

Grigory L Dianov

List of Publications by Year in descending order

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57
papers

4,313
citations

101384

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h-index

143772

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59
all docs

59
docs citations

59
times ranked

5377
citing authors

#	ARTICLE	IF	CITATIONS
1	Mammalian Base Excision Repair: the Forgotten Archangel. <i>Nucleic Acids Research</i> , 2013, 41, 3483-3490.	6.5	306
2	Reconstitution of the DNA base excision repair pathway. <i>Current Biology</i> , 1994, 4, 1069-1076.	1.8	245
3	Inhibiting WEE1 Selectively Kills Histone H3K36me3-Deficient Cancers by dNTP Starvation. <i>Cancer Cell</i> , 2015, 28, 557-568.	7.7	244
4	Repair Pathways for Processing of 8-Oxoguanine in DNA by Mammalian Cell Extracts. <i>Journal of Biological Chemistry</i> , 1998, 273, 33811-33816.	1.6	220
5	Role of DNA Polymerase β in the Excision Step of Long Patch Mammalian Base Excision Repair. <i>Journal of Biological Chemistry</i> , 1999, 274, 13741-13743.	1.6	202
6	Repair of abasic sites in DNA. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2003, 531, 157-163.	0.4	170
7	Human DNA polymerase β initiates DNA synthesis during long-patch repair of reduced AP sites in DNA. <i>EMBO Journal</i> , 2001, 20, 1477-1482.	3.5	159
8	CHIP-Mediated Degradation and DNA Damage-Dependent Stabilization Regulate Base Excision Repair Proteins. <i>Molecular Cell</i> , 2008, 29, 477-487.	4.5	155
9	Poly ADP-ribose polymerase-1: An international molecule of mystery. <i>DNA Repair</i> , 2008, 7, 1077-1086.	1.3	148
10	Repair of uracil residues closely spaced on the opposite strands of plasmid DNA results in double-strand break and deletion formation. <i>Molecular Genetics and Genomics</i> , 1991, 225, 448-452.	2.4	146
11	ATM-Dependent Downregulation of USP7/HAUSP by PPM1G Activates p53 Response to DNA Damage. <i>Molecular Cell</i> , 2012, 45, 801-813.	4.5	145
12	Interaction of Human AP Endonuclease 1 with Flap Endonuclease 1 and Proliferating Cell Nuclear Antigen Involved in Long-Patch Base Excision Repair. <i>Biochemistry</i> , 2001, 40, 12639-12644.	1.2	136
13	XRCC1-DNA polymerase β interaction is required for efficient base excision repair. <i>Nucleic Acids Research</i> , 2004, 32, 2550-2555.	6.5	120
14	Ubiquitin ligase ARF-BP1/Mule modulates base excision repair. <i>EMBO Journal</i> , 2009, 28, 3207-3215.	3.5	119
15	ATM prevents DSB formation by coordinating SSB repair and cell cycle progression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 3997-4002.	3.3	105
16	USP47 Is a Deubiquitylating Enzyme that Regulates Base Excision Repair by Controlling Steady-State Levels of DNA Polymerase β . <i>Molecular Cell</i> , 2011, 41, 609-615.	4.5	102
17	A unified model for the G1/S cell cycle transition. <i>Nucleic Acids Research</i> , 2020, 48, 12483-12501.	6.5	96
18	Co-ordination of DNA single strand break repair. <i>DNA Repair</i> , 2007, 6, 454-460.	1.3	94

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19	Poly(ADP-ribose) polymerase-1 modulates DNA repair capacity and prevents formation of DNA double strand breaks. <i>DNA Repair</i> , 2008, 7, 932-940.	1.3	89
20	Poly(ADP-ribose) polymerase-1 protects excessive DNA strand breaks from deterioration during repair in human cell extracts. <i>FEBS Journal</i> , 2005, 272, 2012-2021.	2.2	85
21	Poly(ADP-ribose) polymerase in base excision repair: always engaged, but not essential for DNA damage processing. <i>Acta Biochimica Polonica</i> , 2003, 50, 169-179.	0.3	78
22	RASSF1A uncouples Wnt from Hippo signalling and promotes YAP mediated differentiation via p73. <i>Nature Communications</i> , 2018, 9, 424.	5.8	72
23	Single Nucleotide Patch Base Excision Repair Is the Major Pathway for Removal of Thymine Glycol from DNA in Human Cell Extracts. <i>Journal of Biological Chemistry</i> , 2000, 275, 11809-11813.	1.6	70
24	Co-ordination of base excision repair and genome stability. <i>DNA Repair</i> , 2013, 12, 326-333.	1.3	68
25	Regulation of oxidative DNA damage repair by DNA polymerase β and MutYH by cross-talk of phosphorylation and ubiquitination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 437-442.	3.3	67
26	DNA Damage and Repair in Schizophrenia and Autism: Implications for Cancer Comorbidity and Beyond. <i>International Journal of Molecular Sciences</i> , 2016, 17, 856.	1.8	66
27	Base excision repair fidelity in normal and cancer cells. <i>Mutagenesis</i> , 2006, 21, 173-178.	1.0	62
28	Ubiquitin ligase UBR3 regulates cellular levels of the essential DNA repair protein APE1 and is required for genome stability. <i>Nucleic Acids Research</i> , 2012, 40, 701-711.	6.5	53
29	DNA Polymerase β Promotes Recruitment of DNA Ligase III α XRCC1 to Sites of Base Excision Repair. <i>Biochemistry</i> , 2005, 44, 10613-10619.	1.2	49
30	Targeting base excision repair to improve cancer therapies. <i>Molecular Aspects of Medicine</i> , 2007, 28, 345-374.	2.7	46
31	Overexpression of DNA polymerase β results in an increased rate of frameshift mutations during base excision repair. <i>Mutagenesis</i> , 2007, 22, 183-188.	1.0	43
32	Phosphorylation of PNKP by ATM prevents its proteasomal degradation and enhances resistance to oxidative stress. <i>Nucleic Acids Research</i> , 2012, 40, 11404-11415.	6.5	42
33	ATMIN is a transcriptional regulator of both lung morphogenesis and ciliogenesis. <i>Development (Cambridge)</i> , 2014, 141, 3966-3977.	1.2	40
34	E2F1 proteolysis via SCF β -cyclin F underlies synthetic lethality between cyclin F loss and Chk1 inhibition. <i>EMBO Journal</i> , 2019, 38, e101443.	3.5	40
35	Cells deficient in base-excision repair reveal cancer hallmarks originating from adjustments to genetic instability. <i>Nucleic Acids Research</i> , 2015, 43, 3667-3679.	6.5	39
36	p53 coordinates base excision repair to prevent genomic instability. <i>Nucleic Acids Research</i> , 2016, 44, 3165-3175.	6.5	39

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37	USP7S-dependent inactivation of Mule regulates DNA damage signalling and repair. <i>Nucleic Acids Research</i> , 2013, 41, 1750-1756.	6.5	34
38	Monitoring base excision repair by in vitro assays. <i>Toxicology</i> , 2003, 193, 35-41.	2.0	32
39	AKT regulates NPM dependent ARF localization and p53mut stability in tumors. <i>Oncotarget</i> , 2014, 5, 6142-6167.	0.8	30
40	ARF induction in response to DNA strand breaks is regulated by PARP1. <i>Nucleic Acids Research</i> , 2014, 42, 2320-2329.	6.5	27
41	The NUCKS1-SKP2-p21/p27 axis controls S phase entry. <i>Nature Communications</i> , 2021, 12, 6959.	5.8	24
42	Interplay between base excision repair protein XRCC1 and ALDH2 predicts overall survival in lung and liver cancer patients. <i>Cellular Oncology (Dordrecht)</i> , 2018, 41, 527-539.	2.1	23
43	Sp1 phosphorylation by ATM downregulates BER and promotes cell elimination in response to persistent DNA damage. <i>Nucleic Acids Research</i> , 2018, 46, 1834-1846.	6.5	22
44	Impaired oxidative stress response characterizes HUWE1-promoted X-linked intellectual disability. <i>Scientific Reports</i> , 2017, 7, 15050.	1.6	21
45	End-damage-specific proteins facilitate recruitment or stability of X-ray cross-complementing protein 1 at the sites of DNA single-strand break repair. <i>FEBS Journal</i> , 2005, 272, 5753-5763.	2.2	20
46	Persistent DNA strand breaks induce a CAF-like phenotype in normal fibroblasts. <i>Oncotarget</i> , 2018, 9, 13666-13681.	0.8	20
47	The emerging role of Mule and ARF in the regulation of base excision repair. <i>FEBS Letters</i> , 2011, 585, 2831-2835.	1.3	18
48	2.3 THz radiation: Absence of genotoxicity/mutagenicity in Escherichia coli and Salmonella typhimurium. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2016, 803-804, 34-38.	0.9	15
49	Src-mediated phosphorylation of GAPDH regulates its nuclear localization and cellular response to DNA damage. <i>FASEB Journal</i> , 2020, 34, 10443-10461.	0.2	15
50	Modulation of proteostasis counteracts oxidative stress and affects DNA base excision repair capacity in ATM-deficient cells. <i>Nucleic Acids Research</i> , 2017, 45, 10042-10055.	6.5	13
51	Two-way crosstalk between BER and NHEJ repair pathway is mediated by Pol δ and Ku70. <i>FASEB Journal</i> , 2019, 33, 11668-11681.	0.2	12
52	DNA Base Excision Repair: The Achilles' Heel of Tumour Cells and their Microenvironment?. <i>Current Pharmaceutical Design</i> , 2017, 23, 4758-4772.	0.9	10
53	Base excision repair targets for cancer therapy. <i>American Journal of Cancer Research</i> , 2011, 1, 845-51.	1.4	8
54	Mild phenotype of knockouts of the major apurinic/apyrimidinic endonuclease APEX1 in a non-cancer human cell line. <i>PLoS ONE</i> , 2021, 16, e0257473.	1.1	4

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55	Sp1-independent downregulation of NHEJ in response to BER deficiency. DNA Repair, 2020, 86, 102740.	1.3	2
56	The role of Sp1 in the detection and elimination of cells with persistent DNA strand breaks. NAR Cancer, 2020, 2, zcaa004.	1.6	2
57	Boron-containing nucleosides as tools for boron-neutron capture therapy. American Journal of Cancer Research, 2021, 11, 4668-4682.	1.4	0