

Roberto Ferrari

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

1,989
citations

257357

24
h-index

360920

35
g-index

43
all docs

43
docs citations

43
times ranked

3682
citing authors

#	ARTICLE	IF	CITATIONS
1	Histone Acetylation Regulates Intracellular pH. <i>Molecular Cell</i> , 2013, 49, 310-321.	4.5	210
2	Epigenetic Reprogramming by Adenovirus e1a. <i>Science</i> , 2008, 321, 1086-1088.	6.0	207
3	Effects of Serelaxin in Patients with Acute Heart Failure. <i>New England Journal of Medicine</i> , 2019, 381, 716-726.	13.9	174
4	In vivo targeting of de novo DNA methylation by histone modifications in yeast and mouse. <i>ELife</i> , 2015, 4, e06205.	2.8	146
5	Scl Represses Cardiomyogenesis in Prospective Hemogenic Endothelium and Endocardium. <i>Cell</i> , 2012, 150, 590-605.	13.5	142
6	Adenovirus Small E1A Employs the Lysine Acetylases p300/CBP and Tumor Suppressor Rb to Repress Select Host Genes and Promote Productive Virus Infection. <i>Cell Host and Microbe</i> , 2014, 16, 663-676.	5.1	88
7	New Small Nuclear RNA Gene-Like Transcriptional Units as Sources of Regulatory Transcripts. <i>PLoS Genetics</i> , 2007, 3, e1.	1.5	82
8	Polycomb Repressive Complex 1 (PRC1) Disassembles RNA Polymerase II Preinitiation Complexes. <i>Journal of Biological Chemistry</i> , 2012, 287, 35784-35794.	1.6	66
9	TFIIIC Binding to Alu Elements Controls Gene Expression via Chromatin Looping and Histone Acetylation. <i>Molecular Cell</i> , 2020, 77, 475-487.e11.	4.5	65
10	Scl binds to primed enhancers in mesoderm to regulate hematopoietic and cardiac fate divergence. <i>EMBO Journal</i> , 2015, 34, 759-777.	3.5	64
11	Reorganization of the host epigenome by a viral oncogene. <i>Genome Research</i> , 2012, 22, 1212-1221.	2.4	61
12	Distinct roles of transcription factors TFIIB and TFIIC in RNA polymerase III transcription reinitiation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 13442-13447.	3.3	60
13	A Composite Upstream Sequence Motif Potentiates tRNA Gene Transcription in Yeast. <i>Journal of Molecular Biology</i> , 2003, 333, 1-20.	2.0	54
14	Mef2C is a lineage-restricted target of Scl/Tal1 and regulates megakaryopoiesis and B-cell homeostasis. <i>Blood</i> , 2009, 113, 3461-3471.	0.6	51
15	Transcription reinitiation by RNA polymerase III. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2013, 1829, 331-341.	0.9	50
16	Hormone-control regions mediate steroid receptor-dependent genome organization. <i>Genome Research</i> , 2019, 29, 29-39.	2.4	49
17	Genome-wide location analysis reveals a role for Sub1 in RNA polymerase III transcription. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 14265-14270.	3.3	47
18	Viral manipulation of the host epigenome for oncogenic transformation. <i>Nature Reviews Genetics</i> , 2009, 10, 290-294.	7.7	47

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19	Abf1 and other general regulatory factors control ribosome biogenesis gene expression in budding yeast. <i>Nucleic Acids Research</i> , 2017, 45, 4493-4506.	6.5	41
20	Genome-Wide Binding Map of the HIV-1 Tat Protein to the Human Genome. <i>PLoS ONE</i> , 2011, 6, e26894.	1.1	40
21	Modeling the regulatory network of histone acetylation in <i>Saccharomyces cerevisiae</i> . <i>Molecular Systems Biology</i> , 2007, 3, 153.	3.2	32
22	Mediator and SAGA Have Distinct Roles in Pol II Preinitiation Complex Assembly and Function. <i>Cell Reports</i> , 2012, 2, 1061-1067.	2.9	28
23	Epigenetic Analysis: ChIP-chip and ChIP-seq. <i>Methods in Molecular Biology</i> , 2012, 802, 377-387.	0.4	28
24	A Minimal Promoter for TFIIIC-dependent In Vitro Transcription of snoRNA and tRNA Genes by RNA Polymerase III. <i>Journal of Biological Chemistry</i> , 2006, 281, 23945-23957.	1.6	26
25	Retrotransposons as Drivers of Mammalian Brain Evolution. <i>Life</i> , 2021, 11, 376.	1.1	24
26	A unique epigenetic signature is associated with active DNA replication loci in human embryonic stem cells. <i>Epigenetics</i> , 2014, 9, 257-267.	1.3	23
27	C/EBP β mediates the growth inhibitory effect of progestins on breast cancer cells. <i>EMBO Journal</i> , 2019, 38, e101426.	3.5	15
28	Distinct modes of TATA box utilization by the RNA polymerase III transcription machineries from budding yeast and higher plants. <i>Gene</i> , 2006, 379, 12-25.	1.0	14
29	Transcription reinitiation properties of bacteriophage T7 RNA polymerase. <i>Biochemical and Biophysical Research Communications</i> , 2004, 315, 376-380.	1.0	13
30	A set of accessible enhancers enables the initial response of breast cancer cells to physiological progestin concentrations. <i>Nucleic Acids Research</i> , 2021, 49, 12716-12731.	6.5	13
31	Epigenetic changes mediated by polycomb repressive complex 2 and E2a are associated with drug resistance in a mouse model of lymphoma. <i>Genome Medicine</i> , 2016, 8, 54.	3.6	12
32	The transcription reinitiation properties of RNA polymerase III in the absence of transcription factors. <i>Cellular and Molecular Biology Letters</i> , 2008, 13, 112-8.	2.7	9
33	An RNA Polymerase III General Transcription Factor Engages in Cell Type-Specific Chromatin Looping. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2260.	1.8	4
34	Latent Cardiogenic Potential in Endocardium and Hemogenic Endothelium Revealed in the Absence of Scl/tal1. <i>Blood</i> , 2011, 118, 2362-2362.	0.6	1
35	Scl represses cardiogenesis via distant enhancers during hemogenic endothelium specification. <i>Experimental Hematology</i> , 2013, 41, S16.	0.2	0
36	The third (III) road to cell transformation. <i>Cell Cycle</i> , 2018, 17, 410-411.	1.3	0

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37	Mef2C Is a Lineage-Restricted Target Gene of Scl/Tal1 and Regulates Megakaryopoiesis and B-Cell Homeostasis. Blood, 2008, 112, 278-278.	0.6	0
38	Specification and Maintenance of the Scl Induced Hematopoietic Stem Cell Fate.. Blood, 2009, 114, 1504-1504.	0.6	0
39	Scl/Tal1 Directly Activates Hematopoiesis and Represses Cardiogenesis During Mesodermal Diversification. Blood, 2012, 120, 3446-3446.	0.6	0