

Luciano Calzari

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9232166/publications.pdf>

Version: 2024-02-01

27
papers

674
citations

687220

13
h-index

580701

25
g-index

29
all docs

29
docs citations

29
times ranked

1030
citing authors

#	ARTICLE	IF	CITATIONS
1	(Epi)genotype-phenotype correlations in Beckwith-Wiedemann syndrome. <i>European Journal of Human Genetics</i> , 2016, 24, 183-190.	1.4	113
2	Stochastic epigenetic mutations (DNA methylation) increase exponentially in human aging and correlate with X chromosome inactivation skewing in females. <i>Aging</i> , 2015, 7, 568-578.	1.4	68
3	A multi-method approach to the molecular diagnosis of overt and borderline 11p15.5 defects underlying Silver-Russell and Beckwith-Wiedemann syndromes. <i>Clinical Epigenetics</i> , 2016, 8, 23.	1.8	54
4	Involvement of the yeast metacaspase Yca1 in \dot{I}^n -programmed cell death. <i>FEMS Yeast Research</i> , 2004, 5, 141-147.	1.1	46
5	A KHDC3L mutation resulting in recurrent hydatidiform mole causes genome-wide DNA methylation loss in oocytes and persistent imprinting defects post-fertilisation. <i>Genome Medicine</i> , 2019, 11, 84.	3.6	45
6	Loss-of-function maternal-effect mutations of PADI6 are associated with familial and sporadic Beckwith-Wiedemann syndrome with multi-locus imprinting disturbance. <i>Clinical Epigenetics</i> , 2020, 12, 139.	1.8	40
7	Fetal growth patterns in Beckwith-Wiedemann syndrome. <i>Clinical Genetics</i> , 2016, 90, 21-27.	1.0	34
8	Quantitative DNA methylation analysis improves epigenotype-phenotype correlations in Beckwith-Wiedemann syndrome. <i>Epigenetics</i> , 2013, 8, 1053-1060.	1.3	33
9	DNA Methylation in the Diagnosis of Monogenic Diseases. <i>Genes</i> , 2020, 11, 355.	1.0	28
10	The phenotypic variations of multi-locus imprinting disturbances associated with maternal-effect variants of NLRP5 range from overt imprinting disorder to apparently healthy phenotype. <i>Clinical Epigenetics</i> , 2019, 11, 190.	1.8	22
11	iPSC-derived neurons of CREBBP - and EP300 -mutated Rubinstein-Taybi syndrome patients show morphological alterations and hypoexcitability. <i>Stem Cell Research</i> , 2018, 30, 130-140.	0.3	19
12	Discrepant molecular and clinical diagnoses in Beckwith-Wiedemann and Silver-Russell syndromes. <i>Genetical Research</i> , 2019, 101, e3.	0.3	17
13	Genome-Wide DNA Methylation Analysis of a Cohort of 41 Patients Affected by Oculo-Auriculo-Vertebral Spectrum (OAVS). <i>International Journal of Molecular Sciences</i> , 2021, 22, 1190.	1.8	16
14	Transcriptome Analysis of iPSC-Derived Neurons from Rubinstein-Taybi Patients Reveals Deficits in Neuronal Differentiation. <i>Molecular Neurobiology</i> , 2020, 57, 3685-3701.	1.9	15
15	Taurine Administration Recovers Motor and Learning Deficits in an Angelman Syndrome Mouse Model. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1088.	1.8	14
16	Ring Chromosome 20 Syndrome: Genetics, Clinical Characteristics, and Overlapping Phenotypes. <i>Frontiers in Neurology</i> , 2020, 11, 613035.	1.1	14
17	Trans-acting genetic variants causing multilocus imprinting disturbance (MLID): common mechanisms and consequences. <i>Clinical Epigenetics</i> , 2022, 14, 41.	1.8	14
18	The Histone Deubiquitinating Enzyme Ubp10 Is Involved in rDNA Locus Control in <i>Saccharomyces cerevisiae</i> by Affecting Sir2p Association. <i>Genetics</i> , 2006, 174, 2249-2254.	1.2	13

#	ARTICLE	IF	CITATIONS
19	Epigenome Wide Association and Stochastic Epigenetic Mutation Analysis on Cord Blood of Preterm Birth. <i>International Journal of Molecular Sciences</i> , 2020, 21, 5044.	1.8	12
20	Blood oxidative stress and metallothionein expression in Rett syndrome: Probing for markers. <i>World Journal of Biological Psychiatry</i> , 2016, 17, 198-209.	1.3	11
21	Molecular Etiology Disclosed by Array CGH in Patients With Silver-Russell Syndrome or Similar Phenotypes. <i>Frontiers in Genetics</i> , 2019, 10, 955.	1.1	11
22	Extensive Placental Methylation Profiling in Normal Pregnancies. <i>International Journal of Molecular Sciences</i> , 2021, 22, 2136.	1.8	8
23	Generation of three iPSC lines (IAli002, IAli004, IAli003) from Rubinstein-Taybi syndrome 1 patients carrying CREBBP non sense c.4435G>T, p.(Gly1479*) and c.3474G>A, p.(Trp1158*) and missense c.4627G>T, p.(Asp1543Tyr) mutations. <i>Stem Cell Research</i> , 2019, 40, 101553.	0.3	6
24	Maternal Uniparental Disomy of Chromosome 20 (UPD(20)mat) as Differential Diagnosis of Silver Russell Syndrome: Identification of Three New Cases. <i>Genes</i> , 2021, 12, 588.	1.0	6
25	Germline variants in genes of the subcortical maternal complex and Multilocus Imprinting Disturbance are associated with miscarriage/infertility or Beckwith-Wiedemann progeny. <i>Clinical Epigenetics</i> , 2022, 14, 43.	1.8	6
26	Genes for RNA-binding proteins involved in neural-specific functions and diseases are downregulated in Rubinstein-Taybi iNeurons. <i>Neural Regeneration Research</i> , 2022, 17, 5.	1.6	5
27	Interconnected Gene Networks Underpin the Clinical Overlap of HNRNPH1-Related and Rubinstein-Taybi Intellectual Disability Syndromes. <i>Frontiers in Neuroscience</i> , 2021, 15, 745684.	1.4	0