

# Anja K Bielinsky

## List of Publications by Year in descending order

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58  
papers

2,095  
citations

236925

25  
h-index

243625

44  
g-index

112  
all docs

112  
docs citations

112  
times ranked

2199  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mcm10 Regulates the Stability and Chromatin Association of DNA Polymerase- $\delta$ . <i>Molecular Cell</i> , 2004, 16, 173-185.	9.7	190
2	Chromosomal ARS1 Has a Single Leading Strand Start Site. <i>Molecular Cell</i> , 1999, 3, 477-486.	9.7	106
3	RNF4 and PLK1 are required for replication fork collapse in ATR-deficient cells. <i>Genes and Development</i> , 2013, 27, 2259-2273.	5.9	98
4	Replication Initiation Point Mapping. <i>Methods</i> , 1997, 13, 271-280.	3.8	89
5	Rapid DNA replication origin licensing protects stem cell pluripotency. <i>ELife</i> , 2017, 6, .	6.0	79
6	Human Mcm10 Regulates the Catalytic Subunit of DNA Polymerase- $\delta$ and Prevents DNA Damage during Replication. <i>Molecular Biology of the Cell</i> , 2007, 18, 4085-4095.	2.1	78
7	Origin recognition complex binding to a metazoan replication origin. <i>Current Biology</i> , 2001, 11, 1427-1431.	3.9	71
8	Ubiquitinated-PCNA protects replication forks from DNA2-mediated degradation by regulating Okazaki fragment maturation and chromatin assembly. <i>Nature Communications</i> , 2020, 11, 2147.	12.8	71
9	Interaction between PCNA and Diubiquitinated Mcm10 Is Essential for Cell Growth in Budding Yeast. <i>Molecular and Cellular Biology</i> , 2006, 26, 4806-4817.	2.3	69
10	Mechanisms of DNA Damage Tolerance: Post-Translational Regulation of PCNA. <i>Genes</i> , 2019, 10, 10.	2.4	69
11	Genome-wide replication profiles of S-phase checkpoint mutants reveal fragile sites in yeast. <i>EMBO Journal</i> , 2006, 25, 3627-3639.	7.8	68
12	Mcm10: A Dynamic Scaffold at Eukaryotic Replication Forks. <i>Genes</i> , 2017, 8, 73.	2.4	67
13	Mg <sup>2+</sup> and Ca <sup>2+</sup> Differentially Regulate $\beta$ 1 Integrin-Mediated Adhesion of Dermal Fibroblasts and Keratinocytes to Various Extracellular Matrix Proteins. <i>Experimental Cell Research</i> , 1994, 214, 381-388.	2.6	64
14	Enigmatic roles of Mcm10 in DNA replication. <i>Trends in Biochemical Sciences</i> , 2013, 38, 184-194.	7.5	64
15	Defects in DNA ligase I trigger PCNA ubiquitylation at Lys 107. <i>Nature Cell Biology</i> , 2010, 12, 74-79.	10.3	63
16	Antigen presentation function of brain-derived dendriform cells depends on astrocyte help. <i>International Immunology</i> , 1999, 11, 1265-1274.	4.0	57
17	DNA replication and chromatin. <i>Current Opinion in Genetics and Development</i> , 2002, 12, 243-248.	3.3	56
18	Structural Basis for DNA Binding by Replication Initiator Mcm10. <i>Structure</i> , 2008, 16, 1892-1901.	3.3	53

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19	A Conserved Hsp10-like Domain in Mcm10 Is Required to Stabilize the Catalytic Subunit of DNA Polymerase- $\delta$ in Budding Yeast. <i>Journal of Biological Chemistry</i> , 2006, 281, 18414-18425.	3.4	49
20	MCM10: One tool for all—Integrity, maintenance and damage control. <i>Seminars in Cell and Developmental Biology</i> , 2014, 30, 121-130.	5.0	45
21	Human NK cell deficiency as a result of biallelic mutations in MCM10. <i>Journal of Clinical Investigation</i> , 2020, 130, 5272-5286.	8.2	44
22	BRCA2 associates with MCM10 to suppress PRIMPOL-mediated repriming and single-stranded gap formation after DNA damage. <i>Nature Communications</i> , 2021, 12, 5966.	12.8	39
23	Flap endonuclease overexpression drives genome instability and DNA damage hypersensitivity in a PCNA-dependent manner. <i>Nucleic Acids Research</i> , 2018, 46, 5634-5650.	14.5	35
24	Easy detection of chromatin binding proteins by the histone association assay. <i>Biological Procedures Online</i> , 2005, 7, 60-69.	2.9	28
25	Mcm10 deficiency causes defective-replisome-induced mutagenesis and a dependency on error-free postreplicative repair. <i>Cell Cycle</i> , 2014, 13, 1737-1748.	2.6	26
26	Slx5/Slx8 Promotes Replication Stress Tolerance by Facilitating Mitotic Progression. <i>Cell Reports</i> , 2016, 15, 1254-1265.	6.4	26
27	EXO1 resection at G-quadruplex structures facilitates resolution and replication. <i>Nucleic Acids Research</i> , 2020, 48, 4960-4975.	14.5	26
28	SUMO-Targeted Ubiquitin Ligases and Their Functions in Maintaining Genome Stability. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5391.	4.1	25
29	Genetic Interactions Implicating Postreplicative Repair in Okazaki Fragment Processing. <i>PLoS Genetics</i> , 2015, 11, e1005659.	3.5	24
30	Replication Origins: Why Do We Need So Many?. <i>Cell Cycle</i> , 2003, 2, 306-308.	2.6	23
31	Ubc4 and Not4 Regulate Steady-State Levels of DNA Polymerase- $\delta$ to Promote Efficient and Accurate DNA Replication. <i>Molecular Biology of the Cell</i> , 2010, 21, 3205-3219.	2.1	23
32	Congenital Diseases of DNA Replication: Clinical Phenotypes and Molecular Mechanisms. <i>International Journal of Molecular Sciences</i> , 2021, 22, 911.	4.1	23
33	Bi-allelic MCM10 variants associated with immune dysfunction and cardiomyopathy cause telomere shortening. <i>Nature Communications</i> , 2021, 12, 1626.	12.8	22
34	Unligated Okazaki Fragments Induce PCNA Ubiquitination and a Requirement for Rad59-Dependent Replication Fork Progression. <i>PLoS ONE</i> , 2013, 8, e66379.	2.5	21
35	The N-terminus of Mcm10 is important for interaction with the 9-1-1 clamp and in resistance to DNA damage. <i>Nucleic Acids Research</i> , 2014, 42, 8389-8404.	14.5	21
36	Functional cross talk between the Fanconi anemia and ATRX/DAXX histone chaperone pathways promotes replication fork recovery. <i>Human Molecular Genetics</i> , 2020, 29, 1083-1095.	2.9	21

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37	<i>SLFN11</i> promotes stalled fork degradation that underlies the phenotype in Fanconi anemia cells. <i>Blood</i> , 2021, 137, 336-348.	1.4	17
38	Divalent cations (Mg <sup>2+</sup> , Ca <sup>2+</sup> ) differentially influence the beta1 integrin-mediated migration of human fibroblasts and keratinocytes to different extracellular matrix proteins. <i>Experimental Dermatology</i> , 1995, 4, 130-137.	2.9	16
39	The spatial arrangement of ORC binding modules determines the functionality of replication origins in budding yeast. <i>Nucleic Acids Research</i> , 2006, 34, 5069-5080.	14.5	16
40	Mcm10 Self-Association Is Mediated by an N-Terminal Coiled-Coil Domain. <i>PLoS ONE</i> , 2013, 8, e70518.	2.5	16
41	The anti-parasitic agent suramin and several of its analogues are inhibitors of the DNA binding protein Mcm10. <i>Open Biology</i> , 2019, 9, 190117.	3.6	15
42	Replication origins: why do we need so many?. <i>Cell Cycle</i> , 2003, 2, 307-9.	2.6	15
43	Damage-specific modification of PCNA. <i>Cell Cycle</i> , 2010, 9, 3698-3703.	2.6	12
44	eIF4E Threshold Levels Differ in Governing Normal and Neoplastic Expansion of Mammary Stem and Luminal Progenitor Cells. <i>Cancer Research</i> , 2015, 75, 687-697.	0.9	12
45	Mcm10: The glue at replication forks. <i>Cell Cycle</i> , 2016, 15, 3024-3025.	2.6	12
46	Replication Initiation Point Mapping: Approach and Implications. <i>Methods in Molecular Biology</i> , 2009, 521, 105-120.	0.9	4
47	Termination at sTop2. <i>Molecular Cell</i> , 2010, 39, 487-489.	9.7	3
48	Mapping ubiquitination sites of <i>S. cerevisiae</i> Mcm10. <i>Biochemistry and Biophysics Reports</i> , 2016, 8, 212-218.	1.3	3
49	Scarce but scary. <i>Nature Genetics</i> , 2007, 39, 707-708.	21.4	2
50	HDM2 ERKs PCNA. <i>Journal of Cell Biology</i> , 2010, 190, 487-489.	5.2	2
51	Crystal Structure of <i>Entamoeba histolytica</i> Cdc45 Suggests a Conformational Switch that May Regulate DNA Replication. <i>IScience</i> , 2018, 3, 102-109.	4.1	2
52	Encircled: Large-Scale Purification of Replication Origins from Mammalian Chromosomes. <i>Molecular Cell</i> , 2006, 21, 735-736.	9.7	0
53	Penetrating enemy territory: Soluble PCNA-peptides stress out MYCN-overexpressing neuroblastomas. <i>EBioMedicine</i> , 2015, 2, 1844-1845.	6.1	0
54	Not just for coding: a new role for histone tails in replication enzyme activation. <i>FEBS Journal</i> , 2016, 283, 4244-4246.	4.7	0

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55	12621 Targeted Chemical-Genetic Screen Platform for Identifying Drug Modes-of-Action. Journal of Clinical and Translational Science, 2021, 5, 101-102.	0.6	0
56	Multiple roles for Mcm10 during lagging strand synthesis. FASEB Journal, 2008, 22, 111.2.	0.5	0
57	Analyzing Origin Activation Patterns by Copy Number Change Experiments. Methods in Molecular Biology, 2009, 521, 279-294.	0.9	0
58	Defects in DNA Ligase I Trigger PCNA Ubiquitination at Lysine 107. FASEB Journal, 2010, 24, 492.5.	0.5	0