## **Achim Breiling**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6189542/publications.pdf

Version: 2024-02-01

26 papers 2,837 citations

361045 20 h-index 28 g-index

28 all docs

28 docs citations

times ranked

28

5582 citing authors

#	Article	IF	CITATIONS
1	Combined Deficiency of Tet1 and Tet2 Causes Epigenetic Abnormalities but Is Compatible with Postnatal Development. Developmental Cell, 2013, 24, 310-323.	3.1	379
2	Loss of Tet Enzymes Compromises Proper Differentiation of Embryonic Stem Cells. Developmental Cell, 2014, 29, 102-111.	3.1	274
3	Epigenetically Deregulated microRNA-375 Is Involved in a Positive Feedback Loop with Estrogen Receptor α in Breast Cancer Cells. Cancer Research, 2010, 70, 9175-9184.	0.4	260
4	Epigenetic regulatory functions of DNA modifications: 5-methylcytosine and beyond. Epigenetics and Chromatin, 2015, 8, 24.	1.8	249
5	General transcription factors bind promoters repressed by Polycomb group proteins. Nature, 2001, 412, 651-655.	13.7	231
6	Chromatin-associated RNA interference components contribute to transcriptional regulation in Drosophila. Nature, 2011, 480, 391-395.	13.7	203
7	Genome-wide DNA-methylation landscape defines specialization of regulatory T cells in tissues. Nature Immunology, 2017, 18, 1160-1172.	7.0	193
8	Genome-wide promoter DNA methylation dynamics of human hematopoietic progenitor cells during differentiation and aging. Blood, 2011, 117, e182-e189.	0.6	177
9	Noncoding RNA synthesis and loss of Polycomb group repression accompanies the colinear activation of the human HOXA cluster. Rna, 2006, 13, 223-239.	1.6	113
10	Drosophila Chromosome Condensation Proteins Topoisomerase II and Barren Colocalize with Polycomb and Maintain Fab-7 PRE Silencing. Molecular Cell, 2001, 7, 127-136.	4.5	110
11	Chronic Inflammation Induces a Novel Epigenetic Program That Is Conserved in Intestinal Adenomas and in Colorectal Cancer. Cancer Research, 2015, 75, 2120-2130.	0.4	91
12	Transcriptional programs that control expression of the autoimmune regulator gene Aire. Nature Immunology, 2017, 18, 161-172.	7.0	81
13	DNA (de)methylation in embryonic stem cells controls CTCF-dependent chromatin boundaries. Genome Research, 2019, 29, 750-761.	2.4	76
14	Hydroxylation of 5-methylcytosine by TET2 maintains the active state of the mammalian HOXA cluster. Nature Communications, 2012, 3, 818.	5.8	65
15	Tet1 and Tet2 Protect DNA Methylation Canyons against Hypermethylation. Molecular and Cellular Biology, 2016, 36, 452-461.	1.1	54
16	The <i>Drosophila</i> Polycomb Protein Interacts with Nucleosomal Core Particles In Vitro via Its Repression Domain. Molecular and Cellular Biology, 1999, 19, 8451-8460.	1.1	47
17	Epigenome changes in active and inactive Polycombâ€groupâ€controlled regions. EMBO Reports, 2004, 5, 976-982.	2.0	46
18	Biology of Polycomb and Trithorax Group Proteins. International Review of Cytology, 2007, 258, 83-136.	6.2	42

## ACHIM BREILING

#	Article	lF	CITATION
19	Rbpj expression in regulatory T cells is critical for restraining TH2 responses. Nature Communications, 2019, 10, 1621.	5.8	41
20	Nucleoside Drugs Induce Cellular Differentiation by Caspase-Dependent Degradation of Stem Cell Factors. PLoS ONE, 2010, 5, e10726.	1.1	38
21	MYC-Induced Epigenetic Activation of GATA4 in Lung Adenocarcinoma. Molecular Cancer Research, 2013, 11, 161-172.	1.5	15
22	SET domain proteins reSET gene expression. Nature Structural Biology, 2002, 9, 894-896.	9.7	12
23	Embryonic Carcinoma Cells Show Specific Dielectric Resistance Profiles during Induced Differentiation. PLoS ONE, 2013, 8, e59895.	1.1	12
24	Chromatin Immunoprecipitation. Methods in Molecular Biology, 2016, 1480, 7-21.	0.4	11
25	RNA-Interference Components Are Dispensable for Transcriptional Silencing of the Drosophila Bithorax-Complex. PLoS ONE, 2013, 8, e65740.	1.1	7
26	Measurement of Cellular Behavior by Electrochemical Impedance Sensing. Methods in Molecular Biology, 2017, 1601, 267-273.	0.4	2