

Fei-Long Meng

List of Publications by Year in descending order

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Version: 2024-02-01

49
papers

2,567
citations

304743

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243625

44
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53
all docs

53
docs citations

53
times ranked

3968
citing authors

#	ARTICLE	IF	CITATIONS
1	Mechanisms of Programmed DNA Lesions and Genomic Instability in the Immune System. <i>Cell</i> , 2013, 152, 417-429.	28.9	407
2	The RNA Exosome Targets the AID Cytidine Deaminase to Both Strands of Transcribed Duplex DNA Substrates. <i>Cell</i> , 2011, 144, 353-363.	28.9	275
3	Convergent Transcription at Intragenic Super-Enhancers Targets AID-Initiated Genomic Instability. <i>Cell</i> , 2014, 159, 1538-1548.	28.9	221
4	Chromosomal Loop Domains Direct the Recombination of Antigen Receptor Genes. <i>Cell</i> , 2015, 163, 947-959.	28.9	140
5	Sequence-Intrinsic Mechanisms that Target AID Mutational Outcomes on Antibody Genes. <i>Cell</i> , 2015, 163, 1124-1137.	28.9	136
6	AID Recognizes Structured DNA for Class Switch Recombination. <i>Molecular Cell</i> , 2017, 67, 361-373.e4.	9.7	136
7	Tid-CRISPR Allows for Efficient and Precise Gene Knockin in Mouse and Human Cells. <i>Developmental Cell</i> , 2018, 45, 526-536.e5.	7.0	123
8	Transcriptional landscape of the human cell cycle. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 3473-3478.	7.1	110
9	Phosphatidylinositol 3-kinase $\hat{\imath}$ blockade increases genomic instability in B cells. <i>Nature</i> , 2017, 542, 489-493.	27.8	105
10	Pooled CRISPR screening identifies m ⁶ A as a positive regulator of macrophage activation. <i>Science Advances</i> , 2021, 7, .	10.3	102
11	Orientation-specific joining of AID-initiated DNA breaks promotes antibody class switching. <i>Nature</i> , 2015, 525, 134-139.	27.8	93
12	The mTOR $\hat{\imath}$ S6K pathway links growth signalling to DNA damage response by targeting RNF168. <i>Nature Cell Biology</i> , 2018, 20, 320-331.	10.3	86
13	<i>Saccharomyces cerevisiae</i> Est3p dimerizes in vitro and dimerization contributes to efficient telomere replication in vivo. <i>Nucleic Acids Research</i> , 2006, 34, 407-416.	14.5	65
14	Telomerase-Null Survivor Screening Identifies Novel Telomere Recombination Regulators. <i>PLoS Genetics</i> , 2013, 9, e1003208.	3.5	52
15	Global detection of DNA repair outcomes induced by CRISPR $\hat{\imath}$ Cas9. <i>Nucleic Acids Research</i> , 2021, 49, 8732-8742.	14.5	52
16	<i>Candida</i> $\hat{\imath}$ albicans $\hat{\imath}$, a distinctive fungal model for cellular aging study. <i>Aging Cell</i> , 2008, 7, 746-757.	6.7	42
17	ERCC6L2 promotes DNA orientation-specific recombination in mammalian cells. <i>Cell Research</i> , 2020, 30, 732-744.	12.0	41
18	Sua5p a single-stranded telomeric DNA-binding protein facilitates telomere replication. <i>EMBO Journal</i> , 2009, 28, 1466-1478.	7.8	34

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19	SWR1 Complex Poises Heterochromatin Boundaries for Antisilencing Activity Propagation. <i>Molecular and Cellular Biology</i> , 2010, 30, 2391-2400.	2.3	34
20	Genome-wide mutational signatures revealed distinct developmental paths for human B cell lymphomas. <i>Journal of Experimental Medicine</i> , 2021, 218, .	8.5	29
21	Intrinsic Nucleotide Preference of Diversifying Base Editors Guides Antibody Ex Vivo Affinity Maturation. <i>Cell Reports</i> , 2018, 25, 884-892.e3.	6.4	28
22	Cis- and trans-factors affecting AID targeting and mutagenic outcomes in antibody diversification. <i>Advances in Immunology</i> , 2019, 141, 51-103.	2.2	26
23	AMPK-mediated phosphorylation on 53BP1 promotes c-NHEJ. <i>Cell Reports</i> , 2021, 34, 108713.	6.4	23
24	Telomere Recombination Accelerates Cellular Aging in <i>Saccharomyces cerevisiae</i> . <i>PLoS Genetics</i> , 2009, 5, e1000535.	3.5	17
25	A systematic dissection of the epigenomic heterogeneity of lung adenocarcinoma reveals two different subclasses with distinct prognosis and core regulatory networks. <i>Genome Biology</i> , 2021, 22, 156.	8.8	17
26	Uncovering a conserved vulnerability site in SARS-CoV-2 by a human antibody. <i>EMBO Molecular Medicine</i> , 2021, 13, e14544.	6.9	17
27	Genome integrity and neurogenesis of postnatal hippocampal neural stem/progenitor cells require a unique regulator Filia. <i>Science Advances</i> , 2020, 6, .	10.3	14
28	Sua5p is required for telomere recombination in <i>Saccharomyces cerevisiae</i> . <i>Cell Research</i> , 2010, 20, 495-498.	12.0	13
29	The 3'-flap endonuclease XPF-ERCC1 promotes alternative end joining and chromosomal translocation during B cell class switching. <i>Cell Reports</i> , 2021, 36, 109756.	6.4	13
30	The development of neutralizing antibodies against SARS-CoV-2 and their common features. <i>Journal of Molecular Cell Biology</i> , 2021, 12, 980-986.	3.3	13
31	Ig Enhancers Increase RNA Polymerase II Stalling at Somatic Hypermutation Target Sequences. <i>Journal of Immunology</i> , 2022, 208, 143-154.	0.8	13
32	Parp3 promotes long-range end joining in murine cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 10076-10081.	7.1	11
33	Generation of Genomic Alteration from Cytidine Deamination. <i>Advances in Experimental Medicine and Biology</i> , 2018, 1044, 49-64.	1.6	11
34	REV7 is required for processing AID initiated DNA lesions in activated B cells. <i>Nature Communications</i> , 2020, 11, 2812.	12.8	9
35	Targeting HSPA1A in ARID2-deficient lung adenocarcinoma. <i>National Science Review</i> , 2021, 8, nwab014.	9.5	9
36	Telomere Recombination Preferentially Occurs at Short Telomeres in Telomerase-Null Type II Survivors. <i>PLoS ONE</i> , 2014, 9, e90644.	2.5	8

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37	Swc4 positively regulates telomere length independently of its roles in NuA4 and SWR1 complexes. <i>Nucleic Acids Research</i> , 2020, 48, 12792-12803.	14.5	8
38	UdgX-Mediated Uracil Sequencing at Single-Nucleotide Resolution. <i>Journal of the American Chemical Society</i> , 2022, 144, 1323-1331.	13.7	8
39	B cell receptor signatures associated with strong and poor SARS-CoV-2 vaccine responses. <i>Emerging Microbes and Infections</i> , 2022, 11, 452-464.	6.5	8
40	C-terminal deletion-induced condensation sequesters AID from IgH targets in immunodeficiency. <i>EMBO Journal</i> , 2022, 41, e109324.	7.8	5
41	A Rapid Embryonic Stem Cell-Based Mouse Model for B-cell Lymphomas Driven by Epstein-Barr Virus Protein LMP1. <i>Cancer Immunology Research</i> , 2015, 3, 641-649.	3.4	3
42	Repair of programmed DNA lesions in antibody class switch recombination: common and unique features. <i>Genome Instability & Disease</i> , 2021, 2, 115-125.	1.1	3
43	New Chromatin Run-On Reaction Enables Global Mapping of Active RNA Polymerase Locations in an Enrichment-free Manner. <i>ACS Chemical Biology</i> , 2022, 17, 768-775.	3.4	3
44	PI3Kdelta Inhibitors Increase Genomic Instability By Upregulating AID Expression. <i>Blood</i> , 2015, 126, 164-164.	1.4	1
45	Evaluation of a Novel Missense Activation-Induced Deaminase AID Mutation in a Child with Hyper IgM Syndrome: Is it a Pathogenic Mutation?. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 133, AB70.	2.9	0
46	The Mechanism of IgH Class Switch Recombination. , 2015, , 345-362.		0
47	Abstract A174: Mechanistic elucidation of activation-induced deaminase (AID) in immunity and cancer. , 2016, , .		0
48	Abstract A180: Topologically associated domains genome-wide restrict the off-target activity of recombination activating gene 1/2 endonuclease. , 2016, , .		0
49	AMPK-Mediated Phosphorylation on 53BP1 Promotes NHEJ. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0