poly(ADP-Ribose) Polymerase 1 Complex in the Poly(ADP-Ribosyl)ation Pathway after DNA Damage. Molecular and Cellular Biology, 2003, 23, 8601-8613.	1.1	140
69	Gene expression profiling in Werner syndrome closely resembles that of normal aging. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 12259-12264.	3.3	140
70	Primary fibroblasts of Cockayne syndrome patients are defective in cellular repair of 8â€hydroxyguanine and 8â€hydroxyadenine resulting from oxidative stress. FASEB Journal, 2003, 17, 668-674.	0.2	140
71	The Werner Syndrome Protein Is Involved in RNA Polymerase II Transcription. Molecular Biology of the Cell, 1999, 10, 2655-2668.	0.9	139
72	The Cockayne Syndrome Group B Gene Product Is Involved in General Genome Base Excision Repair of 8-Hydroxyguanine in DNA. Journal of Biological Chemistry, 2001, 276, 45772-45779.	1.6	138

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73	Interaction of Human AP Endonuclease 1 with Flap Endonuclease 1 and Proliferating Cell Nuclear Antigen Involved in Long-Patch Base Excision Repair. Biochemistry, 2001, 40, 12639-12644.	1.2	136
74	Mitochondrial DNA repair and association with aging â€“ An update. Experimental Gerontology, 2010, 45, 478-488.	1.2	134
75	Genomic heterogeneity of nucleotide excision repair. Gene, 2000, 250, 15-30.	1.0	129
76	A Small Molecule Inhibitor of the BLM Helicase Modulates Chromosome Stability in Human Cells. Chemistry and Biology, 2013, 20, 55-62.	6.2	128
77	The role of DNA repair in brain related disease pathology. DNA Repair, 2013, 12, 578-587.	1.3	127
78	The role of Cockayne Syndrome group B (CSB) protein in base excision repair and aging. Mechanisms of Ageing and Development, 2008, 129, 441-448.	2.2	126
79	Oxidative DNA damage processing in nuclear and mitochondrial DNA. Biochimie, 1999, 81, 155-160.	1.3	125
80	Cockayne syndrome group B protein promotes mitochondrial DNA stability by supporting the DNA repair association with the mitochondrial membrane. FASEB Journal, 2010, 24, 2334-2346.	0.2	124
81	Repair of 8-oxoguanine in DNA is deficient in Cockayne syndrome group B cells. Nucleic Acids Research, 1999, 27, 1365-1368.	6.5	123
82	BDNF and Exercise Enhance Neuronal DNA Repair by Stimulating CREB-Mediated Production of Apurinic/Apyrimidinic Endonuclease 1. NeuroMolecular Medicine, 2014, 16, 161-174.	1.8	121
83	Signaling by cGASâ€“STING in Neurodegeneration, Neuroinflammation, and Aging. Trends in Neurosciences, 2021, 44, 83-96.	4.2	121
84	The mitochondrial transcription factor A functions in mitochondrial base excision repair. DNA Repair, 2010, 9, 1080-1089.	1.3	120
85	Colocalization, Physical, and Functional Interaction between Werner and Bloom Syndrome Proteins. Journal of Biological Chemistry, 2002, 277, 22035-22044.	1.6	119
86	Mitochondria in the signaling pathways that control longevity and health span. Ageing Research Reviews, 2019, 54, 100940.	5.0	118
87	Tomatidine enhances lifespan and healthspan in C. elegans through mitophagy induction via the SKN-1/Nrf2 pathway. Scientific Reports, 2017, 7, 46208.	1.6	116
88	Enzymatic and DNA binding properties of purified WRN protein: high affinity binding to single-stranded DNA but not to DNA damage induced by 4NQO. Nucleic Acids Research, 1999, 27, 3557-3566.	6.5	114
89	The Werner Syndrome Protein Stimulates DNA Polymerase Î² Strand Displacement Synthesis via Its Helicase Activity. Journal of Biological Chemistry, 2003, 278, 22686-22695.	1.6	113
90	DNA repair and aging in mouse liver: 8-oxodG glycosylase activity increase in mitochondrial but not in nuclear extracts. Free Radical Biology and Medicine, 2001, 30, 916-923.	1.3	112

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91	DNA polymerase Î² deficiency leads to neurodegeneration and exacerbates Alzheimer disease phenotypes. Nucleic Acids Research, 2015, 43, 943-959.	6.5	110
92	Werner Syndrome Protein Contains Three Structure-specific DNA Binding Domains. Journal of Biological Chemistry, 2003, 278, 52997-53006.	1.6	109
93	Mitochondrial and nuclear DNA base excision repair are affected differently by caloric restriction. FASEB Journal, 2004, 18, 595-597.	0.2	109
94	Natural polyphenols as sirtuin 6 modulators. Scientific Reports, 2018, 8, 4163.	1.6	109
95	Unwinding of a DNA Triple Helix by the Werner and Bloom Syndrome Helicases. Journal of Biological Chemistry, 2001, 276, 3024-3030.	1.6	108
96	The Processing of Holliday Junctions by BLM and WRN Helicases Is Regulated by p53. Journal of Biological Chemistry, 2002, 277, 31980-31987.	1.6	107
97	Base excision repair capacity in mitochondria and nuclei: tissueâ€specific variations. FASEB Journal, 2002, 16, 1895-1902.	0.2	105
98	WRN Interacts Physically and Functionally with the Recombination Mediator Protein RAD52. Journal of Biological Chemistry, 2003, 278, 36476-36486.	1.6	105
99	Cooperation of the Cockayne Syndrome Group B Protein and Poly(ADP-Ribose) Polymerase 1 in the Response to Oxidative Stress. Molecular and Cellular Biology, 2005, 25, 7625-7636.	1.1	104
100	Cockayne syndrome B protein stimulates apurinic endonuclease 1 activity and protects against agents that introduce base excision repair intermediates. Nucleic Acids Research, 2007, 35, 4103-4113.	6.5	104
101	JNK Phosphorylates SIRT6 to Stimulate DNA Double-Strand Break Repair in Response to Oxidative Stress by Recruiting PARP1 to DNA Breaks. Cell Reports, 2016, 16, 2641-2650.	2.9	104
102	Factors that influence telomeric oxidative base damage and repair by DNA glycosylase OGG1. DNA Repair, 2011, 10, 34-44.	1.3	103
103	Linkage between Werner Syndrome Protein and the Mre11 Complex via Nbs1. Journal of Biological Chemistry, 2004, 279, 21169-21176.	1.6	102
104	Evidence that OGG1 Glycosylase Protects Neurons against Oxidative DNA Damage and Cell Death under Ischemic Conditions. Journal of Cerebral Blood Flow and Metabolism, 2011, 31, 680-692.	2.4	101
105	Inhibition of the Bloom's and Werner's Syndrome Helicases by G-Quadruplex Interacting Ligands. Biochemistry, 2001, 40, 15194-15202.	1.2	100
106	The Human Werner Syndrome Protein Stimulates Repair of Oxidative DNA Base Damage by the DNA Glycosylase NEIL1. Journal of Biological Chemistry, 2007, 282, 26591-26602.	1.6	100
107	Mitochondrial repair of 8-oxoguanine is deficient in Cockayne syndrome group B. Oncogene, 2002, 21, 8675-8682.	2.6	99
108	RECQL4, the Protein Mutated in Rothmund-Thomson Syndrome, Functions in Telomere Maintenance. Journal of Biological Chemistry, 2012, 287, 196-209.	1.6	99

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109	RECQL4 localizes to mitochondria and preserves mitochondrial DNA integrity. Aging Cell, 2012, 11, 456-466.	3.0	97
110	Coordinate Action of the Helicase and 3â€² to 5â€² Exonuclease of Werner Syndrome Protein. Journal of Biological Chemistry, 2001, 276, 44677-44687.	1.6	96
111	DNA repair fine structure and its relations to genomic instability. Carcinogenesis, 1995, 16, 2885-2892.	1.3	95
112	p53 Modulates the Exonuclease Activity of Werner Syndrome Protein. Journal of Biological Chemistry, 2001, 276, 35093-35102.	1.6	95
113	Mitochondrial endogenous oxidative damage has been overestimated. FASEB Journal, 2000, 14, 355-360.	0.2	94
114	Cockayne Syndrome Group B Protein Stimulates Repair of Formamidopyrimidines by NEIL1 DNA Glycosylase. Journal of Biological Chemistry, 2009, 284, 9270-9279.	1.6	92
115	Poly(ADP-ribose) polymerase 1 regulates both the exonuclease and helicase activities of the Werner syndrome protein. Nucleic Acids Research, 2004, 32, 4003-4014.	6.5	89
116	p53 functions in the incorporation step in DNA base excision repair in mouse liver mitochondria. Oncogene, 2004, 23, 6559-6568.	2.6	89
117	Mitochondrial DNA, base excision repair and neurodegeneration. DNA Repair, 2008, 7, 1098-1109.	1.3	89
118	The Cockayne Syndrome Group B Gene Product Is Involved in Cellular Repair of 8-Hydroxyadenine in DNA. Journal of Biological Chemistry, 2002, 277, 30832-30837.	1.6	88
119	Ku heterodimer binds to both ends of the Werner protein and functional interaction occurs at the Werner N-terminus. Nucleic Acids Research, 2002, 30, 3583-3591.	6.5	86
120	Functional crosstalk between hOgg1 and the helicase domain of Cockayne syndrome group B protein. DNA Repair, 2002, 1, 913-927.	1.3	85
121	Stimulation of Flap Endonuclease-1 by the Bloom's Syndrome Protein. Journal of Biological Chemistry, 2004, 279, 9847-9856.	1.6	85
122	Junction of RecQ Helicase Biochemistry and Human Disease. Journal of Biological Chemistry, 2004, 279, 18099-18102.	1.6	85
123	Mitochondria-targeted Ogg1 and Aconitase-2 Prevent Oxidant-induced Mitochondrial DNA Damage in Alveolar Epithelial Cells. Journal of Biological Chemistry, 2014, 289, 6165-6176.	1.6	85
124	Single-molecule imaging reveals a common mechanism shared by G-quadruplexâ€“resolving helicases. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 8448-8453.	3.3	85
125	Roles of the Werner syndrome protein in pathways required for maintenance of genome stability. Experimental Gerontology, 2002, 37, 491-506.	1.2	84
126	Neurons Efficiently Repair Glutamate-induced Oxidative DNA Damage by a Process Involving CREB-mediated Up-regulation of Apurinic Endonuclease 1. Journal of Biological Chemistry, 2010, 285, 28191-28199.	1.6	84

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127	Genome instability in Alzheimer disease. Mechanisms of Ageing and Development, 2017, 161, 83-94.	2.2	83
128	Collaboration of Werner syndrome protein and BRCA1 in cellular responses to DNA interstrand cross-links. Nucleic Acids Research, 2006, 34, 2751-2760.	6.5	82
129	Pathways and functions of the Werner syndrome protein. Mechanisms of Ageing and Development, 2005, 126, 79-86.	2.2	81
130	WRN regulates pathway choice between classical and alternative non-homologous end joining. Nature Communications, 2016, 7, 13785.	5.8	81
131	RECQL4 Promotes DNA End Resection in Repair of DNA Double-Strand Breaks. Cell Reports, 2016, 16, 161-173.	2.9	81
132	Homogenous repair of singlet oxygen-induced DNA damage in differentially transcribed regions and strands of human mitochondrial DNA. Nucleic Acids Research, 1998, 26, 662-1997.	6.5	80
133	Loss of ARID1A in Tumor Cells Renders Selective Vulnerability to Combined Ionizing Radiation and PARP Inhibitor Therapy. Clinical Cancer Research, 2019, 25, 5584-5594.	3.2	80
134	Repair of 8-oxoG is slower in endogenous nuclear genes than in mitochondrial DNA and is without strand bias. DNA Repair, 2002, 1, 261-273.	1.3	78
135	Heterochromatin: an epigenetic point of view in aging. Experimental and Molecular Medicine, 2020, 52, 1466-1474.	3.2	78
136	Werner syndrome protein participates in a complex with RAD51, RAD54, RAD54B and ATR in response to ICL-induced replication arrest. Journal of Cell Science, 2006, 119, 5137-5146.	1.2	77
137	Inhibition of RNA Polymerase II Transcription in Human Cell Extracts by Cisplatin DNA Damage. Biochemistry, 1999, 38, 6204-6212.	1.2	76
138	Oxidized guanine lesions and hOgg1 activity in lung cancer. Oncogene, 2005, 24, 4496-4508.	2.6	76
139	The involvement of human RECQL4 in DNA doubleâ€strand break repair. Aging Cell, 2010, 9, 358-371.	3.0	76
140	Aprataxin localizes to mitochondria and preserves mitochondrial function. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 7437-7442.	3.3	76
141	Regulation of WRN Helicase Activity in Human Base Excision Repair. Journal of Biological Chemistry, 2004, 279, 53465-53474.	1.6	75
142	Direct and indirect roles of RECQL4 in modulating base excision repair capacity. Human Molecular Genetics, 2009, 18, 3470-3483.	1.4	75
143	Roles of RECQ helicases in recombination based DNA repair, genomic stability and aging. Biogerontology, 2009, 10, 235-252.	2.0	75
144	Reduced RNA polymerase II transcription in extracts of Cockayne syndrome and xeroderma pigmentosum/Cockayne syndrome cells. Nucleic Acids Research, 1997, 25, 3636-3642.	6.5	74

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145	A nucleolar targeting sequence in the Werner syndrome protein resides within residues 949-1092. Journal of Cell Science, 2002, 115, 3901-3907.	1.2	74
146	Gene-specific nuclear and mitochondrial repair of formamidopyrimidine DNA glycosylase-sensitive sites in Chinese hamster ovary cells. Mutation Research DNA Repair, 1996, 364, 183-192.	3.8	72
147	WRN Is Required for ATM Activation and the S-Phase Checkpoint in Response to Interstrand Cross-Linkâ€“Induced DNA Double-Strand Breaks. Molecular Biology of the Cell, 2008, 19, 3923-3933.	0.9	72
148	Cockayne syndrome group A and B proteins converge on transcription-linked resolution of non-B DNA. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 12502-12507.	3.3	72
149	Third complementarity-determining region of mutated VH immunoglobulin genes contains shorter V, D, J, P, and N components than non-mutated genes. Immunology, 2001, 103, 179-187.	2.0	71
150	Cell cycle-dependent phosphorylation regulates RECQL4 pathway choice and ubiquitination in DNA double-strand break repair. Nature Communications, 2017, 8, 2039.	5.8	71
151	Single Nucleotide Patch Base Excision Repair Is the Major Pathway for Removal of Thymine Glycol from DNA in Human Cell Extracts. Journal of Biological Chemistry, 2000, 275, 11809-11813.	1.6	70
152	DNA repair in the metallothionein gene increases with transcriptional activation. Nucleic Acids Research, 1987, 15, 10021-10030.	6.5	67
153	The transcriptional response after oxidative stress is defective in Cockayne syndrome group B cells. Oncogene, 2003, 22, 1135-1149.	2.6	66
154	Werner syndrome cells escape hydrogen peroxideâ€induced cell proliferation arrest. FASEB Journal, 2004, 18, 1970-1972.	0.2	66
155	RecQ helicases in DNA double strand break repair and telomere maintenance. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2012, 736, 15-24.	0.4	66
156	Mitochondrial deficiency in Cockayne syndrome. Mechanisms of Ageing and Development, 2013, 134, 275-283.	2.2	66
157	Werner Syndrome Protein Phosphorylation by Abl Tyrosine Kinase Regulates Its Activity and Distribution. Molecular and Cellular Biology, 2003, 23, 6385-6395.	1.1	65
158	The excitatory neurotransmitter glutamate stimulates DNA repair to increase neuronal resiliency. Mechanisms of Ageing and Development, 2011, 132, 405-411.	2.2	65
159	Endonuclease VIII-like 1 (NEIL1) promotes short-term spatial memory retention and protects from ischemic stroke-induced brain dysfunction and death in mice. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 14948-14953.	3.3	64
160	RECQL4 in genomic instability and aging. Trends in Genetics, 2012, 28, 624-631.	2.9	64
161	Differential requirement for the ATPase domain of the Cockayne syndrome group B gene in the processing of UV-induced DNA damage and 8-oxoguanine lesions in human cells. Nucleic Acids Research, 2002, 30, 782-793.	6.5	63
162	Mitochondrial repair of 8-oxoguanine and changes with aging. Experimental Gerontology, 2002, 37, 1189-1196.	1.2	63

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163	Mitochondrial DNA repair pathways. , 1999, 31, 391-398.		62
164	A role for WRN in telomere-based DNA damage responses. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 15073-15078.	3.3	62
165	Longevity and resistance to stress correlate with DNA repair capacity in Caenorhabditis elegans. Nucleic Acids Research, 2008, 36, 1380-1389.	6.5	62
166	Conserved helicase domain of human RecQ4 is required for strand annealing-independent DNA unwinding. DNA Repair, 2010, 9, 796-804.	1.3	61
167	Spatial Transcriptomics Reveals Genes Associated with Dysregulated Mitochondrial Functions and Stress Signaling in Alzheimer Disease. IScience, 2020, 23, 101556.	1.9	61
168	Gene expression and DNA repair in progeroid syndromes and human aging. Ageing Research Reviews, 2005, 4, 579-602.	5.0	60
169	Werner Protein Cooperates with the XRCC4-DNA Ligase IV Complex in End-Processing. Biochemistry, 2008, 47, 7548-7556.	1.2	59
170	The impact of base excision DNA repair in age-related neurodegenerative diseases. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2015, 776, 31-39.	0.4	59
171	Base excision DNA repair levels in mitochondrial lysates ofÂAlzheimer's disease. Neurobiology of Aging, 2014, 35, 1293-1300.	1.5	58
172	Recent Advances in Understanding Werner Syndrome. F1000Research, 2017, 6, 1779.	0.8	58
173	DNA damage and mitochondria in cancer and aging. Carcinogenesis, 2020, 41, 1625-1634.	1.3	58
174	NAD⁺ supplementation prevents STINGâ€induced senescence in ataxia telangiectasia by improving mitophagy. Aging Cell, 2021, 20, e13329.	3.0	58
175	Werner protein stimulates topoisomerase I DNA relaxation activity. Cancer Research, 2003, 63, 7136-46.	0.4	58
176	Oxidative DNA damage processing and changes with aging. Toxicology Letters, 1998, 102-103, 47-52.	0.4	57
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