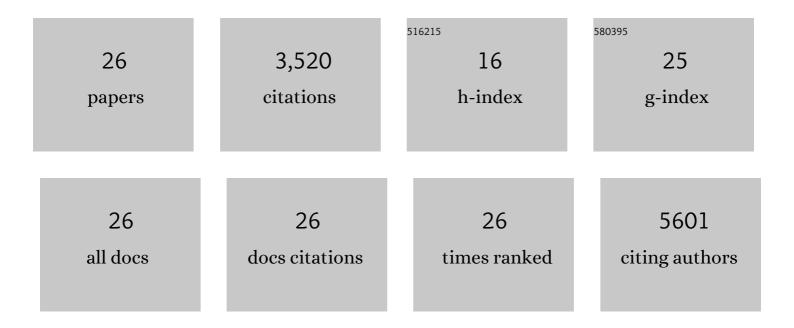
Mireia JordÃ

List of Publications by Year in descending order

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Μιρειλ ΙοροÃ

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

Mireia JordÃ

#	Article	IF	CITATIONS
19	Analysis of DNA Methylation by Amplification of Intermethylated Sites (AIMS). Methods in Molecular Biology, 2009, 507, 107-116.	0.4	11
20	Genome-wide tracking of unmethylated DNA Alu repeats in normal and cancer cells. Nucleic Acids Research, 2008, 36, 770-784.	6.5	94
21	Snail silencing effectively suppresses tumour growth and invasiveness. Oncogene, 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. Experimental Cell Research, 2007, 313, 2389-2403.	1.2	34
23	Functional CpG Methylation System in a Social Insect. Science, 2006, 314, 645-647.	6.0	331
24	Insights into social insects from the genome of the honeybee Apis mellifera. Nature, 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. Cancer Research, 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. Journal of Cell Science, 2005, 118, 3371-3385.	1.2	200