

Andrew Flaus

List of Publications by Year in descending order

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39
papers

4,090
citations

236925

25
h-index

330143

37
g-index

42
all docs

42
docs citations

42
times ranked

3847
citing authors

#	ARTICLE	IF	CITATIONS
1	Histone isoforms and the oncohistone code. <i>Current Opinion in Genetics and Development</i> , 2021, 67, 61-66.	3.3	15
2	Survival outcomes are associated with genomic instability in luminal breast cancers. <i>PLoS ONE</i> , 2021, 16, e0245042.	2.5	8
3	Cryo-EM structure of the nucleosome core particle containing <i>Giardia lamblia</i> histones. <i>Nucleic Acids Research</i> , 2021, 49, 8934-8946.	14.5	20
4	The Face of Chromatin Variants. <i>Cell</i> , 2019, 178, 1284-1286.	28.9	0
5	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.	10.3	233
6	Viral proteins as a potential driver of histone depletion in dinoflagellates. <i>Nature Communications</i> , 2018, 9, 1535.	12.8	33
7	Unlocking the nucleosome. <i>Science</i> , 2017, 355, 245-246.	12.6	4
8	Histone H2AX Y142 phosphorylation is a low abundance modification. <i>International Journal of Mass Spectrometry</i> , 2015, 391, 139-145.	1.5	12
9	Life at the mesoscale: the self-organised cytoplasm and nucleoplasm. <i>BMC Biophysics</i> , 2015, 8, 4.	4.4	16
10	A chromatin-independent role of Polycomb-like 1 to stabilize p53 and promote cellular quiescence. <i>Genes and Development</i> , 2015, 29, 2231-2243.	5.9	32
11	Principles and practice of nucleosome positioning <i>in vitro</i> . <i>Frontiers in Life Science: Frontiers of Interdisciplinary Research in the Life Sciences</i> , 2011, 5, 5-27.	1.1	16
12	Mechanisms for ATP-dependent chromatin remodelling: the means to the end. <i>FEBS Journal</i> , 2011, 278, 3579-3595.	4.7	102
13	Structure and Function of Histone H2AX. <i>Sub-Cellular Biochemistry</i> , 2010, 50, 55-78.	2.4	47
14	Nucleosomes can invade DNA territories occupied by their neighbors. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 151-158.	8.2	95
15	Histone Tails and the H3 \pm N Helix Regulate Nucleosome Mobility and Stability. <i>Molecular and Cellular Biology</i> , 2007, 27, 4037-4048.	2.3	122
16	Histone Modifications Influence the Action of Snf2 Family Remodelling Enzymes by Different Mechanisms. <i>Journal of Molecular Biology</i> , 2007, 374, 563-579.	4.2	121
17	Chromatin modulation and the DNA damage response. <i>Experimental Cell Research</i> , 2006, 312, 2677-2686.	2.6	33
18	Identification of multiple distinct Snf2 subfamilies with conserved structural motifs. <i>Nucleic Acids Research</i> , 2006, 34, 2887-2905.	14.5	612

#	ARTICLE	IF	CITATIONS
19	Snf2 family ATPases and DExx box helicases: differences and unifying concepts from high-resolution crystal structures. <i>Nucleic Acids Research</i> , 2006, 34, 4160-4167.	14.5	93
20	Analysis of Nucleosome Repositioning by Yeast ISWI and Chd1 Chromatin Remodeling Complexes*. <i>Journal of Biological Chemistry</i> , 2006, 281, 16279-16288.	3.4	167
21	Nucleosome dynamics. <i>Biochemical Society Symposia</i> , 2006, 73, 109-119.	2.7	2
22	Sin mutations alter inherent nucleosome mobility. <i>EMBO Journal</i> , 2004, 23, 343-353.	7.8	114
23	Mechanisms for ATP-dependent chromatin remodelling: farewell to the tuna-can octamer?. <i>Current Opinion in Genetics and Development</i> , 2004, 14, 165-173.	3.3	138
24	Mechanisms for nucleosome mobilization. <i>Biopolymers</i> , 2003, 68, 563-578.	2.4	74
25	Site-Specific Attachment of Reporter Compounds to Recombinant Histones. <i>Methods in Enzymology</i> , 2003, 375, 211-228.	1.0	10
26	Histone H2A/H2B Dimer Exchange by ATP-Dependent Chromatin Remodeling Activities. <i>Molecular Cell</i> , 2003, 12, 1599-1606.	9.7	175
27	Dynamic Properties of Nucleosomes during Thermal and ATP-Driven Mobilization. <i>Molecular and Cellular Biology</i> , 2003, 23, 7767-7779.	2.3	94
28	Evidence for DNA Translocation by the ISWI Chromatin-Remodeling Enzyme. <i>Molecular and Cellular Biology</i> , 2003, 23, 1935-1945.	2.3	131
29	Mechanisms for ATP-dependent chromatin remodelling. <i>Current Opinion in Genetics and Development</i> , 2001, 11, 148-154.	3.3	157
30	Generation of Superhelical Torsion by ATP-Dependent Chromatin Remodeling Activities. <i>Cell</i> , 2000, 103, 1133-1142.	28.9	241
31	Base-Pair Resolution Mapping of Nucleosomes In Vitro. , 1999, 119, 45-60.		7
32	Base-pair resolution mapping of nucleosome positions using site-directed hydroxy radicals. <i>Methods in Enzymology</i> , 1999, 304, 251-263.	1.0	12
33	Nucleosome mobilization catalysed by the yeast SWI/SNF complex. <i>Nature</i> , 1999, 400, 784-787.	27.8	306
34	Positioning and stability of nucleosomes on MMTV 3' LTR sequences. <i>Journal of Molecular Biology</i> , 1998, 275, 427-441.	4.2	120
35	The mouse mammary tumour virus promoter positioned on a tetramer of histones H3 and H4 binds nuclear factor 1 and OTF1. <i>Journal of Molecular Biology</i> , 1998, 278, 725-739.	4.2	54
36	Differential nucleosome positioning on <i>Xenopus</i> oocyte and somatic 5 s RNA genes determines both TFIIIA and H1 binding: a mechanism for selective H1 repression 1 Edited by J. Karn. <i>Journal of Molecular Biology</i> , 1998, 282, 683-697.	4.2	58

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37	Characterization of nucleosome core particles containing histone proteins made in bacteria 1 1Edited by A. Klug. Journal of Molecular Biology, 1997, 272, 301-311.	4.2	446
38	Mapping nucleosome position at single base-pair resolution by using site-directed hydroxyl radicals.. Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 1370-1375.	7.1	157
39	Purification and Crystallization of the Endoglycosidase PNGase F, a Peptide:N-glycosidase from Flavobacterium meningosepticum. Journal of Molecular Biology, 1994, 241, 624-626.	4.2	6