

# Maria L Kireeva

## List of Publications by Year in descending order

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57  
papers

4,633  
citations

147801

31  
h-index

149698

56  
g-index

58  
all docs

58  
docs citations

58  
times ranked

3721  
citing authors

#	ARTICLE	IF	CITATIONS
1	BNT162b2, mRNA-1273, and Sputnik V Vaccines Induce Comparable Immune Responses on a Par With Severe Course of COVID-19. <i>Frontiers in Immunology</i> , 2022, 13, 797918.	4.8	1
2	Dual-Antigen System Allows Elimination of False Positive Results in COVID-19 Serological Testing. <i>Diagnostics</i> , 2021, 11, 102.	2.6	8
3	Expression of SARS-CoV-2 surface glycoprotein fragment 319â€“640 in <i>E. coli</i> , and its refolding and purification. <i>Protein Expression and Purification</i> , 2021, 183, 105861.	1.3	25
4	IgG Antibodies Develop to Spike but Not to the Nucleocapsid Viral Protein in Many Asymptomatic and Light COVID-19 Cases. <i>Viruses</i> , 2021, 13, 1945.	3.3	16
5	NusG controls transcription pausing and RNA polymerase translocation throughout the <i>Bacillus subtilis</i> genome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 21628-21636.	7.1	38
6	The Role of Pyrophosphorolysis in the Initiation-to-Elongation Transition by <i>E. coli</i> RNA Polymerase. <i>Journal of Molecular Biology</i> , 2019, 431, 2528-2542.	4.2	7
7	RNAâ€“DNA and DNAâ€“DNA base-pairing at the upstream edge of the transcription bubble regulate translocation of RNA polymerase and transcription rate. <i>Nucleic Acids Research</i> , 2018, 46, 5764-5775.	14.5	12
8	RNA Polymerases and Transcription. , 2018, , 1-9.		0
9	Production and characterization of a highly pure RNA polymerase holoenzyme from <i>Mycobacterium tuberculosis</i> . <i>Protein Expression and Purification</i> , 2017, 134, 1-10.	1.3	7
10	A <i>Cre</i> Transcription Fidelity Reporter Identifies GreA as a Major RNA Proofreading Factor in <i>Escherichia coli</i> . <i>Genetics</i> , 2017, 206, 179-187.	2.9	26
11	Cotranscriptional Production of Chemically Modified RNA Nanoparticles. <i>Methods in Molecular Biology</i> , 2017, 1632, 91-105.	0.9	4
12	Productive mRNA stem loop-mediated transcriptional slippage: Crucial features in common with intrinsic terminators. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E1984-93.	7.1	20
13	Computational and Experimental Studies of Reassociating RNA/DNA Hybrids Containing Split Functionalities. <i>Methods in Enzymology</i> , 2015, 553, 313-334.	1.0	12
14	Triggering of RNA Interference with RNAâ€“RNA, RNAâ€“DNA, and DNAâ€“RNA Nanoparticles. <i>ACS Nano</i> , 2015, 9, 251-259.	14.6	100
15	Direct Competition Assay for Transcription Fidelity. <i>Methods in Molecular Biology</i> , 2015, 1276, 153-164.	0.9	2
16	Co-transcriptional production of RNAâ€“DNA hybrids for simultaneous release of multiple split functionalities. <i>Nucleic Acids Research</i> , 2014, 42, 2085-2097.	14.5	54
17	A Genetic Assay for Transcription Errors Reveals Multilayer Control of RNA Polymerase II Fidelity. <i>PLoS Genetics</i> , 2014, 10, e1004532.	3.5	26
18	Coliphage HK022 Nun protein inhibits RNA polymerase translocation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E2368-75.	7.1	28

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19	<i>In Silico</i> Design and Enzymatic Synthesis of Functional RNA Nanoparticles. <i>Accounts of Chemical Research</i> , 2014, 47, 1731-1741.	15.6	80
20	Intrinsic Translocation Barrier as an Initial Step in Pausing by RNA Polymerase II. <i>Journal of Molecular Biology</i> , 2013, 425, 697-712.	4.2	38
21	RNA Polymerase Structure, Function, Regulation, Dynamics, Fidelity, and Roles in GENE EXPRESSION. <i>Chemical Reviews</i> , 2013, 113, 8325-8330.	47.7	8
22	The RNA polymerase bridge helix YFI motif in catalysis, fidelity and translocation. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2013, 1829, 187-198.	1.9	31
23	The Fidelity of Transcription. <i>Journal of Biological Chemistry</i> , 2013, 288, 2689-2699.	3.4	30
24	Complete dissection of transcription elongation reveals slow translocation of RNA polymerase II in a linear ratchet mechanism. <i>ELife</i> , 2013, 2, e00971.	6.0	111
25	Sensitivity of Mitochondrial Transcription and Resistance of RNA Polymerase II Dependent Nuclear Transcription to Antiviral Ribonucleosides. <i>PLoS Pathogens</i> , 2012, 8, e1003030.	4.7	119
26	Co-transcriptional Assembly of Chemically Modified RNA Nanoparticles Functionalized with siRNAs. <i>Nano Letters</i> , 2012, 12, 5192-5195.	9.1	117
27	Mechanism of Translesion Transcription by RNA Polymerase II and Its Role in Cellular Resistance to DNA Damage. <i>Molecular Cell</i> , 2012, 46, 18-29.	9.7	104
28	Molecular dynamics and mutational analysis of the catalytic and translocation cycle of RNA polymerase. <i>BMC Biophysics</i> , 2012, 5, 11.	4.4	35
29	Interaction of RNA Polymerase II Fork Loop 2 with Downstream Non-template DNA Regulates Transcription Elongation. <i>Journal of Biological Chemistry</i> , 2011, 286, 30898-30910.	3.4	25
30	Synergistic action of RNA polymerases in overcoming the nucleosomal barrier. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 745-752.	8.2	114
31	Translocation by multi-subunit RNA polymerases. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2010, 1799, 389-401.	1.9	43
32	Conformational coupling, bridge helix dynamics and active site dehydration in catalysis by RNA polymerase. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2010, 1799, 575-587.	1.9	34
33	Site-directed mutagenesis, purification and assay of <i>Saccharomyces cerevisiae</i> RNA polymerase II. <i>Protein Expression and Purification</i> , 2010, 69, 83-90.	1.3	17
34	Mechanism of sequence-specific pausing of bacterial RNA polymerase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 8900-8905.	7.1	100
35	Rpb9 Subunit Controls Transcription Fidelity by Delaying NTP Sequestration in RNA Polymerase II. <i>Journal of Biological Chemistry</i> , 2009, 284, 19601-19612.	3.4	74
36	Millisecond phase kinetic analysis of elongation catalyzed by human, yeast, and <i>Escherichia coli</i> RNA polymerase. <i>Methods</i> , 2009, 48, 333-345.	3.8	27

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37	Transient Reversal of RNA Polymerase II Active Site Closing Controls Fidelity of Transcription Elongation. <i>Molecular Cell</i> , 2008, 30, 557-566.	9.7	154
38	Effects of Friedreich's ataxia (GAA) <sub>n</sub> (TTC) <sub>n</sub> repeats on RNA synthesis and stability. <i>Nucleic Acids Research</i> , 2007, 35, 1075-1084.	14.5	49
39	Mutations in the <i>Saccharomyces cerevisiae</i> RPB1 Gene Conferring Hypersensitivity to 6-Azauracil. <i>Genetics</i> , 2006, 172, 2201-2209.	2.9	96
40	Nature of the Nucleosomal Barrier to RNA Polymerase II. <i>Molecular Cell</i> , 2005, 18, 97-108.	9.7	198
41	Chromatin remodeling by RNA polymerases. <i>Trends in Biochemical Sciences</i> , 2004, 29, 127-135.	7.5	88
42	Assay of the Fate of the Nucleosome During Transcription by RNA Polymerase II. <i>Methods in Enzymology</i> , 2003, 371, 564-577.	1.0	15
43	Assays and Affinity Purification of Biotinylated and Nonbiotinylated Forms of Double-Tagged Core RNA Polymerase II from <i>Saccharomyces cerevisiae</i> . <i>Methods in Enzymology</i> , 2003, 370, 138-155.	1.0	45
44	Engineering of Elongation Complexes of Bacterial and Yeast RNA Polymerases. <i>Methods in Enzymology</i> , 2003, 371, 233-251.	1.0	78
45	Bacterial Polymerase and Yeast Polymerase II Use Similar Mechanisms for Transcription through Nucleosomes. <i>Journal of Biological Chemistry</i> , 2003, 278, 36148-36156.	3.4	51
46	Nucleosome Remodeling Induced by RNA Polymerase II. <i>Molecular Cell</i> , 2002, 9, 541-552.	9.7	419
47	Shortening of RNA:DNA Hybrid in the Elongation Complex of RNA Polymerase Is a Prerequisite for Transcription Termination. <i>Molecular Cell</i> , 2002, 10, 1151-1162.	9.7	130
48	The 8-Nucleotide-long RNA:DNA Hybrid Is a Primary Stability Determinant of the RNA Polymerase II Elongation Complex. <i>Journal of Biological Chemistry</i> , 2000, 275, 6530-6536.	3.4	200
49	Overextended RNA:DNA hybrid as a negative regulator of RNA polymerase II processivity 1 Edited by R. Ebright. <i>Journal of Molecular Biology</i> , 2000, 299, 325-335.	4.2	52
50	Activation-dependent Adhesion of Human Platelets to Cyr61 and Fisp12/Mouse Connective Tissue Growth Factor Is Mediated through Integrin $\alpha 5 \beta 1$ . <i>Journal of Biological Chemistry</i> , 1999, 274, 24321-24327.	3.4	196
51	Adhesion of Human Umbilical Vein Endothelial Cells to the Immediate-Early Gene Product Cyr61 Is Mediated through Integrin $\alpha 5 \beta 1$ . <i>Journal of Biological Chemistry</i> , 1998, 273, 3090-3096.	3.4	192
52	CYR61, a product of a growth factor-inducible immediate early gene, promotes angiogenesis and tumor growth. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 6355-6360.	7.1	432
53	Cyr61, Product of a Growth Factor-Inducible Immediate-Early Gene, Regulates Chondrogenesis in Mouse Limb Bud Mesenchymal Cells. <i>Developmental Biology</i> , 1997, 192, 492-508.	2.0	140
54	Cyr61 and Fisp12 Are Both ECM-Associated Signaling Molecules: Activities, Metabolism, and Localization during Development. <i>Experimental Cell Research</i> , 1997, 233, 63-77.	2.6	243

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55	Cyr61, a Product of a Growth Factor-Inducible Immediate-Early Gene, Promotes Cell Proliferation, Migration, and Adhesion. <i>Molecular and Cellular Biology</i> , 1996, 16, 1326-1334.	2.3	309
56	Circularly permuted dihydrofolate reductase of E.coli has functional activity and a destabilized tertiary structure. <i>Protein Engineering, Design and Selection</i> , 1994, 7, 1373-1377.	2.1	38
57	Novel data on interactions of elongation factor Ts. <i>Biochimie</i> , 1992, 74, 419-425.	2.6	15