

Anna K Naumova

List of Publications by Year in descending order

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46
papers

2,184
citations

304743

22
h-index

243625

44
g-index

48
all docs

48
docs citations

48
times ranked

3775
citing authors

#	ARTICLE	IF	CITATIONS
1	Distinct roles of androgen receptor, estrogen receptor alpha, and BCL6 in the establishment of sex-biased DNA methylation in mouse liver. <i>Scientific Reports</i> , 2021, 11, 13766.	3.3	7
2	Treatment of Allergic Asthma with Fenretinide Formulation (LAU-7b) Downregulates ORMDL Sphingolipid Biosynthesis Regulator 3 (<i>Ormdl3</i>) Expression and Normalizes Ceramide Imbalance. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2020, 373, 476-487.	2.5	5
3	Sex Chromosomes and Sex Phenotype Contribute to Biased DNA Methylation in Mouse Liver. <i>Cells</i> , 2020, 9, 1436.	4.1	13
4	A novel exomal ATRX mutation with preferential transmission to offspring: A case report and review of the literature for transmission ratio distortion in ATRX families. <i>Molecular Medicine Reports</i> , 2020, 22, 4561-4566.	2.4	1
5	A novel exomal ATRX mutation with preferential transmission to offspring: A case report and review of the literature for transmission ratio distortion in ATRX families. <i>Molecular Medicine Reports</i> , 2020, 22, 4561-4566.	2.4	2
6	Regulatory interaction between the ZBP2-ORMDL3/Zbp2-Ormdl3 region and the circadian clock. <i>PLoS ONE</i> , 2019, 14, e0223212.	2.5	3
7	Loss of the zona pellucida-binding protein 2 (<i>Zbp2</i>) gene in mice impacts airway hypersensitivity and lung lipid metabolism in a sex-dependent fashion. <i>Mammalian Genome</i> , 2018, 29, 281-298.	2.2	7
8	X chromosome dosage and presence of SRY shape sex-specific differences in DNA methylation at an autosomal region in human cells. <i>Biology of Sex Differences</i> , 2018, 9, 10.	4.1	20
9	When Does the Epigenome Become "Sexy"? <i>BioEssays</i> , 2018, 40, e1800120.	2.5	0
10	Genetic and Epigenetic Variation at the H19 Imprinted Region and Its Effect on Birth Weight. , 2017, , 185-207.		0
11	Role of DNA methylation in expression control of the IKZF3-GSDMA region in human epithelial cells. <i>PLoS ONE</i> , 2017, 12, e0172707.	2.5	31
12	Local genotype influences DNA methylation at two asthma-associated regions, 5q31 and 17q21, in a founder effect population. <i>Journal of Medical Genetics</i> , 2016, 53, 232-241.	3.2	17
13	Robertsonian translocations modify genomic distribution of ³ H2AFX and H3.3 in mouse germ cells. <i>Mammalian Genome</i> , 2016, 27, 225-236.	2.2	5
14	Genomic Imprinting Variations in the Mouse Type 3 Deiodinase Gene Between Tissues and Brain Regions. <i>Molecular Endocrinology</i> , 2014, 28, 1875-1886.	3.7	34
15	Transgenerational Epigenetic Effects and Complex Inheritance Patterns. , 2013, , 107-129.		1
16	Sex- and age-dependent DNA methylation at the 17q12-q21 locus associated with childhood asthma. <i>Human Genetics</i> , 2013, 132, 811-822.	3.8	59
17	Oocyte heterogeneity with respect to the meiotic silencing of unsynapsed X chromosomes in the XY female mouse. <i>Chromosoma</i> , 2013, 122, 337-349.	2.2	11
18	Dynamics of Response to Asynapsis and Meiotic Silencing in Spermatocytes from Robertsonian Translocation Carriers. <i>PLoS ONE</i> , 2013, 8, e75970.	2.5	15

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19	<i>DNA methyltransferase 1 (Dnmt1) mutation affects Snrpn imprinting in the mouse male germ line. Genome, 2012, 55, 673-682.</i>	2.0	5
20	Interaction between genetic and epigenetic variation defines gene expression patterns at the asthma-associated locus 17q12-q21 in lymphoblastoid cell lines. <i>Human Genetics, 2012, 131, 1161-1171.</i>	3.8	55
21	Defective imprint resetting in carriers of Robertsonian translocation Rb (8.12). <i>Mammalian Genome, 2010, 21, 377-387.</i>	2.2	11
22	Cell culture-induced aberrant methylation of the imprinted IG DMR in human lymphoblastoid cell lines. <i>Epigenetics, 2010, 5, 50-60.</i>	2.7	30
23	Global patterns of cis variation in human cells revealed by high-density allelic expression analysis. <i>Nature Genetics, 2009, 41, 1216-1222.</i>	21.4	206
24	Allele-Specific Chromatin Remodeling in the ZBP2/GSDMB/ORMDL3 Locus Associated with the Risk of Asthma and Autoimmune Disease. <i>American Journal of Human Genetics, 2009, 85, 377-393.</i>	6.2	262
25	Novel imprinted transcripts from the <i>Dlk1-Gtl2</i> intergenic region, <i>Mico1</i> and <i>Mico1os</i> , show circadian oscillations. <i>Epigenetics, 2008, 3, 322-329.</i>	2.7	20
26	Coordinated diurnal regulation of genes from the <i>Dlk1-Dio3</i> imprinted domain: implications for regulation of clusters of non-paralogous genes. <i>Human Molecular Genetics, 2008, 17, 15-26.</i>	2.9	21
27	Parental Effect of DNA (Cytosine-5) Methyltransferase 1 on Grandparental-Origin-Dependent Transmission Ratio Distortion in Mouse Crosses and Human Families. <i>Genetics, 2008, 178, 35-45.</i>	2.9	12
28	Long-QT Syndrome. <i>New England Journal of Medicine, 2007, 356, 1680-1680.</i>	27.0	1
29	Screening for pain phenotypes: Analysis of three congenic mouse strains on a battery of nine nociceptive assays. <i>Pain, 2006, 126, 24-34.</i>	4.2	70
30	Transmission ratio distortion in the myotonic dystrophy locus in human preimplantation embryos. <i>European Journal of Human Genetics, 2006, 14, 299-306.</i>	2.8	26
31	Twin study of genetic and aging effects on X chromosome inactivation. <i>European Journal of Human Genetics, 2005, 13, 599-606.</i>	2.8	100
32	Increased plasticity of genomic imprinting of <i>Dlk1</i> in brain is due to genetic and epigenetic factors. <i>Mammalian Genome, 2005, 16, 127-135.</i>	2.2	22
33	A longitudinal study of X-inactivation ratio in human females. <i>Human Genetics, 2004, 115, 387-92.</i>	3.8	56
34	A survey of genetic and epigenetic variation affecting human gene expression. <i>Physiological Genomics, 2004, 16, 184-193.</i>	2.3	228
35	Alternative splicing and imprinting control of the <i>Meg3/Gtl2-Dlk1</i> locus in mouse embryos. <i>Mammalian Genome, 2003, 14, 231-241.</i>	2.2	32
36	Inheritance patterns of maternal alleles in imprinted regions of the mouse genome at different stages of development. <i>Mammalian Genome, 2002, 13, 24-29.</i>	2.2	19

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37	Imprinting and deviation from Mendelian transmission ratios. <i>Genome</i> , 2001, 44, 311-320.	2.0	37
38	Imprinting defects in mouse embryos: stochastic errors or polymorphic phenotype?. <i>Genesis</i> , 2001, 31, 11-16.	1.6	18
39	Report of 33 Novel AVPR2 Mutations and Analysis of 117 Families with X-Linked Nephrogenic Diabetes Insipidus. <i>Journal of the American Society of Nephrology: JASN</i> , 2000, 11, 1044-1054.	6.1	156
40	Pigmentary mosaicism in hypomelanosis of Ito. <i>Human Genetics</i> , 1998, 103, 441-449.	3.8	38
41	X Chromosome Inactivation in Carriers of Barth Syndrome. <i>American Journal of Human Genetics</i> , 1998, 63, 1457-1463.	6.2	71
42	A promoter mutation in the XIST gene in two unrelated families with skewed X-chromosome inactivation. <i>Nature Genetics</i> , 1997, 17, 353-356.	21.4	279
43	Confirmation of maternal transmission ratio distortion at Om and direct evidence that the maternal and paternal "DDK syndrome" genes are linked. <i>Mammalian Genome</i> , 1997, 8, 642-646.	2.2	44
44	Chromosomal Localization of Mouse and Human Genes Encoding the Splicing Factors ASF/SF2 (SFRS1) and SC-35 (SFRS2). <i>Genomics</i> , 1995, 29, 70-79.	2.9	20
45	Preferential amplification of the paternal allele of the "myc gene in human neuroblastomas. <i>Nature Genetics</i> , 1993, 4, 191-194.	21.4	65
46	A cloned unique gene of drosophila melanogaster contains a repetitive 3' end exon whose sequence is present at the 3' ends of many different mRNAs. <i>Cell</i> , 1982, 28, 365-373.	28.9	48