

Fabrizio d'Adda di Fagagna

List of Publications by Year in descending order

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90
papers

18,317
citations

38660

50
h-index

45213

90
g-index

98
all docs

98
docs citations

98
times ranked

20410
citing authors

#	ARTICLE	IF	CITATIONS
1	Cellular senescence: when bad things happen to good cells. <i>Nature Reviews Molecular Cell Biology</i> , 2007, 8, 729-740.	16.1	3,502
2	A DNA damage checkpoint response in telomere-initiated senescence. <i>Nature</i> , 2003, 426, 194-198.	13.7	2,381
3	Oncogene-induced senescence is a DNA damage response triggered by DNA hyper-replication. <i>Nature</i> , 2006, 444, 638-642.	13.7	1,576
4	Chemokine Signaling via the CXCR2 Receptor Reinforces Senescence. <i>Cell</i> , 2008, 133, 1006-1018.	13.5	1,446
5	Living on a break: cellular senescence as a DNA-damage response. <i>Nature Reviews Cancer</i> , 2008, 8, 512-522.	12.8	866
6	Cellular senescence in ageing: from mechanisms to therapeutic opportunities. <i>Nature Reviews Molecular Cell Biology</i> , 2021, 22, 75-95.	16.1	812
7	Telomeric DNA damage is irreparable and causes persistent DNA-damage-response activation. <i>Nature Cell Biology</i> , 2012, 14, 355-365.	4.6	646
8	Site-specific DICER and DROSHA RNA products control the DNA-damage response. <i>Nature</i> , 2012, 488, 231-235.	13.7	460
9	Human cell senescence as a DNA damage response. <i>Mechanisms of Ageing and Development</i> , 2005, 126, 111-117.	2.2	383
10	Interplay between oncogene-induced DNA damage response and heterochromatin in senescence and cancer. <i>Nature Cell Biology</i> , 2011, 13, 292-302.	4.6	294
11	Damage-induced lncRNAs control the DNA damage response through interaction with DDRNAs at individual double-strand breaks. <i>Nature Cell Biology</i> , 2017, 19, 1400-1411.	4.6	288
12	Identification of a DNA Nonhomologous End-Joining Complex in Bacteria. <i>Science</i> , 2002, 297, 1686-1689.	6.0	284
13	Effects of DNA nonhomologous end-joining factors on telomere length and chromosomal stability in mammalian cells. <i>Current Biology</i> , 2001, 11, 1192-1196.	1.8	260
14	Oncogene-induced reactive oxygen species fuel hyperproliferation and DNA damage response activation. <i>Cell Death and Differentiation</i> , 2014, 21, 998-1012.	5.0	254
15	Functional links between telomeres and proteins of the DNA-damage response. <i>Genes and Development</i> , 2004, 18, 1781-1799.	2.7	244
16	Functional transcription promoters at DNA double-strand breaks mediate RNA-driven phase separation of damage-response factors. <i>Nature Cell Biology</i> , 2019, 21, 1286-1299.	4.6	233
17	Functions of poly(ADP-ribose) polymerase in controlling telomere length and chromosomal stability. <i>Nature Genetics</i> , 1999, 23, 76-80.	9.4	218
18	Oncogene-induced telomere dysfunction enforces cellular senescence in human cancer precursor lesions. <i>EMBO Journal</i> , 2012, 31, 2839-2851.	3.5	200

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19	Telomere dysfunction in ageing and age-related diseases. <i>Nature Cell Biology</i> , 2022, 24, 135-147.	4.6	194
20	Crosstalk between chromatin state and DNA damage response in cellular senescence and cancer. <i>Nature Reviews Cancer</i> , 2012, 12, 709-720.	12.8	181
21	Interaction of HIV-1 Tat Protein with Heparin. <i>Journal of Biological Chemistry</i> , 1997, 272, 11313-11320.	1.6	179
22	BRCA2 controls DNA:RNA hybrid level at DSBs by mediating RNase H2 recruitment. <i>Nature Communications</i> , 2018, 9, 5376.	5.8	176
23	Activation of transcription factor NF-kappaB by the Tat protein of human immunodeficiency virus type 1. <i>Journal of Virology</i> , 1996, 70, 4427-4437.	1.5	136
24	A direct role for small non-coding RNAs in DNA damage response. <i>Trends in Cell Biology</i> , 2014, 24, 171-178.	3.6	114
25	Stable Cellular Senescence Is Associated with Persistent DDR Activation. <i>PLoS ONE</i> , 2014, 9, e110969.	1.1	110
26	Irreparable telomeric DNA damage and persistent DDR signalling as a shared causative mechanism of cellular senescence and ageing. <i>Current Opinion in Genetics and Development</i> , 2014, 26, 89-95.	1.5	106
27	Cellular senescence: hot or what?. <i>Current Opinion in Genetics and Development</i> , 2009, 19, 25-31.	1.5	103
28	DICER, DROSHA and DNA damage-response RNAs are necessary for the secondary recruitment of DNA damage response factors. <i>Journal of Cell Science</i> , 2016, 129, 1468-76.	1.2	99
29	SASPense and DDRama in cancer and ageing. <i>Nature Cell Biology</i> , 2009, 11, 921-923.	4.6	96
30	Cleavage and Inactivation of ATM during Apoptosis. <i>Molecular and Cellular Biology</i> , 1999, 19, 6076-6084.	1.1	95
31	A human binding site for transcription factor USF/MLTF mimics the negative regulatory element of human immunodeficiency virus type 1. <i>Virology</i> , 1992, 186, 133-147.	1.1	94
32	A damaged genome's transcriptional landscape through multilayered expression profiling around in situ-mapped DNA double-strand breaks. <i>Nature Communications</i> , 2017, 8, 15656.	5.8	89
33	Inhibition of DNA damage response at telomeres improves the detrimental phenotypes of Hutchinson Gilford Progeria Syndrome. <i>Nature Communications</i> , 2019, 10, 4990.	5.8	85
34	Yeast Nhp6A/B and Mammalian Hmgb1 Facilitate the Maintenance of Genome Stability. <i>Current Biology</i> , 2005, 15, 68-72.	1.8	84
35	DNA Damage in Mammalian Neural Stem Cells Leads to Astrocytic Differentiation Mediated by BMP2 Signaling through JAK-STAT. <i>Stem Cell Reports</i> , 2013, 1, 123-138.	2.3	79
36	Polycomb proteins control proliferation and transformation independently of cell cycle checkpoints by regulating DNA replication. <i>Nature Communications</i> , 2014, 5, 3649.	5.8	79

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37	Expression of H-RASV12 in a zebrafish model of Costello syndrome causes cellular senescence in adult proliferating cells. <i>DMM Disease Models and Mechanisms</i> , 2009, 2, 56-67.	1.2	77
38	The Gam protein of bacteriophage Mu is an orthologue of eukaryotic Ku. <i>EMBO Reports</i> , 2003, 4, 47-52.	2.0	76
39	DNA damage response activation in mouse embryonic fibroblasts undergoing replicative senescence and following spontaneous immortalization. <i>Cell Cycle</i> , 2008, 7, 3601-3606.	1.3	76
40	DNA damage response inhibition at dysfunctional telomeres by modulation of telomeric DNA damage response RNAs. <i>Nature Communications</i> , 2017, 8, 13980.	5.8	76
41	Complex engagement of DNA damage response pathways in human cancer and in lung tumor progression. <i>Carcinogenesis</i> , 2007, 28, 2082-2088.	1.3	74
42	Breaking news: high-speed race ends in arrest – how oncogenes induce senescence. <i>Trends in Cell Biology</i> , 2007, 17, 529-536.	3.6	73
43	Notch is a direct negative regulator of the DNA-damage response. <i>Nature Structural and Molecular Biology</i> , 2015, 22, 417-424.	3.6	68
44	Differential regulation of DNA damage response activation between somatic and germline cells in <i>Caenorhabditis elegans</i> . <i>Cell Death and Differentiation</i> , 2012, 19, 1847-1855.	5.0	65
45	Telomerase abrogates aneuploidy-induced telomere replication stress, senescence and cell depletion. <i>EMBO Journal</i> , 2015, 34, 1371-1384.	3.5	65
46	From “Cellular” RNA to “Smart” RNA: Multiple Roles of RNA in Genome Stability and Beyond. <i>Chemical Reviews</i> , 2018, 118, 4365-4403.	23.0	63
47	Is cellular senescence an example of antagonistic pleiotropy?. <i>Aging Cell</i> , 2012, 11, 378-383.	3.0	62
48	Activation of the DNA Damage Response by Telomere Attrition: A Passage to Cellular Senescence. <i>Cell Cycle</i> , 2004, 3, 541-544.	1.3	61
49	Terminally differentiated astrocytes lack DNA damage response signaling and are radioresistant but retain DNA repair proficiency. <i>Cell Death and Differentiation</i> , 2012, 19, 582-591.	5.0	58
50	A novel single-cell method provides direct evidence of persistent DNA damage in senescent cells and aged mammalian tissues. <i>Aging Cell</i> , 2017, 16, 422-427.	3.0	56
51	Molecular and functional interactions of transcription factor USF with the long terminal repeat of human immunodeficiency virus type 1. <i>Journal of Virology</i> , 1995, 69, 2765-2775.	1.5	54
52	NOTCH1 Inhibits Activation of ATM by Impairing the Formation of an ATM-FOXO3a-KAT5/Tip60 Complex. <i>Cell Reports</i> , 2016, 16, 2068-2076.	2.9	53
53	Transcription and DNA Damage: Holding Hands or Crossing Swords?. <i>Journal of Molecular Biology</i> , 2017, 429, 3215-3229.	2.0	52
54	Telomere and Telomerase Modulation by the Mammalian Rad9/Rad1/Hus1 DNA-Damage-Checkpoint Complex. <i>Current Biology</i> , 2006, 16, 1551-1558.	1.8	50

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55	Pharmacological boost of DNA damage response and repair by enhanced biogenesis of DNA damage response RNAs. <i>Scientific Reports</i> , 2019, 9, 6460.	1.6	49
56	Telomere transcription in ageing. <i>Ageing Research Reviews</i> , 2020, 62, 101115.	5.0	44
57	MRE11-RAD50-NBS1 Complex Is Sufficient to Promote Transcription by RNA Polymerase II at Double-Strand Breaks by Melting DNA Ends. <i>Cell Reports</i> , 2021, 34, 108565.	2.9	43
58	Human replication protein Cdc6 is selectively cleaved by caspase 3 during apoptosis. <i>EMBO Reports</i> , 2002, 3, 780-784.	2.0	39
59	Recent Advancements in DNA Damage—Transcription Crosstalk and High-Resolution Mapping of DNA Breaks. <i>Annual Review of Genomics and Human Genetics</i> , 2017, 18, 87-113.	2.5	37
60	DNA Damage Triggers a New Phase in Neurodegeneration. <i>Trends in Genetics</i> , 2021, 37, 337-354.	2.9	37
61	Epigenetic alterations associated with cellular senescence: A barrier against tumorigenesis or a red carpet for cancer?. <i>Seminars in Cancer Biology</i> , 2011, 21, 360-366.	4.3	35
62	RNA processing proteins regulate Mec1/ATR activation by promoting generation of RPA-coated ssDNA. <i>EMBO Reports</i> , 2015, 16, 221-231.	2.0	32
63	Telomere damage promotes vascular smooth muscle cell senescence and immune cell recruitment after vessel injury. <i>Communications Biology</i> , 2021, 4, 611.	2.0	32
64	Transcriptional and post-transcriptional regulation of the ionizing radiation response by ATM and p53. <i>Scientific Reports</i> , 2017, 7, 43598.	1.6	31
65	Express or repress? The transcriptional dilemma of damaged chromatin. <i>FEBS Journal</i> , 2017, 284, 2133-2147.	2.2	28
66	Neural stem cells exposed to BrdU lose their global DNA methylation and undergo astrocytic differentiation. <i>Nucleic Acids Research</i> , 2012, 40, 5332-5342.	6.5	26
67	Resection is responsible for loss of transcription around a double-strand break in <i>Saccharomyces cerevisiae</i> . <i>ELife</i> , 2015, 4, .	2.8	26
68	Cleavage of the Bloom's syndrome gene product during apoptosis by caspase-3 results in an impaired interaction with topoisomerase IIIalpha. <i>Nucleic Acids Research</i> , 2001, 29, 3172-3180.	6.5	25
69	Stimulation of the adenovirus major late promoter in vitro by transcription factor USF is enhanced by the adenovirus DNA binding protein. <i>Journal of Virology</i> , 1994, 68, 8288-8295.	1.5	25
70	Never-ageing cellular senescence. <i>European Journal of Cancer</i> , 2011, 47, 1616-1622.	1.3	24
71	DNA damage response at telomeres boosts the transcription of SARS-CoV-2 receptor ACE2 during aging. <i>EMBO Reports</i> , 2022, 23, e53658.	2.0	24
72	PREP1 tumor suppressor protects the late-replicating DNA by controlling its replication timing and symmetry. <i>Scientific Reports</i> , 2018, 8, 3198.	1.6	18

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73	The prolyl-isomerase PIN1 is essential for nuclear Lamin-B structure and function and protects heterochromatin under mechanical stress. <i>Cell Reports</i> , 2021, 36, 109694.	2.9	15
74	Need telomere maintenance? Call 911. <i>Cell Division</i> , 2007, 2, 3.	1.1	11
75	The cohesin complex prevents Myc-induced replication stress. <i>Cell Death and Disease</i> , 2017, 8, e2956-e2956.	2.7	11
76	BRCA1 deficiency specific base substitution mutagenesis is dependent on translesion synthesis and regulated by 53BP1. <i>Nature Communications</i> , 2022, 13, 226.	5.8	11
77	TGS1 mediates 2,2,7-trimethyl guanosine capping of the human telomerase RNA to direct telomerase dependent telomere maintenance. <i>Nature Communications</i> , 2022, 13, 2302.	5.8	11
78	Target-enrichment sequencing for detailed characterization of small RNAs. <i>Nature Protocols</i> , 2018, 13, 768-786.	5.5	9
79	DROSHA is recruited to DNA damage sites by the MRN complex to promote non-homologous end joining. <i>Journal of Cell Science</i> , 2021, 134, .	1.2	9
80	DNA Damage In Situ Ligation Followed by Proximity Ligation Assay (DI-PLA). <i>Methods in Molecular Biology</i> , 2019, 1896, 11-20.	0.4	8
81	Long non-coding RNA in the control of genome stability and cancer phenotypes. <i>Non-coding RNA Investigation</i> , 2018, 2, 13-13.	0.6	7
82	RNase A treatment and reconstitution with DNA damage response RNA in living cells as a tool to study the role of non-coding RNA in the formation of DNA damage response foci. <i>Nature Protocols</i> , 2019, 14, 1489-1508.	5.5	7
83	Cellular senescence and cellular longevity: Nearly 50 years on and still working on it. <i>Experimental Cell Research</i> , 2008, 314, 1907-1908.	1.2	6
84	NOTCH1 modulates activity of DNA-PKcs. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2018, 808, 20-27.	0.4	4
85	A Role for Human DNA Polymerase β in Alternative Lengthening of Telomeres. <i>International Journal of Molecular Sciences</i> , 2021, 22, 2365.	1.8	3
86	Human nuclear ARGONAUTE 2 interacts in vivo only with small RNAs and not with DNA. <i>Cell Cycle</i> , 2015, 14, 2001-2002.	1.3	2
87	Detection of Telomeric DNA:RNA Hybrids Using TeloDRIP-qPCR. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9774.	1.8	1
88	Genetics and development. <i>Current Opinion in Genetics and Development</i> , 2002, 12, 621-627.	1.5	0
89	Cell biology. <i>Current Opinion in Cell Biology</i> , 2002, 14, 661-670.	2.6	0
90	In Vitro Detection of Long Noncoding RNA Generated from DNA Double-Strand Breaks. <i>Methods in Molecular Biology</i> , 2019, 2004, 209-219.	0.4	0