John Arne Dahl

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	ALKBH5 Is a Mammalian RNA Demethylase that Impacts RNA Metabolism and Mouse Fertility. Molecular Cell, 2013, 49, 18-29.	9.7	2,549
2	Broad histone H3K4me3 domains in mouse oocytes modulate maternal-to-zygotic transition. Nature, 2016, 537, 548-552.	27.8	484
3	A rapid micro chromatin immunoprecipitation assay (ChIP). Nature Protocols, 2008, 3, 1032-1045.	12.0	259
4	Highâ€resolution analysis of genetic stability of human adipose tissue stem cells cultured to senescence. Journal of Cellular and Molecular Medicine, 2008, 12, 553-563.	3.6	148
5	Q2ChIP, a Quick and Quantitative Chromatin Immunoprecipitation Assay, Unravels Epigenetic Dynamics of Developmentally Regulated Genes in Human Carcinoma Cells. Stem Cells, 2007, 25, 1037-1046.	3.2	137
6	ALKBH1 is a Histone H2A Dioxygenase Involved in Neural Differentiation. Stem Cells, 2012, 30, 2672-2682.	3.2	97
7	Dynamic RNA modifications in disease. Current Opinion in Genetics and Development, 2014, 26, 47-52.	3.3	92
8	Histone H3 Lysine 27 Methylation Asymmetry on Developmentally-Regulated Promoters Distinguish the First Two Lineages in Mouse Preimplantation Embryos. PLoS ONE, 2010, 5, e9150.	2.5	91
9	A novel method for the efficient and selective identification of 5-hydroxymethylcytosine in genomic DNA. Nucleic Acids Research, 2011, 39, e55-e55.	14.5	88
10	Sprouts of RNA epigenetics. RNA Biology, 2013, 10, 915-918.	3.1	85
11	μChIP—a rapid micro chromatin immunoprecipitation assay for small cell samples and biopsies. Nucleic Acids Research, 2008, 36, e15.	14.5	78
12	KDM4A regulates the maternal-to-zygotic transition by protecting broad H3K4me3 domains from H3K9me3 invasion in oocytes. Nature Cell Biology, 2020, 22, 380-388.	10.3	77
13	Pull-down of 5-hydroxymethylcytosine DNA using JBP1-coated magnetic beads. Nature Protocols, 2012, 7, 340-350.	12.0	56
14	Parental micronutrient deficiency distorts liver DNA methylation and expression of lipid genes associated with a fatty-liver-like phenotype in offspring. Scientific Reports, 2018, 8, 3055.	3.3	50
15	A quick and quantitative chromatin immunoprecipitation assay for small cell samples. Frontiers in Bioscience - Landmark, 2007, 12, 4925.	3.0	44
16	Persisting symptoms three to eight months after non-hospitalized COVID-19, a prospective cohort study. PLoS ONE, 2021, 16, e0256142.	2.5	39
17	Epigenetic age is a cellâ€intrinsic property in transplanted human hematopoietic cells. Aging Cell, 2019, 18, e12897.	6.7	39
18	Parental vitamin deficiency affects the embryonic gene expression of immune-, lipid transport- and apolipoprotein genes. Scientific Reports, 2016, 6, 34535.	3.3	37

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19	Fast genomic μChIP-chip from 1,000 cells. Genome Biology, 2009, 10, R13.	9.6	35
20	Reversible RNA modifications in meiosis and pluripotency. Nature Methods, 2017, 14, 18-22.	19.0	33
21	Histone modifications and mRNA expression in the inner cell mass and trophectoderm of bovine blastocysts. Epigenetics, 2013, 8, 281-289.	2.7	32
22	μChIP: Chromatin Immunoprecipitation for Small Cell Numbers. Methods in Molecular Biology, 2009, 567, 59-74.	0.9	27
23	On the way to reprogramming cells to pluripotency using cell-free extracts. Reproductive BioMedicine Online, 2006, 12, 762-770.	2.4	19
24	Sensitive on-chip quantitative real-time PCR performed on an adaptable and robust platform. Biomedical Microdevices, 2008, 10, 769-776.	2.8	19
25	Positioning Europe for the EPITRANSCRIPTOMICS challenge. RNA Biology, 2018, 15, 1-3.	3.1	18
26	LSD1 represses a neonatal/reparative gene program in adult intestinal epithelium. Science Advances, 2020, 6, .	10.3	18
27	Persistence of Collagen Type II Synthesis and Secretion in Rapidly Proliferating Human Articular Chondrocytes <i>In Vitro</i> . Tissue Engineering - Part A, 2008, 14, 1999-2007.	3.1	16
28	DNA base modifications in honey bee and fruit fly genomes suggest an active demethylation machinery with species- and tissue-specific turnover rates. Biochemistry and Biophysics Reports, 2016, 6, 9-15.	1.3	16
29	How low can you go? Pushing the limits of low-input ChIP-seq. Briefings in Functional Genomics, 2018, 17, 89-95.	2.7	15
30	Intestinal-epithelial LSD1 controls goblet cell maturation and effector responses required for gut immunity to bacterial and helminth infection. PLoS Pathogens, 2021, 17, e1009476.	4.7	13
31	Micro Chromatin Immunoprecipitation (μChIP) from Early Mammalian Embryos. Methods in Molecular Biology, 2015, 1222, 227-245.	0.9	11
32	Histone Methylations Define Neural Stem/Progenitor Cell Subtypes in the Mouse Subventricular Zone. Molecular Neurobiology, 2020, 57, 997-1008.	4.0	10
33	Genome-wide profiling of DNA 5-hydroxymethylcytosine during rat Sertoli cell maturation. Cell Discovery, 2017, 3, 17013.	6.7	8
34	5-hydroxymethylcytosine Marks Mammalian Origins Acting as a Barrier to Replication. Scientific Reports, 2019, 9, 11065.	3.3	8
35	Going low to reach high: Smallâ€scale ChIPâ€seq maps new terrain. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 2020, 12, e1465.	6.6	8
36	The use of public transport and contraction of SARS-CoV-2 in a large prospective cohort in Norway. BMC Infectious Diseases, 2022, 22, 252.	2.9	8

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37	Analysis of epigenetic aging <i>in vivo</i> and <i>in vitro</i> : Factors controlling the speed and direction. Experimental Biology and Medicine, 2020, 245, 1543-1551.	2.4	7
38	Bases of DNA repair and regulation. Nature Chemical Biology, 2014, 10, 487-488.	8.0	5
39	Screening bioactive food compounds in honey bees suggests curcumin blocks alcohol-induced damage to longevity and DNA methylation. Scientific Reports, 2021, 11, 19156.	3.3	5
40	ALKBH5 regulates somatic cell reprogramming in a phase-specific manner. Journal of Cell Science, 2022, 135, .	2.0	3