

Susan S Wallace

List of Publications by Year in descending order

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66
papers

4,783
citations

101543

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67
docs citations

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times ranked

3550
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#	ARTICLE	IF	CITATIONS
1	Biological consequences of free radical-damaged DNA bases ^{1,2} 1Guest Editor: Miral Dizdaroglu 2This article is part of a series of reviews on "Oxidative DNA Damage and Repair."The full list of papers may be found on the homepage of the journal.. Free Radical Biology and Medicine, 2002, 33, 1-14.	2.9	375
2	Base excision repair: A critical player in many games. DNA Repair, 2014, 19, 14-26.	2.8	327
3	A novel human DNA glycosylase that removes oxidative DNA damage and is homologous to Escherichia coli endonuclease VIII. DNA Repair, 2002, 1, 517-529.	2.8	298
4	Thymine glycols and urea residues in M13 DNA constitute replicative blocks in vitro. Nucleic Acids Research, 1985, 13, 8035-8052.	14.5	267
5	Base excision repair and cancer. Cancer Letters, 2012, 327, 73-89.	7.2	257
6	Characterization of the Escherichia coli x-ray endonuclease, endonuclease III. Biochemistry, 1983, 22, 4071-4081.	2.5	201
7	Isolation and Characterization of Endonuclease VIII from Escherichia coli. Biochemistry, 1994, 33, 1255-1264.	2.5	175
8	The mouse ortholog of NEIL3 is a functional DNA glycosylase in vitro and in vivo. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 4925-4930.	7.1	169
9	Characterization of Escherichia coli Endonuclease VIII. Journal of Biological Chemistry, 1997, 272, 32230-32239.	3.4	146
10	The crystal structure of human endonuclease VIII-like 1 (NEIL1) reveals a zincless finger motif required for glycosylase activity. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 10284-10289.	7.1	132
11	The NEIL glycosylases remove oxidized guanine lesions from telomeric and promoter quadruplex DNA structures. Nucleic Acids Research, 2015, 43, 4039-4054.	14.5	129
12	A Role for the Fifth G-Track in G-Quadruplex Forming Oncogene Promoter Sequences during Oxidative Stress: Do These "Spare Tires" Have an Evolved Function?. ACS Central Science, 2015, 1, 226-233.	11.3	125
13	Saccharomyces cerevisiae Ntg1p and Ntg2p: Broad Specificity N-Glycosylases for the Repair of Oxidative DNA Damage in the Nucleus and Mitochondria. Biochemistry, 1999, 38, 11298-11306.	2.5	110
14	Single Qdot-labeled glycosylase molecules use a wedge amino acid to probe for lesions while scanning along DNA. Nucleic Acids Research, 2011, 39, 7487-7498.	14.5	107
15	Multiply damaged sites in DNA: interactions with Escherichia coli endonucleases III and VIII. Nucleic Acids Research, 1998, 26, 932-941.	14.5	103
16	Neil3 and NEIL1 DNA Glycosylases Remove Oxidative Damages from Quadruplex DNA and Exhibit Preferences for Lesions in the Telomeric Sequence Context. Journal of Biological Chemistry, 2013, 288, 27263-27272.	3.4	103
17	Two glycosylase families diffusively scan DNA using a wedge residue to probe for and identify oxidatively damaged bases. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E2091-9.	7.1	92
18	The enigma of endonuclease VIII. DNA Repair, 2003, 2, 441-453.	2.8	89

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19	DNA glycosylases search for and remove oxidized DNA bases. <i>Environmental and Molecular Mutagenesis</i> , 2013, 54, 691-704.	2.2	87
20	Nucleosome Disruption by DNA Ligase III-XRCC1 Promotes Efficient Base Excision Repair. <i>Molecular and Cellular Biology</i> , 2011, 31, 4623-4632.	2.3	80
21	The Fpg/Nei Family of DNA Glycosylases. <i>Progress in Molecular Biology and Translational Science</i> , 2012, 110, 71-91.	1.7	78
22	Rules of engagement for base excision repair in chromatin. <i>Journal of Cellular Physiology</i> , 2013, 228, 258-266.	4.1	75
23	Initiation of Base Excision Repair of Oxidative Lesions in Nucleosomes by the Human, Bifunctional DNA Glycosylase NTH1. <i>Molecular and Cellular Biology</i> , 2007, 27, 8442-8453.	2.3	72
24	Neil3, the final frontier for the DNA glycosylases that recognize oxidative damage. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2013, 743-744, 4-11.	1.0	72
25	NEIL3 Repairs Telomere Damage during S Phase to Secure Chromosome Segregation at Mitosis. <i>Cell Reports</i> , 2017, 20, 2044-2056.	6.4	57
26	Nucleosomes Suppress the Formation of Double-strand DNA Breaks during Attempted Base Excision Repair of Clustered Oxidative Damages. <i>Journal of Biological Chemistry</i> , 2014, 289, 19881-19893.	3.4	56
27	Cooperation of the NEIL3 and Fanconi anemia/BRCA pathways in interstrand crosslink repair. <i>Nucleic Acids Research</i> , 2020, 48, 3014-3028.	14.5	53
28	Non-specific DNA binding interferes with the efficient excision of oxidative lesions from chromatin by the human DNA glycosylase, NEIL1. <i>DNA Repair</i> , 2010, 9, 134-143.	2.8	51
29	Structural Characterization of a Mouse Ortholog of Human NEIL3 with a Marked Preference for Single-Stranded DNA. <i>Structure</i> , 2013, 21, 247-256.	3.3	51
30	Studies on Escherichia coli X-Ray Endonuclease Specificity: Roles of Hydroxyl and Reducing Radicals in the Production of DNA Lesions. <i>Radiation Research</i> , 1977, 69, 328.	1.5	48
31	Structural Characterization of Viral Ortholog of Human DNA Glycosylase NEIL1 Bound to Thymine Glycol or 5-Hydroxyuracil-containing DNA. <i>Journal of Biological Chemistry</i> , 2012, 287, 4288-4298.	3.4	48
32	Hide and seek: How do DNA glycosylases locate oxidatively damaged DNA bases amidst a sea of undamaged bases?. <i>Free Radical Biology and Medicine</i> , 2017, 107, 170-178.	2.9	48
33	Germ-line variant of human NTH1 DNA glycosylase induces genomic instability and cellular transformation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 14314-14319.	7.1	47
34	Structural and biochemical studies of a plant formamidopyrimidine-DNA glycosylase reveal why eukaryotic Fpg glycosylases do not excise 8-oxoguanine. <i>DNA Repair</i> , 2012, 11, 714-725.	2.8	46
35	The oxidative DNA glycosylases of Mycobacterium tuberculosis exhibit different substrate preferences from their Escherichia coli counterparts. <i>DNA Repair</i> , 2010, 9, 177-190.	2.8	43
36	Structural Characterization of a Viral NEIL1 Ortholog Unliganded and Bound to Abasic Site-containing DNA. <i>Journal of Biological Chemistry</i> , 2009, 284, 26174-26183.	3.4	41

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37	Genome and cancer single nucleotide polymorphisms of the human NEIL1 DNA glycosylase: Activity, structure, and the effect of editing. <i>DNA Repair</i> , 2014, 14, 17-26.	2.8	38
38	Human endonuclease VIII-like (NEIL) proteins in the giant DNA Mimivirus. <i>DNA Repair</i> , 2007, 6, 1629-1641.	2.8	36
39	A functional assay-based procedure to classify mismatch repair gene variants in Lynch syndrome. <i>Genetics in Medicine</i> , 2019, 21, 1486-1496.	2.4	36
40	Contribution of DNA unwrapping from histone octamers to the repair of oxidatively damaged DNA in nucleosomes. <i>DNA Repair</i> , 2013, 12, 964-971.	2.8	31
41	Insights into the glycosylase search for damage from single-molecule fluorescence microscopy. <i>DNA Repair</i> , 2014, 20, 23-31.	2.8	30
42	The Human Ligase III β -XRCC1 Protein Complex Performs DNA Nick Repair after Transient Unwrapping of Nucleosomal DNA. <i>Journal of Biological Chemistry</i> , 2017, 292, 5227-5238.	3.4	29
43	The Tumor-Associated Variant RAD51 G151D Induces a Hyper-Recombination Phenotype. <i>PLoS Genetics</i> , 2016, 12, e1006208.	3.5	26
44	Base Excision Repair Variants in Cancer. <i>Methods in Enzymology</i> , 2017, 591, 119-157.	1.0	26
45	Oxidative DNA Glycosylases: Recipes from Cloning to Characterization. <i>Methods in Enzymology</i> , 2006, 408, 15-33.	1.0	23
46	The lyase activity of bifunctional DNA glycosylases and the 3 β -diesterase activity of APE1 contribute to the repair of oxidized bases in nucleosomes. <i>Nucleic Acids Research</i> , 2019, 47, 2922-2931.	14.5	23
47	Unhooking of an interstrand cross-link at DNA fork structures by the DNA glycosylase NEIL3. <i>DNA Repair</i> , 2020, 86, 102752.	2.8	23
48	<i>Clostridium acetobutylicum</i> 8-Oxoguanine DNA Glycosylase (Ogg) Differs from Eukaryotic Oggs with Respect to Opposite Base Discrimination. <i>Biochemistry</i> , 2008, 47, 7626-7636.	2.5	22
49	The C-terminal Lysine of Ogg2 DNA Glycosylases is a Major Molecular Determinant for Guanine/8-Oxoguanine Distinction. <i>Journal of Molecular Biology</i> , 2010, 397, 46-56.	4.2	21
50	Structural investigation of a viral ortholog of human NEIL2/3 DNA glycosylases. <i>DNA Repair</i> , 2013, 12, 1062-1071.	2.8	21
51	Crystal Structures of Two Archaeal 8-Oxoguanine DNA Glycosylases Provide Structural Insight into Guanine/8-Oxoguanine Distinction. <i>Structure</i> , 2009, 17, 703-712.	3.3	20
52	The NEIL1 G83D germline DNA glycosylase variant induces genomic instability and cellular transformation. <i>Oncotarget</i> , 2017, 8, 85883-85895.	1.8	17
53	Structural Characterization of <i>Clostridium acetobutylicum</i> 8-Oxoguanine DNA Glycosylase in Its Apo Form and in Complex with 8-Oxodeoxyguanosine. <i>Journal of Molecular Biology</i> , 2009, 387, 669-679.	4.2	14
54	Structural basis for the lack of opposite base specificity of <i>Clostridium acetobutylicum</i> 8-oxoguanine DNA glycosylase. <i>DNA Repair</i> , 2009, 8, 1283-1289.	2.8	13

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55	Visualizing the search for radiation-damaged DNA bases in real time. <i>Radiation Physics and Chemistry</i> , 2016, 128, 126-133.	2.8	13
56	Single molecule glycosylase studies with engineered 8-oxoguanine DNA damage sites show functional defects of a MUTYH polyposis variant. <i>Nucleic Acids Research</i> , 2019, 47, 3058-3071.	14.5	13
57	Destabilization of the PCNA trimer mediated by its interaction with the NEIL1 DNA glycosylase. <i>Nucleic Acids Research</i> , 2016, 45, gkw1282.	14.5	11
58	Caught in motion: human NTHL1 undergoes interdomain rearrangement necessary for catalysis. <i>Nucleic Acids Research</i> , 2021, 49, 13165-13178.	14.5	11
59	Consequences and Repair of Oxidative DNA Damage. <i>Issues in Toxicology</i> , 2012, , 115-159.	0.1	10
60	The Phosphodiester Bond 3â€™ to a Deoxyuridine Residue Is Crucial for Substrate Binding for Uracil DNAN-Glycosylaseâ€™. <i>Biochemistry</i> , 1996, 35, 16630-16637.	2.5	6
61	Probing the activity of NTHL1 orthologs by targeting conserved amino acid residues. <i>DNA Repair</i> , 2017, 53, 43-51.	2.8	5
62	Molecular radiobiology and the origins of the base excision repair pathway: an historical perspective. <i>International Journal of Radiation Biology</i> , 2023, 99, 891-902.	1.8	5
63	Personal reflections of a woman scientist growing up in a man's world. <i>DNA Repair</i> , 2013, 12, 313-325.	2.8	1
64	Consequences and repair of radiation-induced DNA damage: fifty years of fun questions and answers. <i>International Journal of Radiation Biology</i> , 2022, 98, 367-382.	1.8	1
65	The DNA Glycosylases that Recognize and Remove Free Radical-damaged Pyrimidines. , 2017, , 117-188.		0
66	Mouse Embryonic Fibroblasts Isolated From Nthl1 D227Y Knockin Mice Exhibit Defective DNA Repair and Increased Genome Instability. <i>DNA Repair</i> , 2022, 109, 103247.	2.8	0