

Roberto Ferrari

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

36
papers

1,494
citations

21
h-index

38
g-index

43
ext. papers

1,790
ext. citations

11.7
avg, IF

4.01
L-index

#	Paper	IF	Citations
36	A set of accessible enhancers enables the initial response of breast cancer cells to physiological progestin concentrations. <i>Nucleic Acids Research</i> , 2021 ,	20.1	2
35	Retrotransposons as Drivers of Mammalian Brain Evolution. <i>Life</i> , 2021 , 11,	3	8
34	TFIIIC Binding to Alu Elements Controls Gene Expression via Chromatin Looping and Histone Acetylation. <i>Molecular Cell</i> , 2020 , 77, 475-487.e11	17.6	29
33	C/EBP β mediates the growth inhibitory effect of progestins on breast cancer cells. <i>EMBO Journal</i> , 2019 , 38, e101426	13	10
32	Effects of Serelaxin in Patients with Acute Heart Failure. <i>New England Journal of Medicine</i> , 2019 , 381, 716-726	59.2	92
31	Hormone-control regions mediate steroid receptor-dependent genome organization. <i>Genome Research</i> , 2019 , 29, 29-39	9.7	28
30	Abf1 and other general regulatory factors control ribosome biogenesis gene expression in budding yeast. <i>Nucleic Acids Research</i> , 2017 , 45, 4493-4506	20.1	19
29	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	14.4	7
28	Scl binds to primed enhancers in mesoderm to regulate hematopoietic and cardiac fate divergence. <i>EMBO Journal</i> , 2015 , 34, 759-77	13	50
27	In vivo targeting of de novo DNA methylation by histone modifications in yeast and mouse. <i>ELife</i> , 2015 , 4, e06205	8.9	107
26	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, 663-76	23.4	72
25	A unique epigenetic signature is associated with active DNA replication loci in human embryonic stem cells. <i>Epigenetics</i> , 2014 , 9, 257-67	5.7	17
24	Transcription reinitiation by RNA polymerase III. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2013 , 1829, 331-41	6	36
23	Histone acetylation regulates intracellular pH. <i>Molecular Cell</i> , 2013 , 49, 310-21	17.6	171
22	Scl represses cardiomyogenesis in prospective hemogenic endothelium and endocardium. <i>Cell</i> , 2012 , 150, 590-605	56.2	121
21	Mediator and SAGA have distinct roles in Pol II preinitiation complex assembly and function. <i>Cell Reports</i> , 2012 , 2, 1061-7	10.6	24
20	Polycomb repressive complex 1 (PRC1) disassembles RNA polymerase II preinitiation complexes. <i>Journal of Biological Chemistry</i> , 2012 , 287, 35784-94	5.4	50

19	Epigenetic analysis: CHIP-chip and CHIP-seq. <i>Methods in Molecular Biology</i> , 2012 , 802, 377-87	1.4	21
18	Reorganization of the host epigenome by a viral oncogene. <i>Genome Research</i> , 2012 , 22, 1212-21	9.7	51
17	Scl/Tal1 Directly Activates Hematopoiesis and Represses Cardiogenesis During Mesodermal Diversification. <i>Blood</i> , 2012 , 120, 3446-3446	2.2	
16	Genome-wide binding map of the HIV-1 Tat protein to the human genome. <i>PLoS ONE</i> , 2011 , 6, e26894	3.7	27
15	Latent Cardiogenic Potential in Endocardium and Hemogenic Endothelium Revealed in the Absence of Scl/tal1. <i>Blood</i> , 2011 , 118, 2362-2362	2.2	
14	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, 14265-70	11.5	43
13	Viral manipulation of the host epigenome for oncogenic transformation. <i>Nature Reviews Genetics</i> , 2009 , 10, 290-4	30.1	44
12	Mef2C is a lineage-restricted target of Scl/Tal1 and regulates megakaryopoiesis and B-cell homeostasis. <i>Blood</i> , 2009 , 113, 3461-71	2.2	40
11	Specification and Maintenance of the Scl Induced Hematopoietic Stem Cell Fate.. <i>Blood</i> , 2009 , 114, 1504-1504		
10	Epigenetic reprogramming by adenovirus e1a. <i>Science</i> , 2008 , 321, 1086-8	33.3	179
9	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	8.1	8
8	Mef2C Is a Lineage-Restricted Target Gene of Scl/Tal1 and Regulates Megakaryopoiesis and B-Cell Homeostasis. <i>Blood</i> , 2008 , 112, 278-278	2.2	
7	New small nuclear RNA gene-like transcriptional units as sources of regulatory transcripts. <i>PLoS Genetics</i> , 2007 , 3, e1	6	76
6	Modeling the regulatory network of histone acetylation in <i>Saccharomyces cerevisiae</i> . <i>Molecular Systems Biology</i> , 2007 , 3, 153	12.2	18
5	A minimal promoter for TFIIC-dependent in vitro transcription of snoRNA and tRNA genes by RNA polymerase III. <i>Journal of Biological Chemistry</i> , 2006 , 281, 23945-57	5.4	22
4	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	3.8	13
3	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, 13442-7	11.5	48
2	Transcription reinitiation properties of bacteriophage T7 RNA polymerase. <i>Biochemical and Biophysical Research Communications</i> , 2004 , 315, 376-80	3.4	11

1 A composite upstream sequence motif potentiates tRNA gene transcription in yeast. *Journal of Molecular Biology*, **2003**, 333, 1-20

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