

Mireia Jordà

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

3,520
citations

516215

16
h-index

580395

25
g-index

26
all docs

26
docs citations

26
times ranked

5601
citing authors

#	ARTICLE	IF	CITATIONS
1	Lipid Metabolism and Epigenetics Crosstalk in Prostate Cancer. <i>Nutrients</i> , 2022, 14, 851.	1.7	17
2	Laparoscopic sleeve gastrectomy induces molecular changes in peripheral white blood cells. <i>Clinical Nutrition</i> , 2020, 39, 592-598.	2.3	6
3	Hsa-miR-139-5p is a prognostic thyroid cancer marker involved in HNRNPF-mediated alternative splicing. <i>International Journal of Cancer</i> , 2020, 146, 521-530.	2.3	29
4	Tissue and cancer-specific expression of DIXF is epigenetically mediated by an Alu repeat. <i>Epigenetics</i> , 2020, 15, 765-779.	1.3	4
5	Epigenetic footprint enables molecular risk stratification of hepatoblastoma with clinical implications. <i>Journal of Hepatology</i> , 2020, 73, 328-341.	1.8	82
6	Molecular profiling for acromegaly treatment: a validation study. <i>Endocrine-Related Cancer</i> , 2020, 27, 375-389.	1.6	39
7	DNA methylation in thyroid cancer. <i>Endocrine-Related Cancer</i> , 2019, 26, R415-R439.	1.6	72
8	Kallikreins Stepwise Scoring Reveals Three Subtypes of Papillary Thyroid Cancer with Prognostic Implications. <i>Thyroid</i> , 2018, 28, 601-612.	2.4	13
9	Increased Global DNA Hypomethylation in Distant Metastatic and Dedifferentiated Thyroid Cancer. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2018, 103, 397-406.	1.8	20
10	The epigenetic landscape of <i>Alu</i> repeats delineates the structural and functional genomic architecture of colon cancer cells. <i>Genome Research</i> , 2017, 27, 118-132.	2.4	51
11	A knowledgebase of the human Alu repetitive elements. <i>Journal of Biomedical Informatics</i> , 2016, 60, 77-83.	2.5	10
12	Quantification of Unmethylated Alu (QUAlu): a tool to assess global hypomethylation in routine clinical samples. <i>Oncotarget</i> , 2016, 7, 10536-10546.	0.8	14
13	Overlapping DNA Methylation Dynamics in Mouse Intestinal Cell Differentiation and Early Stages of Malignant Progression. <i>PLoS ONE</i> , 2015, 10, e0123263.	1.1	14
14	DNA methylation profiling of well-differentiated thyroid cancer uncovers markers of recurrence free survival. <i>International Journal of Cancer</i> , 2014, 135, 598-610.	2.3	66
15	Long range epigenetic silencing is a trans-species mechanism that results in cancer specific deregulation by overriding the chromatin domains of normal cells. <i>Molecular Oncology</i> , 2013, 7, 1129-1141.	2.1	13
16	Epigenetics of Host-Pathogen Interactions. , 2013, , 1-22.		0
17	Epigenetics of Host-Pathogen Interactions: The Road Ahead and the Road Behind. <i>PLoS Pathogens</i> , 2012, 8, e1003007.	2.1	205
18	Methods for DNA methylation analysis and applications in colon cancer. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2010, 693, 84-93.	0.4	23

#	ARTICLE	IF	CITATIONS
19	Analysis of DNA Methylation by Amplification of Intermethylated Sites (AIMS). <i>Methods in Molecular Biology</i> , 2009, 507, 107-116.	0.4	11
20	Genome-wide tracking of unmethylated DNA Alu repeats in normal and cancer cells. <i>Nucleic Acids Research</i> , 2008, 36, 770-784.	6.5	94
21	Snail silencing effectively suppresses tumour growth and invasiveness. <i>Oncogene</i> , 2007, 26, 1862-1874.	2.6	239
22	Id-1 is induced in MDCK epithelial cells by activated Erk/MAPK pathway in response to expression of the Snail and E47 transcription factors. <i>Experimental Cell Research</i> , 2007, 313, 2389-2403.	1.2	34
23	Functional CpG Methylation System in a Social Insect. <i>Science</i> , 2006, 314, 645-647.	6.0	331
24	Insights into social insects from the genome of the honeybee <i>Apis mellifera</i> . <i>Nature</i> , 2006, 443, 931-949.	13.7	1,648
25	Genetic Profiling of Epithelial Cells Expressing E-Cadherin Repressors Reveals a Distinct Role for Snail, Slug, and E47 Factors in Epithelial-Mesenchymal Transition. <i>Cancer Research</i> , 2006, 66, 9543-9556.	0.4	285
26	Upregulation of MMP-9 in MDCK epithelial cell line in response to expression of the Snail transcription factor. <i>Journal of Cell Science</i> , 2005, 118, 3371-3385.	1.2	200