

François Lecoquierre

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

134
citations

1307594

7
h-index

1372567

10
g-index

13
all docs

13
docs citations

13
times ranked

248
citing authors

#	ARTICLE	IF	CITATIONS
1	Detection of copy-number variations from NGS data using read depth information: a diagnostic performance evaluation. <i>European Journal of Human Genetics</i> , 2021, 29, 99-109.	2.8	23
2	Variant recurrence in neurodevelopmental disorders: the use of publicly available genomic data identifies clinically relevant pathogenic missense variants. <i>Genetics in Medicine</i> , 2019, 21, 2504-2511.	2.4	21
3	MYT1L-associated neurodevelopmental disorder: description of 40 new cases and literature review of clinical and molecular aspects. <i>Human Genetics</i> , 2022, 141, 65-80.	3.8	14
4	De novo variants in <i>TCF7L2</i> are associated with a syndromic neurodevelopmental disorder. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 2384-2390.	1.2	13
5	De novo coding variants in the <i>AGO1</i> gene cause a neurodevelopmental disorder with intellectual disability. <i>Journal of Medical Genetics</i> , 2022, 59, 965-975.	3.2	13
6	Optimization of the diagnosis of inherited colorectal cancer using NGS and capture of exonic and intronic sequences of panel genes. <i>European Journal of Human Genetics</i> , 2018, 26, 1597-1602.	2.8	12
7	Confirmation and further delineation of the <i>SMC9</i> deficiency syndrome, a rare and severe developmental disorder. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 2257-2262.	1.2	9
8	Haploinsufficiency of the Primary Familial Brain Calcification Gene <i>SLC20A2</i> Mediated by Disruption of a Regulatory Element. <i>Movement Disorders</i> , 2020, 35, 1336-1345.	3.9	9
9	Excess of de novo variants in genes involved in chromatin remodelling in patients with marfanoid habitus and intellectual disability. <i>Journal of Medical Genetics</i> , 2020, 57, 466-474.	3.2	7
10	uORF-introducing variants in the 5'UTR of the <i>NIPBL</i> gene as a cause of Cornelia de Lange syndrome. <i>Human Mutation</i> , 2022, 43, 1239-1248.	2.5	6
11	Heterozygous variants in <i>PRPF8</i> are associated with neurodevelopmental disorders. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 2750-2759.	1.2	4
12	Patients with 10q22.3q23.1 recurrent deletion syndrome are at risk for juvenile polyposis. <i>European Journal of Medical Genetics</i> , 2020, 63, 103773.	1.3	3