Kohzoh Mitsuya

List of Publications by Year in descending order

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Κομζομ Μιτειιγλ

#	Article	IF	CITATIONS
1	2-Hydroxyglutarate destabilizes chromatin regulatory landscape and lineage fidelity to promote cellular heterogeneity. Cell Reports, 2022, 38, 110220.	6.4	8
2	Menin and Menin-Associated Proteins Coregulate Cancer Energy Metabolism. Cancers, 2020, 12, 2715.	3.7	7
3	ERα-related chromothripsis enhances concordant gene transcription on chromosome 17q11.1-q24.1 in luminal breast cancer. BMC Medical Genomics, 2020, 13, 69.	1.5	6
4	Alterations in the placental methylome with maternal obesity and evidence for metabolic regulation. PLoS ONE, 2017, 12, e0186115.	2.5	89
5	Epigenetics of Human Myometrium: DNA Methylation of Genes Encoding Contraction-Associated Proteins in Term and Preterm Labor. Biology of Reproduction, 2014, 90, 98.	2.7	31
6	Role for piRNAs and Noncoding RNA in de Novo DNA Methylation of the Imprinted Mouse <i>Rasgrf1</i> Locus. Science, 2011, 332, 848-852.	12.6	341
7	Epigenetic modification of fetal baboon hepatic phosphoenolpyruvate carboxykinase following exposure to moderately reduced nutrient availability. Journal of Physiology, 2010, 588, 1349-1359.	2.9	111
8	Food Restriction in Pregnant Mice Can Induce Changes in Histone Modifications and Suppress Gene Expression in Fetus. Nucleic Acids Symposium Series, 2007, 51, 125-126.	0.3	15
9	The SRA protein Np95 mediates epigenetic inheritance by recruiting Dnmt1 to methylated DNA. Nature, 2007, 450, 908-912.	27.8	1,096
10	Narrowed abrogation of the Angelman syndrome critical interval on human chromosome 15 does not interfere with epigenotype maintenance in somatic cells. Journal of Human Genetics, 2005, 50, 124-132.	2.3	1
11	Tandem Repeat Hypothesis in Imprinting: Deletion of a Conserved Direct Repeat Element Upstream of H19 Has No Effect on Imprinting in the Igf2-H19 Region. Molecular and Cellular Biology, 2004, 24, 5650-5656.	2.3	36
12	Imprinting on distal chromosome 7 in the placenta involves repressive histone methylation independent of DNA methylation. Nature Genetics, 2004, 36, 1291-1295.	21.4	394
13	Imprinting analysis of 10 genes and/or transcripts in a 1.5-Mb MEST-flanking region at human chromosome 7q32. Genomics, 2004, 83, 402-412.	2.9	17
14	Chromosome Loops, Insulators, and Histone Methylation: New Insights into Regulation of Imprinting in Clusters. Cold Spring Harbor Symposia on Quantitative Biology, 2004, 69, 29-38.	1.1	16
15	Epigenetic asymmetry in the mammalian zygote and early embryo: relationship to lineage commitment?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2003, 358, 1403-1409.	4.0	96
16	Paternal imprints can be established on the maternal Igf2-H19 locus without altering replication timing of DNA. Human Molecular Genetics, 2003, 12, 3123-3132.	2.9	19
17	Identification and characterization of an imprinted antisense RNA (MESTIT1) in the human MEST locus on chromosome 7q32. Human Molecular Genetics, 2002, 11, 1743-1756.	2.9	54
18	Molecular Genetic Studies of Human Chromosome 7 in Russell–Silver Syndrome. Genomics, 2002, 79, 186-196.	2.9	38

Конгон Мітѕича

#	Article	IF	CITATIONS
19	Epigenetic Heterogeneity at Imprinted Loci in Normal Populations. Biochemical and Biophysical Research Communications, 2001, 283, 1124-1130.	2.1	93
20	Loss of Imprinting of Long QT Intronic Transcript 1 in Colorectal Cancer. Oncology, 2001, 60, 268-273.	1.9	37
21	Construction of 700 human/mouse A9 monochromosomal hybrids and analysis of imprinted genes on human chromosome 6. Journal of Human Genetics, 2001, 46, 137-145.	2.3	33
22	A novel maternally expressed gene, ATP10C, encodes a putative aminophospholipid translocase associated with Angelman syndrome. Nature Genetics, 2001, 28, 19-20.	21.4	136
23	Large-scale evaluation of imprinting status in the Prader-Willi syndrome region: an imprinted direct repeat cluster resembling small nucleolar RNA genes. Human Molecular Genetics, 2001, 10, 383-394.	2.9	67
24	Title is missing!. Nature Genetics, 2001, 28, 19-20.	21.4	57
25	Targeted disruption of the human LIT1 locus defines a putative imprinting control element playing an essential role in Beckwith-Wiedemann syndrome. Human Molecular Genetics, 2000, 9, 2075-2083.	2.9	159
26	Loss of imprinting of a paternally expressed transcript, with antisense orientation to K _V LQT1, occurs frequently in Beckwith–Wiedemann syndrome and is independent of insulin-like growth factor II imprinting. Proceedings of the National Academy of Sciences of the United States of America, 1999, 96, 5203-5208.	7.1	350
27	LIT1, an imprinted antisense RNA in the human KvLQT1 locus identified by screening for differentially expressed transcripts using monochromosomal hybrids. Human Molecular Genetics, 1999, 8, 1209-1217.	2.9	214
28	Epigenetic reprogramming of the humanH19gene in mouse embryonic cells does not erase the primary parental imprint. Genes To Cells, 1998, 3, 245-255.	1.2	15
29	Evidence for Uniparental, Paternal Expression of the Human GABAA Receptor Subunit Genes, Using Microcell-Mediated Chromosome Transfer. Human Molecular Genetics, 1997, 6, 2127-2133.	2.9	85
30	Paternal expression of WT1 in human fibroblasts and lymphocytes. Human Molecular Genetics, 1997, 6, 2243-2246.	2.9	39
31	Normal human chromosome 2 induces cellular senescence in the human cervical carcinoma cell line SiHa. Genes Chromosomes and Cancer, 1995, 14, 120-127.	2.8	46
32	Isolation and Mapping of 186 New DNA Markers on Human Chromosome 1. Genomics, 1995, 27, 207-210.	2.9	28
33	Isolation and Mapping of Human Homologues of an Imprinted Mouse GeneU2af1-rs1. Genomics, 1995, 30, 257-263.	2.9	35