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List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8491316/publications.pdf

Version: 2024-02-01

38 papers 2,345 citations

304743 22 h-index 377865 34 g-index

40 all docs

40 docs citations

40 times ranked

4963 citing authors

#	Article	IF	CITATIONS
1	Epigenetic mechanisms in neurological diseases: genes, syndromes, and therapies. Lancet Neurology, The, 2009, 8, 1056-1072.	10.2	528
2	Quantitative comparison of DNA methylation assays for biomarker development and clinical applications. Nature Biotechnology, 2016, 34, 726-737.	17.5	270
3	Aberrant DNA methylation patterns of spermatozoa in men with unexplained infertility. Human Reproduction, 2015, 30, 1014-1028.	0.9	144
4	DNA methylation map of mouse and human brain identifies target genes in Alzheimer's disease. Brain, 2013, 136, 3018-3027.	7.6	129
5	Disrupted microRNA expression caused by Mecp2 loss in a mouse model of Rett syndrome. Epigenetics, 2010, 5, 656-663.	2.7	125
6	H3K4me1 marks DNA regions hypomethylated during aging in human stem and differentiated cells. Genome Research, 2015, 25, 27-40.	5.5	119
7	Mecp2-Null Mice Provide New Neuronal Targets for Rett Syndrome. PLoS ONE, 2008, 3, e3669.	2.5	106
8	DNA Methylation Signatures Identify Biologically Distinct Thyroid Cancer Subtypes. Journal of Clinical Endocrinology and Metabolism, 2013, 98, 2811-2821.	3.6	100
9	A promoter DNA demethylation landscape of human hematopoietic differentiation. Nucleic Acids Research, 2012, 40, 116-131.	14.5	97
10	Immune-Dependent and Independent Antitumor Activity of GM-CSF Aberrantly Expressed by Mouse and Human Colorectal Tumors. Cancer Research, 2013, 73, 395-405.	0.9	69
11	Multiâ€omic rejuvenation of naturally aged tissues by a single cycle of transient reprogramming. Aging Cell, 2022, 21, e13578.	6.7	60
12	The RNA-binding protein HuR regulates DNA methylation through stabilization of DNMT3b mRNA. Nucleic Acids Research, 2009, 37, 2658-2671.	14.5	56
13	Aberrant epigenetic regulation of bromodomain Brd4 in human colon cancer. Journal of Molecular Medicine, 2012, 90, 587-595.	3.9	50
14	Epigenetic downregulation of TET3 reduces genomeâ€wide 5hmC levels and promotes glioblastoma tumorigenesis. International Journal of Cancer, 2020, 146, 373-387.	5.1	45
15	Longitudinal genome-wide DNA methylation analysis uncovers persistent early-life DNA methylation changes. Journal of Translational Medicine, 2019, 17, 15.	4.4	44
16	Frequent aberrant expression of the human ether \tilde{A} go-go (hEAG1) potassium channel in head and neck cancer: pathobiological mechanisms and clinical implications. Journal of Molecular Medicine, 2012, 90, 1173-1184.	3.9	43
17	DNA methylation changes in human lung epithelia cells exposed to multi-walled carbon nanotubes. Nanotoxicology, 2017, 11, 857-870.	3.0	36
18	Age-associated hydroxymethylation in human bone-marrow mesenchymal stem cells. Journal of Translational Medicine, 2016, 14, 207.	4.4	33

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19	A DNA methylation signature associated with the epigenetic repression of glycine N-methyltransferase in human hepatocellular carcinoma. Journal of Molecular Medicine, 2013, 91, 939-950.	3.9	29
20	Longitudinal study of DNA methylation during the first 5Âyears of life. Journal of Translational Medicine, 2016, 14, 160.	4.4	29
21	Epigenetic Profiling and Response to CD19 Chimeric Antigen Receptor T-Cell Therapy in B-Cell Malignancies. Journal of the National Cancer Institute, 2022, 114, 436-445.	6.3	29
22	Epigenetic dysregulation of <i>TET2</i> ii human glioblastoma. Oncotarget, 2018, 9, 25922-25934.	1.8	29
23	Loss of 5hmC identifies a new type of aberrant DNA hypermethylation in glioma. Human Molecular Genetics, 2018, 27, 3046-3059.	2.9	26
24	A DNA methylation signature associated with aberrant promoter DNA hypermethylation of DNMT3B in human colorectal cancer. European Journal of Cancer, 2012, 48, 2270-2281.	2.8	23
25	Chromatin regulation by Histone H4 acetylation at Lysine 16 during cell death and differentiation in the myeloid compartment. Nucleic Acids Research, 2019, 47, 5016-5037.	14.5	23
26	LINE-1 methylation in leukocyte DNA, interaction with phosphatidylethanolamine N-methyltransferase variants and bladder cancer risk. British Journal of Cancer, 2014, 110, 2123-2130.	6.4	17
27	The Epigenetic Basis of Adaptation and Responses to Environmental Change: Perspective on Human Reproduction. Advances in Experimental Medicine and Biology, 2014, 753, 97-117.	1.6	15
28	Histone H3 and H4 Modification Profiles in a Rett Syndrome Mouse Model. Epigenetics, 2007, 2, 11-14.	2.7	12
29	Physical exercise shapes the mouse brain epigenome. Molecular Metabolism, 2021, 54, 101398.	6.5	12
30	Epigenome-wide analysis reveals specific DNA hypermethylation of T cells during human hematopoietic differentiation. Epigenomics, 2018, 10, 903-923.	2.1	11
31	HERG1A potassium channel is the predominant isoform in head and neck squamous cell carcinomas: evidence for regulation by epigenetic mechanisms. Scientific Reports, 2016, 6, 19666.	3.3	10
32	Deregulation of the imprinted DLK1-DIO3 locus ncRNAs is associated with replicative senescence of human adipose-derived stem cells. PLoS ONE, 2018, 13, e0206534.	2.5	9
33	Epigenetic Deregulation of the Histone Methyltransferase KMT5B Contributes to Malignant Transformation in Glioblastoma. Frontiers in Cell and Developmental Biology, 2021, 9, 671838.	3.7	6
34	Conservation of Aging and Cancer Epigenetic Signatures across Human and Mouse. Molecular Biology and Evolution, 2021, 38, 3415-3435.	8.9	5
35	Epigenetics, Inflammation, and Aging. , 2014, , 85-101.		4
36	Role of Epigenetics in Neural Differentiation: Implications for Health and Disease., 2014,, 63-79.		2

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37	801 Frequent Aberrant Expression of the Human Ether a Go-go (hEAG1) Potassium Channel in Head and Neck Cancer –Pathobiological Mechanisms and Clinical Implications. European Journal of Cancer, 2012, 48, S191.	2.8	0
38	Stability of Imprinting and Differentiation Capacity in NaÃ-ve Human Cells Induced by Chemical Inhibition of CDK8 and CDK19. Cells, 2021, 10, 876.	4.1	0