

Natalie S Hauser

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

583
citations

758635

12
h-index

676716

22
g-index

23
all docs

23
docs citations

23
times ranked

1453
citing authors

#	ARTICLE	IF	CITATIONS
1	Novel pathogenic variants and quantitative phenotypic analyses of Robinow syndrome: WNT signaling perturbation and phenotypic variability. <i>Human Genetics and Genomics Advances</i> , 2022, 3, 100074.	1.0	14
2	Expanding the genotypic and phenotypic spectrum in a diverse cohort of 104 individuals with Wiedemann-Steiner syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 1649-1665.	0.7	34
3	Expanding the phenotype of <sc><i>ASXL3</i></sc>-related syndrome: A comprehensive description of 45 unpublished individuals with inherited and de novo pathogenic variants in <sc><i>ASXL3</i></sc>. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 3446-3458.	0.7	12
4	The natural history of infantile mitochondrial DNA depletion syndrome due to RRM2B deficiency. <i>Genetics in Medicine</i> , 2020, 22, 199-209.	1.1	14
5	<sc><i>SCN3A</i></sc>-Related Neurodevelopmental Disorder: A Spectrum of Epilepsy and Brain Malformation. <i>Annals of Neurology</i> , 2020, 88, 348-362.	2.8	42
6	Mosaicism in ASXL3-related syndrome: Description of five patients from three families. <i>European Journal of Medical Genetics</i> , 2020, 63, 103925.	0.7	9
7	EPG5 Variants with Modest Functional Impact Result in an Ameliorated and Primarily Neurological Phenotype in a 3.5-Year-Old Patient with Vici Syndrome. <i>Neuropediatrics</i> , 2019, 50, 257-261.	0.3	5
8	Genotype-first analysis of a generally healthy population cohort supports genetic testing for diagnosis of hereditary angioedema of unknown cause. <i>Allergy, Asthma and Clinical Immunology</i> , 2019, 15, 32.	0.9	12
9	Novel GABRA2 variants in epileptic encephalopathy and intellectual disability with seizures. <i>Brain</i> , 2019, 142, e15-e15.	3.7	12
10	Endosomal trafficking defects in patient cells with KIAA1109 biallelic variants. <i>Genes and Diseases</i> , 2019, 6, 56-67.	1.5	22
11	A case study of atypical Larsen syndrome with absent hallmark joint dislocations. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e648.	0.6	2
12	Pitfalls of clinical exome and gene panel testing: alternative transcripts. <i>Genetics in Medicine</i> , 2019, 21, 1240-1245.	1.1	17
13	Biochemical signatures mimicking multiple carboxylase deficiency in children with mutations in MT-ATP6. <i>Mitochondrion</i> , 2019, 44, 58-64.	1.6	19
14	Mutation in an alternative transcript of <i>CDKL5</i> in a boy with early-onset seizures. <i>Journal of Physical Education and Sports Management</i> , 2018, 4, a002360.	0.5	10
15	Experience with genomic sequencing in pediatric patients with congenital cardiac defects in a large community hospital. <i>Molecular Genetics & Genomic Medicine</i> , 2018, 6, 200-212.	0.6	12
16	FOXP1 syndrome: genotype-phenotype association in 83 patients with FOXP1 variants. <i>Genetics in Medicine</i> , 2018, 20, 98-108.	1.1	77
17	Diagnosis of LCHAD/TFP deficiency in an at risk newborn using umbilical cord blood acylcarnitine analysis. <i>Molecular Genetics and Metabolism Reports</i> , 2017, 10, 8-10.	0.4	1
18	Genomic analysis of an infant with intractable diarrhea and dilated cardiomyopathy. <i>Journal of Physical Education and Sports Management</i> , 2017, 3, a002055.	0.5	13

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19	De Novo Mutations in SON Disrupt RNA Splicing of Genes Essential for Brain Development and Metabolism, Causing an Intellectual-Disability Syndrome. American Journal of Human Genetics, 2016, 99, 711-719.	2.6	81
20	De novo missense variants in PPP2R5D are associated with intellectual disability, macrocephaly, hypotonia, and autism. Neurogenetics, 2016, 17, 43-49.	0.7	61
21	Mutation screen reveals novel variants and expands the phenotypes associated with DYNC1H1. Journal of Neurology, 2015, 262, 2124-2134.	1.8	59
22	Regions of homozygosity identified by oligonucleotide SNP arrays: evaluating the incidence and clinical utility. European Journal of Human Genetics, 2015, 23, 663-671.	1.4	54