

# Alejandro Vaquero

## List of Publications by Year in descending order

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Version: 2024-02-01

60  
papers

6,021  
citations

117625

34  
h-index

161849

54  
g-index

65  
all docs

65  
docs citations

65  
times ranked

8516  
citing authors

#	ARTICLE	IF	CITATIONS
1	Human SirT1 Interacts with Histone H1 and Promotes Formation of Facultative Heterochromatin. <i>Molecular Cell</i> , 2004, 16, 93-105.	9.7	796
2	SirT2 is a histone deacetylase with preference for histone H4 Lys 16 during mitosis. <i>Genes and Development</i> , 2006, 20, 1256-1261.	5.9	535
3	SirT3 is a nuclear NAD <sup>+</sup> -dependent histone deacetylase that translocates to the mitochondria upon cellular stress. <i>Genes and Development</i> , 2007, 21, 920-928.	5.9	409
4	SIRT1 regulates the histone methyl-transferase SUV39H1 during heterochromatin formation. <i>Nature</i> , 2007, 450, 440-444.	27.8	380
5	Composition and histone substrates of polycomb repressive group complexes change during cellular differentiation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 1859-1864.	7.1	371
6	L3MBTL1, a Histone-Methylation-Dependent Chromatin Lock. <i>Cell</i> , 2007, 129, 915-928.	28.9	318
7	The Dual Role of Sirtuins in Cancer. <i>Genes and Cancer</i> , 2011, 2, 648-662.	1.9	281
8	NAD <sup>+</sup> -dependent deacetylation of H4 lysine 16 by class III HDACs. <i>Oncogene</i> , 2007, 26, 5505-5520.	5.9	259
9	The tumor suppressor SirT2 regulates cell cycle progression and genome stability by modulating the mitotic deposition of H4K20 methylation. <i>Genes and Development</i> , 2013, 27, 639-653.	5.9	232
10	<sc>SIRT</sc> 7 promotes genome integrity and modulates non-homologous end joining <sc>DNA</sc> repair. <i>EMBO Journal</i> , 2016, 35, 1488-1503.	7.8	211
11	The Constantly Changing Face of Chromatin. <i>Science of Aging Knowledge Environment: SAGE KE</i> , 2003, 2003, 4re-4.	0.8	147
12	Calorie restriction and the exercise of chromatin. <i>Genes and Development</i> , 2009, 23, 1849-1869.	5.9	130
13	The Diversity of Histone Versus Nonhistone Sirtuin Substrates. <i>Genes and Cancer</i> , 2013, 4, 148-163.	1.9	119
14	Stabilization of Suv39H1 by SirT1 Is Part of Oxidative Stress Response and Ensures Genome Protection. <i>Molecular Cell</i> , 2011, 42, 210-223.	9.7	115
15	Sirtuin-dependent epigenetic regulation in the maintenance of genome integrity. <i>FEBS Journal</i> , 2015, 282, 1745-1767.	4.7	114
16	The conserved role of sirtuins in chromatin regulation. <i>International Journal of Developmental Biology</i> , 2009, 53, 303-322.	0.6	102
17	The microRNA-449 family inhibits TGF- $\beta$ -mediated liver cancer cell migration by targeting SOX4. <i>Journal of Hepatology</i> , 2017, 66, 1012-1021.	3.7	102
18	The N-terminal POZ Domain of GAGA Mediates the Formation of Oligomers That Bind DNA with High Affinity and Specificity. <i>Journal of Biological Chemistry</i> , 1999, 274, 16461-16469.	3.4	95

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19	The Embryonic Linker Histone H1 Variant of Drosophila, dBigH1, Regulates Zygotic Genome Activation. <i>Developmental Cell</i> , 2013, 26, 578-590.	7.0	91
20	Sirtuins in stress response: guardians of the genome. <i>Oncogene</i> , 2014, 33, 3764-3775.	5.9	91
21	Sirt7 promotes adipogenesis in the mouse by inhibiting autocatalytic activation of Sirt1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E8352-E8361.	7.1	88
22	KAT6B Is a Tumor Suppressor Histone H3 Lysine 23 Acetyltransferase Undergoing Genomic Loss in Small Cell Lung Cancer. <i>Cancer Research</i> , 2015, 75, 3936-3945.	0.9	65
23	Dietary Restriction: Standing Up for Sirtuins. <i>Science</i> , 2010, 329, 1012-1013.	12.6	63
24	Mammalian HP1 Isoforms Have Specific Roles in Heterochromatin Structure and Organization. <i>Cell Reports</i> , 2017, 21, 2048-2057.	6.4	63
25	Characterization of Drosophila melanogaster JmjC+N histone demethylases. <i>Nucleic Acids Research</i> , 2008, 36, 2852-2863.	14.5	58
26	SIRT7 mediates L1 elements transcriptional repression and their association with the nuclear lamina. <i>Nucleic Acids Research</i> , 2019, 47, 7870-7885.	14.5	55
27	Arachidonic and oleic acid exert distinct effects on the DNA methylome. <i>Epigenetics</i> , 2016, 11, 321-334.	2.7	52
28	Drosophila melanogaster linker histone dH1 is required for transposon silencing and to preserve genome integrity. <i>Nucleic Acids Research</i> , 2012, 40, 5402-5414.	14.5	51
29	Analysis of the Effects of Daunorubicin and WP631 on Transcription. <i>Current Medicinal Chemistry</i> , 2001, 8, 1-8.	2.4	46
30	SIRT6-dependent cysteine monoubiquitination in the PRE-SET domain of Suv39h1 regulates the NF- $\kappa$ B pathway. <i>Nature Communications</i> , 2018, 9, 101.	12.8	46
31	A View on the Role of Epigenetics in the Biology of Malaria Parasites. <i>PLoS Pathogens</i> , 2012, 8, e1002943.	4.7	43
32	The GAGA Factor of Drosophila Binds Triple-stranded DNA. <i>Journal of Biological Chemistry</i> , 1998, 273, 24640-24648.	3.4	41
33	SIRT1/2 orchestrate acquisition of DNA methylation and loss of histone H3 activating marks to prevent premature activation of inflammatory genes in macrophages. <i>Nucleic Acids Research</i> , 2020, 48, 665-681.	14.5	39
34	Combined bottom-up and top-down mass spectrometry analyses of the pattern of post-translational modifications of Drosophila melanogaster linker histone H1. <i>Journal of Proteomics</i> , 2012, 75, 4124-4138.	2.4	38
35	Loss of $\langle \text{sc} \rangle \text{SIRT} \langle / \text{sc} \rangle 2$ leads to axonal degeneration and locomotor disability associated with redox and energy imbalance. <i>Aging Cell</i> , 2017, 16, 1404-1413.	6.7	36
36	Sirt7 auto-ADP-ribosylation regulates glucose starvation response through mH2A1. <i>Science Advances</i> , 2020, 6, eaaz2590.	10.3	33

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37	Functional Mapping of the GAGA Factor Assigns Its Transcriptional Activity to the C-terminal Glutamine-rich Domain. <i>Journal of Biological Chemistry</i> , 2000, 275, 19461-19468.	3.4	32
38	SIRT7-dependent deacetylation of NPM promotes p53 stabilization following UV-induced genotoxic stress. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	30
39	SIRT1 activation with neuroheal is neuroprotective but SIRT2 inhibition with AK7 is detrimental for disconnected motoneurons. <i>Cell Death and Disease</i> , 2018, 9, 531.	6.3	26
40	Chromatin regulation by Histone H4 acetylation at Lysine 16 during cell death and differentiation in the myeloid compartment. <i>Nucleic Acids Research</i> , 2019, 47, 5016-5037.	14.5	23
41	An HP1 isoform-specific feedback mechanism regulates Suv39h1 activity under stress conditions. <i>Epigenetics</i> , 2017, 12, 166-175.	2.7	22
42	At the crossroad of lifespan, calorie restriction, chromatin and disease: Meeting on sirtuins. <i>Cell Cycle</i> , 2010, 9, 1907-1912.	2.6	20
43	Sirtuin 1 Inhibiting Thiocyanates (S1th)â€”A New Class of Isotype Selective Inhibitors of NAD+ Dependent Lysine Deacetylases. <i>Frontiers in Oncology</i> , 2020, 10, 657.	2.8	19
44	Complex role of SIRT6 in NF-Î²B pathway regulation. <i>Molecular and Cellular Oncology</i> , 2018, 5, e1445942.	0.7	16
45	A Big Step for SIRT7, One Giant Leap for Sirtuinsâ€” in Cancer. <i>Cancer Cell</i> , 2012, 21, 719-721.	16.8	15
46	Steps Toward Understanding the Inheritance of Repressive Methyl-Lysine Marks in Histones. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2004, 69, 171-182.	1.1	14
47	A Synthetic mRNA Cell Reprogramming Method Using <i>CYCLIN D1</i> Promotes DNA rEpair, Generating Improved Genetically Stable Human Induced Pluripotent Stem Cells. <i>Stem Cells</i> , 2021, 39, 866-881.	3.2	14
48	Sirtuins in female meiosis and in reproductive longevity. <i>Molecular Reproduction and Development</i> , 2020, 87, 1175-1187.	2.0	12
49	Shikimic acid protects skin cells from UV-induced senescence through activation of the NAD+-dependent deacetylase SIRT1. <i>Aging</i> , 2021, 13, 12308-12333.	3.1	11
50	Histone methyltransferase <i>Suv39h1</i> deficiency prevents <i>Myc</i> -induced chromosomal instability in murine myeloid leukemias. <i>Genes Chromosomes and Cancer</i> , 2013, 52, 423-430.	2.8	10
51	Activation-induced cytidine deaminase targets SUV4-20-mediated histone H4K20 trimethylation to class-switch recombination sites. <i>Scientific Reports</i> , 2017, 7, 7594.	3.3	10
52	Activation properties of GAGA transcription factor. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2008, 1779, 312-317.	1.9	9
53	Raising the list of Sir7 targets to a new level. <i>Proteomics</i> , 2017, 17, 1700137.	2.2	4
54	Niacin. , 2020, , 287-293.		2

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55	Sirtuins in hematopoiesis and blood malignancies. , 2021, , 373-391.		2
56	Sirtuins in Biology and Disease. , 2008, , 73-104.		2
57	Methods to Study the Role of Sirtuins in Genome Stability. Methods in Molecular Biology, 2013, 1077, 273-283.	0.9	2
58	The Histone Code and Disease. , 2016, , 417-445.		1
59	Sirtuins in Biology and Disease. , 2008, , 73-104.		1
60	Sirtuins as a Double-Edged Sword in Cancer: From Molecular Mechanisms to Therapeutic Opportunities. , 2015, , 75-106.		0