

CITATION REPORT

List of articles citing

Genome-wide analysis of differential DNA methylation in Silver-Russell syndrome

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Science China Life Sciences, 2017, 60, 692-699.

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#	Paper	IF	Citations
17	eRAM: encyclopedia of rare disease annotations for precision medicine. <i>Nucleic Acids Research</i> , 2018 , 46, D937-D943	20.1	21
16	RDAD: A Machine Learning System to Support Phenotype-Based Rare Disease Diagnosis. <i>Frontiers in Genetics</i> , 2018 , 9, 587	4.5	17
15	PedAM: a database for Pediatric Disease Annotation and Medicine. <i>Nucleic Acids Research</i> , 2018 , 46, D977-D983	7.0	13
14	Integrative analysis identifies potential DNA methylation biomarkers for pan-cancer diagnosis and prognosis. <i>Epigenetics</i> , 2019 , 14, 67-80	5.7	44
13	DNA methylation signatures in mendelian developmental disorders as a diagnostic bridge between genotype and phenotype. <i>Epigenomics</i> , 2019 , 11, 563-575	4.4	21
12	Epigenetic studies and pediatric research. <i>Pediatric Research</i> , 2020 , 87, 378-384	3.2	6
11	EpiSigns. <i>Advances in Molecular Pathology</i> , 2020 , 3, 29-39	0.3	2
10	Genome-wide methylation analysis in Silver-Russell syndrome, Temple syndrome, and Prader-Willi syndrome. <i>Clinical Epigenetics</i> , 2020 , 12, 159	7.7	4
9	A pilot study of expanded newborn screening for 573 genes related to severe inherited disorders in China: results from 1,127 newborns. <i>Annals of Translational Medicine</i> , 2020 , 8, 1058	3.2	7
8	The role of maternal methylation in the association between prenatal meteorological conditions and neonatal H19/H19-DMR methylation. <i>Ecotoxicology and Environmental Safety</i> , 2020 , 197, 110643	7	5
7	Patients with PWS and related syndromes display differentially methylated regions involved in neurodevelopmental and nutritional trajectory. <i>Clinical Epigenetics</i> , 2021 , 13, 159	7.7	2
6	Pan-Cancer DNA Methylation Analysis and Tumor Origin Identification of Carcinoma of Unknown Primary Site Based on Multi-Omics. <i>Frontiers in Genetics</i> , 2021 , 12, 798748	4.5	0
5	Genome-wide methylomic analyses identify prognostic epigenetic signature in lower grade glioma. <i>Journal of Cellular and Molecular Medicine</i> , 2021 ,	5.6	2
4	Data_Sheet_1.DOCX. 2018 ,		
3	Image_1.PNG. 2018 ,		
2	Image_2.PNG. 2018 ,		
1	Image_3.PNG. 2018 ,		

