

Ana Monteagudo-Sánchez

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

342
citations

1163117

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1281871

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12
docs citations

12
times ranked

744
citing authors

#	ARTICLE	IF	CITATIONS
1	Human Oocyte-Derived Methylation Differences Persist in the Placenta Revealing Widespread Transient Imprinting. <i>PLoS Genetics</i> , 2016, 12, e1006427.	3.5	94
2	PM20D1 is a quantitative trait locus associated with Alzheimer's disease. <i>Nature Medicine</i> , 2018, 24, 598-603.	30.7	73
3	The role of ZFP57 and additional KRAB-zinc finger proteins in the maintenance of human imprinted methylation and multi-locus imprinting disturbances. <i>Nucleic Acids Research</i> , 2020, 48, 11394-11407.	14.5	32
4	Characterization of parent-of-origin methylation using the Illumina Infinium MethylationEPIC array platform. <i>Epigenomics</i> , 2018, 10, 941-954.	2.1	31
5	Differences in expression rather than methylation at placenta-specific imprinted loci is associated with intrauterine growth restriction. <i>Clinical Epigenetics</i> , 2019, 11, 35.	4.1	29
6	Epigenetic and genetic variants in the HTR1B gene and clinical improvement in children and adolescents treated with fluoxetine. <i>Progress in Neuro-Psychopharmacology and Biological Psychiatry</i> , 2017, 75, 28-34.	4.8	28
7	Copy number rather than epigenetic alterations are the major dictator of imprinted methylation in tumors. <i>Nature Communications</i> , 2017, 8, 467.	12.8	27
8	Maternal mutations of <i>FOXF1</i> cause alveolar capillary dysplasia despite not being imprinted. <i>Human Mutation</i> , 2017, 38, 615-620.	2.5	13
9	The hypomethylation of imprinted genes in IVF/ICSI placenta samples is associated with concomitant changes in histone modifications. <i>Epigenetics</i> , 2020, 15, 1386-1395.	2.7	8
10	Preimplantation genetic testing for a chr14q32 microdeletion in a family with Kagami-Ogata syndrome and Temple syndrome. <i>Journal of Medical Genetics</i> , 2022, 59, 253-261.	3.2	5
11	The Use of Methylation-Sensitive Multiplex Ligation-Dependent Probe Amplification for Quantification of Imprinted Methylation. <i>Methods in Molecular Biology</i> , 2018, 1766, 109-121.	0.9	2