

# Thanos D Halazonetis

## List of Publications by Year in descending order

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Version: 2024-02-01

48  
papers

11,929  
citations

172457

29  
h-index

223800

46  
g-index

49  
all docs

49  
docs citations

49  
times ranked

14571  
citing authors

#	ARTICLE	IF	CITATIONS
1	Activation of the DNA damage checkpoint and genomic instability in human precancerous lesions. <i>Nature</i> , 2005, 434, 907-913.	27.8	1,870
2	Genomic instability "an evolving hallmark of cancer. <i>Nature Reviews Molecular Cell Biology</i> , 2010, 11, 220-228.	37.0	1,798
3	Oncogene-induced senescence is part of the tumorigenesis barrier imposed by DNA damage checkpoints. <i>Nature</i> , 2006, 444, 633-637.	27.8	1,777
4	An Oncogene-Induced DNA Damage Model for Cancer Development. <i>Science</i> , 2008, 319, 1352-1355.	12.6	1,612
5	Methylated lysine 79 of histone H3 targets 53BP1 to DNA double-strand breaks. <i>Nature</i> , 2004, 432, 406-411.	27.8	815
6	P53 Binding Protein 1 (53bp1) Is an Early Participant in the Cellular Response to DNA Double-Strand Breaks. <i>Journal of Cell Biology</i> , 2000, 151, 1381-1390.	5.2	801
7	DNA Replication Stress as a Hallmark of Cancer. <i>Annual Review of Pathology: Mechanisms of Disease</i> , 2015, 10, 425-448.	22.4	593
8	Break-Induced Replication Repair of Damaged Forks Induces Genomic Duplications in Human Cells. <i>Science</i> , 2014, 343, 88-91.	12.6	387
9	53BP1 functions in an ATM-dependent checkpoint pathway that is constitutively activated in human cancer. <i>Nature Cell Biology</i> , 2002, 4, 998-1002.	10.3	386
10	Intragenic origins due to short G1 phases underlie oncogene-induced DNA replication stress. <i>Nature</i> , 2018, 555, 112-116.	27.8	303
11	Mammalian RAD52 Functions in Break-Induced Replication Repair of Collapsed DNA Replication Forks. <i>Molecular Cell</i> , 2016, 64, 1127-1134.	9.7	223
12	Alternative lengthening of human telomeres is a conservative DNA replication process with features of break-induced replication. <i>EMBO Reports</i> , 2016, 17, 1731-1737.	4.5	133
13	An Oligomerized 53BP1 Tudor Domain Suffices for Recognition of DNA Double-Strand Breaks. <i>Molecular and Cellular Biology</i> , 2009, 29, 1050-1058.	2.3	104
14	Impaired liver regeneration in aged mice can be rescued by silencing Hippo core kinases MST1 and MST2. <i>EMBO Molecular Medicine</i> , 2017, 9, 46-60.	6.9	98
15	An induced fit mechanism regulates p53 DNA binding kinetics to confer sequence specificity. <i>EMBO Journal</i> , 2011, 30, 2167-2176.	7.8	95
16	ATM signaling and 53BP1. <i>Radiotherapy and Oncology</i> , 2005, 76, 119-122.	0.6	83
17	TopBP1 functions with 53BP1 in the G1 DNA damage checkpoint. <i>EMBO Journal</i> , 2010, 29, 3723-3732.	7.8	79
18	High-resolution mapping of mitotic DNA synthesis regions and common fragile sites in the human genome through direct sequencing. <i>Cell Research</i> , 2020, 30, 997-1008.	12.0	74

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19	Studies of genomic copy number changes in human cancers reveal signatures of DNA replication stress. <i>Molecular Oncology</i> , 2011, 5, 308-314.	4.6	69
20	A multi-pronged approach targeting SARS-CoV-2 proteins using ultra-large virtual screening. <i>IScience</i> , 2021, 24, 102021.	4.1	66
21	A Single-Nucleotide Substitution Mutator Phenotype Revealed by Exome Sequencing of Human Colon Adenomas. <i>Cancer Research</i> , 2012, 72, 6279-6289.	0.9	61
22	R&#euml;spodin 1 and noggin facilitate expansion of resident stem cells from non&#euml;damaged gallbladders. <i>EMBO Reports</i> , 2016, 17, 769-779.	4.5	53
23	Structural Differences in the DNA Binding Domains of Human p53 and Its <i>C. elegans</i> Ortholog Cep-1. <i>Structure</i> , 2004, 12, 1237-1243.	3.3	51
24	DNA replication stress as an Achilles' heel of cancer. <i>Oncotarget</i> , 2015, 6, 1-2.	1.8	50
25	Crystal Structure of a Multidomain Human p53 Tetramer Bound to the Natural <i>CDKN1A</i> (<i>p21</i>) p53-Response Element. <i>Molecular Cancer Research</i> , 2011, 9, 1493-1499.	3.4	49
26	Non-covalent SARS-CoV-2 Mpro inhibitors developed from in silico screen hits. <i>Scientific Reports</i> , 2022, 12, 2505.	3.3	41
27	Constitutively active DNA damage checkpoint pathways as the driving force for the high frequency of p53 mutations in human cancer. <i>DNA Repair</i> , 2004, 3, 1057-1062.	2.8	37
28	Reversal of the DNA-Binding-Induced Loop L1 Conformational Switch in an Engineered Human p53 Protein. <i>Journal of Molecular Biology</i> , 2014, 426, 936-944.	4.2	36
29	POLD3 Is Haploinsufficient for DNA Replication in Mice. <i>Molecular Cell</i> , 2016, 63, 877-883.	9.7	34
30	The helicase domain and C-terminus of human RecQL4 facilitate replication elongation on DNA templates damaged by ionizing radiation. <i>Carcinogenesis</i> , 2012, 33, 1203-1210.	2.8	27
31	Monitoring early S-phase origin firing and replication fork movement by sequencing nascent DNA from synchronized cells. <i>Nature Protocols</i> , 2019, 14, 51-67.	12.0	21
32	Enhanced Rate of Acquisition of Point Mutations in Mouse Intestinal Adenomas Compared to Normal Tissue. <i>Cell Reports</i> , 2017, 19, 2185-2192.	6.4	18
33	Change in oligomerization specificity of the p53 tetramerization domain by hydrophobic amino acid substitutions. <i>Protein Science</i> , 1999, 8, 1773-1779.	7.6	16
34	The role of SMARCAL1 in replication fork stability and telomere maintenance. <i>DNA Repair</i> , 2017, 56, 129-134.	2.8	13
35	Ubiquitin-H2AX fusions render 53BP1 recruitment to DNA damage sites independent of RNF8 or RNF168. <i>Cell Cycle</i> , 2015, 14, 1748-1758.	2.6	10
36	DNA Damage Signaling Recruits the RNA Polymerase II Binding Protein Che-1 to the p53 Promoter. <i>Molecular Cell</i> , 2006, 24, 809-810.	9.7	8

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37	Increased Cell Proliferation and Gene Expression of Genes Related to Bone Remodeling, Cell Adhesion and Collagen Metabolism in the Periodontal Ligament of Unopposed Molars in Growing Rats. <i>Frontiers in Physiology</i> , 2017, 8, 75.	2.8	7
38	A transcription-based mechanism for oncogenic $\beta$ -catenin-induced lethality in BRCA1/2-deficient cells. <i>Nature Communications</i> , 2021, 12, 4919.	12.8	6
39	Genomic Instability Profiles at the Single Cell Level in Mouse Colorectal Cancers of Defined Genotypes. <i>Cancers</i> , 2021, 13, 1267.	3.7	5
40	Conservative DNA Replication. <i>Nature Reviews Molecular Cell Biology</i> , 2014, 15, 300-300.	37.0	4
41	Remodeling Collapsed DNA Replication Forks for Cancer Development. <i>Cancer Research</i> , 2019, 79, 1297-1298.	0.9	4
42	SAHF, to senesce or not to senesce?. <i>Cell Cycle</i> , 2011, 10, 741-740.	2.6	3
43	Delayed DNA break repair for genome stability. <i>Nature Cell Biology</i> , 2021, 23, 1055-1057.	10.3	3
44	A Model to Investigate Single-Strand DNA Responses in G1 Human Cells via a Telomere-Targeted, Nuclease-Deficient CRISPR-Cas9 System. <i>PLoS ONE</i> , 2017, 12, e0169126.	2.5	2
45	Draining the FEN1s for cancer therapy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 21849-21850.	7.1	1
46	A method to sequence genomic sites of mitotic DNA synthesis in mammalian cells. <i>Methods in Enzymology</i> , 2021, 661, 283-304.	1.0	1
47	DNA Replication & DNA Replication Stress. <i>ACS in Focus</i> , 2022, , .	0.6	0
48	Beating cancer one carbon at a time. <i>Nature Cancer</i> , 2022, 3, 141-142.	13.2	0