

Fumio Hanaoka

List of Publications by Year in descending order

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54
papers

5,735
citations

236925

25
h-index

197818

49
g-index

55
all docs

55
docs citations

55
times ranked

3951
citing authors

#	ARTICLE	IF	CITATIONS
1	The XPV (xeroderma pigmentosum variant) gene encodes human DNA polymerase $\hat{\iota}$. <i>Nature</i> , 1999, 399, 700-704.	27.8	1,248
2	UV-Induced Ubiquitylation of XPC Protein Mediated by UV-DDB-Ubiquitin Ligase Complex. <i>Cell</i> , 2005, 121, 387-400.	28.9	517
3	The Molecular Basis of CRL4DDB2/CSA Ubiquitin Ligase Architecture, Targeting, and Activation. <i>Cell</i> , 2011, 147, 1024-1039.	28.9	372
4	Low fidelity DNA synthesis by human DNA polymerase $\hat{\iota}$. <i>Nature</i> , 2000, 404, 1011-1013.	27.8	356
5	Structure and mechanism of human DNA polymerase $\hat{\iota}$. <i>Nature</i> , 2010, 465, 1044-1048.	27.8	300
6	Centrosome Protein Centrin 2/Caltractin 1 Is Part of the Xeroderma Pigmentosum Group C Complex That Initiates Global Genome Nucleotide Excision Repair. <i>Journal of Biological Chemistry</i> , 2001, 276, 18665-18672.	3.4	290
7	Error-prone bypass of certain DNA lesions by the human DNA polymerase $\hat{\iota}$. <i>Genes and Development</i> , 2000, 14, 1589-1594.	5.9	250
8	Interaction of hREV1 with three human Y-family DNA polymerases. <i>Genes To Cells</i> , 2004, 9, 523-531.	1.2	244
9	Interaction of hHR23 with S5a. <i>Journal of Biological Chemistry</i> , 1999, 274, 28019-28025.	3.4	243
10	Efficient Translesion Replication Past Oxaliplatin and Cisplatin GpG Adducts by Human DNA Polymerase $\hat{\iota}$. <i>Biochemistry</i> , 2000, 39, 4575-4580.	2.5	209
11	Misinsertion and bypass of thymine-thymine dimers by human DNA polymerase $\hat{\iota}$. <i>EMBO Journal</i> , 2000, 19, 5259-5266.	7.8	197
12	129-derived Strains of Mice Are Deficient in DNA Polymerase $\hat{\iota}$ and Have Normal Immunoglobulin Hypermutation. <i>Journal of Experimental Medicine</i> , 2003, 198, 635-643.	8.5	169
13	Molecular analysis of mutations in DNA polymerase $\hat{\iota}$ in xeroderma pigmentosum-variant patients. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 815-820.	7.1	165
14	3-Methyladenine-DNA Glycosylase (MPG Protein) Interacts with Human RAD23 Proteins. <i>Journal of Biological Chemistry</i> , 2000, 275, 28433-28438.	3.4	109
15	UV-B Radiation Induces Epithelial Tumors in Mice Lacking DNA Polymerase $\hat{\iota}$ and Mesenchymal Tumors in Mice Deficient for DNA Polymerase $\hat{\iota}$. <i>Molecular and Cellular Biology</i> , 2006, 26, 7696-7706.	2.3	102
16	Developmental Defects and Male Sterility in Mice Lacking the Ubiquitin-Like DNA Repair Gene mHR23B. <i>Molecular and Cellular Biology</i> , 2002, 22, 1233-1245.	2.3	99
17	Systematic identification, classification, and characterization of the open reading frames which encode novel helicase-related proteins in <i>Saccharomyces cerevisiae</i> by gene disruption and Northern analysis. <i>Genetics</i> , 1999, 15, 219-253.		90
18	Cloning, Comparative Mapping, and RNA Expression of the Mouse Homologues of the <i>Saccharomyces cerevisiae</i> Nucleotide Excision Repair Gene RAD23. <i>Genomics</i> , 1996, 31, 20-27.	2.9	66

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19	Relative levels of the two mammalian Rad23 homologs determine composition and stability of the xeroderma pigmentosum group C protein complex. <i>DNA Repair</i> , 2004, 3, 1285-1295.	2.8	63
20	Genomic structure, chromosomal localization and identification of mutations in the xeroderma pigmentosum variant (XPV) gene. <i>Oncogene</i> , 2000, 19, 4721-4728.	5.9	58
21	Sequence context-dependent replication of DNA templates containing UV-induced lesions by human DNA polymerase $\hat{\iota}$. <i>DNA Repair</i> , 2003, 2, 991-1006.	2.8	54
22	Interaction with DNA polymerase $\hat{\iota}$ is required for nuclear accumulation of REV1 and suppression of spontaneous mutations in human cells. <i>DNA Repair</i> , 2009, 8, 585-599.	2.8	53
23	Atomic force microscopy sees nucleosome positioning and histone H1-induced compaction in reconstituted chromatin. <i>FEBS Letters</i> , 1999, 452, 267-271.	2.8	47
24	Comparison of Incorporation and Extension of Nucleotides in vitro opposite 8-Hydroxyguanine (7,8-Dihydro-8-oxoguanine) in Hot Spots of the c-Ha-ras Gene. <i>Japanese Journal of Cancer Research</i> , 1995, 86, 270-276.	1.7	31
25	SUMOylation of xeroderma pigmentosum group C protein regulates DNA damage recognition during nucleotide excision repair. <i>Scientific Reports</i> , 2015, 5, 10984.	3.3	31
26	Chemical synthesis and translesion replication of a cis-syn cyclobutane thymine-uracil dimer. <i>Nucleic Acids Research</i> , 2004, 32, 1738-1745.	14.5	30
27	Expression Profiles of Transcripts from 126 Open Reading Frames in the Entire Chromosome VI of <i>Saccharomyces cerevisiae</i> by Systematic Northern Analyses. , 1997, 13, 1275-1290.		26
28	Two mammalian homologs of yeast Rad23, HR23A and HR23B, as multifunctional proteins. <i>Gene</i> , 2017, 597, 1-9.	2.2	26
29	Eukaryotic DNA Polymerases $\hat{\iota}$, $\hat{\iota}^2$ and $\hat{\iota}^3$ Incorporate Guanine Opposite 2,2,4-triamino-5(2H)-oxazolone. <i>ChemBioChem</i> , 2009, 10, 2613-2616.	2.6	25
30	Novel function of HATs and HDACs in homologous recombination through acetylation of human RAD52 at double-strand break sites. <i>PLoS Genetics</i> , 2018, 14, e1007277.	3.5	25
31	DNA polymerases $\hat{\iota}$ and $\hat{\iota}^2$ are responsible for error-free translesion DNA synthesis activity over a cis-syn thymine dimer in <i>Xenopus laevis</i> oocyte extracts. <i>DNA Repair</i> , 2005, 4, 1252-1269.	2.8	24
32	Cyclobutane thymine dimers in arasproto-oncogene hot spot activate the gene by point mutation. <i>Nucleic Acids Research</i> , 1993, 21, 2355-2361.	14.5	22
33	A DNA oligomer containing 2,2,4-triamino-5(2H)-oxazolone is incised by human NEIL1 and NTH1. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2012, 734, 73-77.	1.0	20
34	UV-induced mutations in epidermal cells of mice defective in DNA polymerase $\hat{\iota}$ and/or $\hat{\iota}^1$. <i>DNA Repair</i> , 2015, 29, 139-146.	2.8	19
35	Molecular cloning of the cDNA for the catalytic subunit of plant DNA polymerase $\hat{\iota}$ and its cell-cycle dependent expression. <i>Genes To Cells</i> , 1997, 2, 695-709.	1.2	17
36	Translesion synthesis by human DNA polymerase $\hat{\iota}$ across oxidative products of guanine. <i>Nucleic Acids Symposium Series</i> , 2004, 48, 171-172.	0.3	17

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37	<i>Schizosaccharomyces pombe</i> Ddb1 Recruits Substrate-Specific Adaptor Proteins through a Novel Protein Motif, the DDB-Box. <i>Molecular and Cellular Biology</i> , 2008, 28, 6746-6756.	2.3	17
38	The Human RNA Helicase A (DDX9) Gene Maps to the Prostate Cancer Susceptibility Locus at Chromosome Band 1q25 and Its Pseudogene (DDX9P) to 13q22, Respectively. <i>Somatic Cell and Molecular Genetics</i> , 1999, 25, 33-39.	0.7	16
39	Stalled Pol β at its cognate substrate initiates an alternative translesion synthesis pathway via interaction with REV1. <i>Genes To Cells</i> , 2012, 17, 98-108.	1.2	16
40	Two budding yeast RAD4 homologs in fission yeast play different roles in the repair of UV-induced DNA damage. <i>DNA Repair</i> , 2002, 1, 833-845.	2.8	12
41	The Protein Level of Rev1, a TLS Polymerase in Fission Yeast, Is Strictly Regulated during the Cell Cycle and after DNA Damage. <i>PLoS ONE</i> , 2015, 10, e0130000.	2.5	11
42	E2F regulates growth-dependent transcription of genes encoding both catalytic and regulatory subunits of mouse primase. <i>Genes To Cells</i> , 2001, 6, 57-70.	1.2	8
43	SOS polymerases. <i>Nature</i> , 2001, 409, 33-34.	27.8	8
44	Cloning and Characterization of Novel Gene, DCRR1, Expressed from Down's Syndrome Critical Region of Human Chromosome 21q22.2. <i>DNA Sequence</i> , 1997, 7, 153-164.	0.7	7
45	Analysis of a 36.2 kb DNA sequence including the right telomere of chromosome VI from <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 1996, 12, 149-167.	1.7	6
46	Hypersensitivity of mouse embryonic fibroblast cells defective for DNA polymerases β , δ and ϵ to various genotoxic compounds: Its potential for application in chemical genotoxic screening. <i>DNA Repair</i> , 2018, 61, 76-85.	2.8	5
47	Fifteen open reading frames in a 30.8 kb region of the right arm of chromosome VI from <i>Saccharomyces cerevisiae</i> . <i>Yeast</i> , 1996, 12, 177-190.	1.7	4
48	Temperature-sensitive Mutation of DNA Polymerase α . Induces Growth-suppressive Phenotypes Involving Retinoblastoma Protein and Cyclin D1. <i>Cell Structure and Function</i> , 1995, 20, 285-291.	1.1	4
49	Translesion DNA Synthesis and Damage Tolerance Pathways. , 2016, , 249-304.		3
50	Identification of the Coding Region of <i>Saccharomyces cerevisiae</i> Chromosome VI Using the Computer Program GenMark. <i>DNA Research</i> , 1995, 2, 247-253.	3.4	2
51	Molecular biology of Fanconi anaemia "an old problem, a new insight. <i>BioEssays</i> , 2002, 24, 439-448.	2.5	1
52	Resolution and Characterization of Polymorphic DNA by SSCP and Chemical Cleavage Methodologies. <i>Journal of Radiation Research</i> , 0, , .	1.6	0
53	Trapping DNA Replication Origins from the Human Genome. <i>Genes</i> , 2013, 4, 198-225.	2.4	0
54	Translesion DNA Synthesis. , 2019, , 169-189.		0