Ana Monteagudo-SÃ¡nchez

List of Publications by Year
in descending order

11 papers	342 citations	1163117 8 h-index	1281871 11 g-index
12 all docs	12 docs citations	12 times ranked	744 citing authors

#	Article	IF	Citations
1	Preimplantation genetic testing for a chr14q32 microdeletion in a family with Kagami-Ogata syndrome and Temple syndrome. Journal of Medical Genetics, 2022, 59, 253-261.	3.2	5
2	The role of ZFP57 and additional KRAB-zinc finger proteins in the maintenance of human imprinted methylation and multi-locus imprinting disturbances. Nucleic Acids Research, 2020, 48, 11394-11407.	14.5	32
3	The hypomethylation of imprinted genes in IVF/ICSI placenta samples is associated with concomitant changes in histone modifications. Epigenetics, 2020, 15, 1386-1395.	2.7	8
4	Differences in expression rather than methylation at placenta-specific imprinted loci is associated with intrauterine growth restriction. Clinical Epigenetics, 2019, 11, 35.	4.1	29
5	The Use of Methylation-Sensitive Multiplex Ligation-Dependent Probe Amplification for Quantification of Imprinted Methylation. Methods in Molecular Biology, 2018, 1766, 109-121.	0.9	2
6	Characterization of parent-of-origin methylation using the Illumina Infinium MethylationEPIC array platform. Epigenomics, 2018, 10, 941-954.	2.1	31
7	PM20D1 is aÂquantitative trait locus associated with Alzheimerâ€™s disease. Nature Medicine, 2018, 24, 598-603.	30.7	73
8	Maternal mutations ofFOXF1cause alveolar capillary dysplasia despite not being imprinted. Human Mutation, 2017, 38, 615-620.	2.5	13
9	Epigenetic and genetic variants in the HTR1B gene and clinical improvement in children and adolescents treated with fluoxetine. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2017, 75, 28-34.	4.8	28
10	Copy number rather than epigenetic alterations are the major dictator of imprinted methylation in tumors. Nature Communications, 2017, 8, 467.	12.8	27
11	Human Oocyte-Derived Methylation Differences Persist in the Placenta Revealing Widespread Transient Imprinting. PLoS Genetics, 2016, 12, e1006427.	3.5	94