

Cecilia Mancini

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7794636/publications.pdf>

Version: 2024-02-01

28
papers

609
citations

567281

15
h-index

610901

24
g-index

28
all docs

28
docs citations

28
times ranked

1361
citing authors

#	ARTICLE	IF	CITATIONS
1	KCNK18 Biallelic Variants Associated with Intellectual Disability and Neurodevelopmental Disorders Alter TRESK Channel Activity. <i>International Journal of Molecular Sciences</i> , 2021, 22, 6064.	4.1	3
2	In vitro dexamethasone treatment does not induce alternative ATM transcripts in cells from Ataxiaâ€“Telangiectasia patients. <i>Scientific Reports</i> , 2020, 10, 20182.	3.3	3
3	Mice harbouring a SCA28 patient mutation in AFG3L2 develop late-onset ataxia associated with enhanced mitochondrial proteotoxicity. <i>Neurobiology of Disease</i> , 2019, 124, 14-28.	4.4	23
4	Spontaneous remission in a Diamondâ€“Blackfan anaemia patient due to a revertant uniparental disomy ablating a <i>de novo</i> RPS19</i> mutation. <i>British Journal of Haematology</i> , 2019, 185, 994-998.	2.5	24
5	A fetal case of microphthalmia and limb anomalies with abnormal neuronal migration associated with SMOG1 biallelic variants. <i>European Journal of Medical Genetics</i> , 2019, 62, 103578.	1.3	4
6	ATXN2 intermediate repeat expansions influence the clinical phenotype in frontotemporal dementia. <i>Neurobiology of Aging</i> , 2019, 73, 231.e7-231.e9.	3.1	21
7	Mitochondrial stress response triggered by defects in protein synthesis quality control. <i>Life Science Alliance</i> , 2019, 2, e201800219.	2.8	26
8	Spinocerebellar Ataxia Tethering PCR. <i>Journal of Molecular Diagnostics</i> , 2018, 20, 289-297.	2.8	16
9	Altered homeostasis of trace elements in the blood of SCA2 patients. <i>Journal of Trace Elements in Medicine and Biology</i> , 2018, 47, 111-114.	3.0	7
10	Updated genetic testing of Italian patients referred with a clinical diagnosis of primary hyperoxaluria. <i>Journal of Nephrology</i> , 2017, 30, 219-225.	2.0	9
11	A case of Feingold type 2 syndrome associated with keratoconus refines keratoconus type 7 locus on chromosome 13q. <i>European Journal of Medical Genetics</i> , 2017, 60, 224-227.	1.3	10
12	A novel homozygous change of <i>CLCN2</i> (p.His590Pro) is associated with a subclinical form of leukoencephalopathy with ataxia (LKPAT). <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2017, 88, 894-896.	1.9	20
13	Exome sequencing in children of women with skewed X-inactivation identifies atypical cases and complex phenotypes. <i>European Journal of Paediatric Neurology</i> , 2017, 21, 475-484.	1.6	14
14	Human canonical CD157/Bst1 is an alternatively spliced isoform masking a previously unidentified primate-specific exon included in a novel transcript. <i>Scientific Reports</i> , 2017, 7, 15923.	3.3	10
15	Cover Image, Volume 170A, Number 7, July 2016. , 2016, 170, i-i.		0
16	Whole exome sequencing is necessary to clarify ID/DD cases with <i>de novo</i> copy number variants of uncertain significance: Two proofâ€“ofâ€“concept examples. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1772-1779.	1.2	26
17	A novel 3q29 deletion associated with autism, intellectual disability, psychiatric disorders, and obesity. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 290-299.	1.7	34
18	Ribosomal <i>scp</i> RNA analysis in the diagnosis of Diamondâ€“Blackfan Anaemia. <i>British Journal of Haematology</i> , 2016, 172, 782-785.	2.5	24

#	ARTICLE	IF	CITATIONS
19	Adult-onset autosomal recessive ataxia associated with neuronal ceroid lipofuscinosis type 5 gene (CLN5) mutations. <i>Journal of Neurology</i> , 2015, 262, 173-178.	3.6	29
20	An atypical form of AOA2 with myoclonus associated with mutations in SETX and AFG3L2. <i>BMC Medical Genetics</i> , 2015, 16, 16.	2.1	12
21	Two families with novel missense mutations in COL4A1: When diagnosis can be missed. <i>Journal of the Neurological Sciences</i> , 2015, 352, 99-104.	0.6	21
22	Blood metal levels and related antioxidant enzyme activities in patients with ataxia telangiectasia. <i>Neurobiology of Disease</i> , 2015, 81, 162-167.	4.4	13
23	Large cryptic genomic rearrangements with apparently normal karyotypes detected by array-CGH. <i>Molecular Cytogenetics</i> , 2014, 7, 82.	0.9	25
24	ELOVL5 Mutations Cause Spinocerebellar Ataxia 38. <i>American Journal of Human Genetics</i> , 2014, 95, 209-217.	6.2	107
25	Genome-wide expression profiling and functional characterization of SCA28 lymphoblastoid cell lines reveal impairment in cell growth and activation of apoptotic pathways. <i>BMC Medical Genomics</i> , 2013, 6, 22.	1.5	14
26	Megalencephalic leukoencephalopathy with subcortical cysts type 1 (MLC1) due to a homozygous deep intronic splicing mutation (c.895-226T>G) abrogated in vitro using an antisense morpholino oligonucleotide. <i>Neurogenetics</i> , 2012, 13, 205-214.	1.4	21
27	Gene-targeted embryonic stem cells: real-time PCR assay for estimation of the number of neomycin selection cassettes. <i>Biological Procedures Online</i> , 2011, 13, 10.	2.9	12
28	Missense mutations in the AFG3L2 proteolytic domain account for ~1.5% of European autosomal dominant cerebellar ataxias. <i>Human Mutation</i> , 2010, 31, 1117-1124.	2.5	81