

Farah R Zahir

List of Publications by Year in descending order

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28
papers

1,519
citations

567281

15
h-index

580821

25
g-index

29
all docs

29
docs citations

29
times ranked

3286
citing authors

#	ARTICLE	IF	CITATIONS
1	Utility of whole-exome sequencing for those near the end of the diagnostic odyssey: time to address gaps in care. <i>Clinical Genetics</i> , 2016, 89, 275-284.	2.0	323
2	Oligonucleotide Microarray Analysis of Genomic Imbalance in Children with Mental Retardation. <i>American Journal of Human Genetics</i> , 2006, 79, 500-513.	6.2	261
3	Mutations in NGLY1 cause an inherited disorder of the endoplasmic reticulum-associated degradation pathway. <i>Genetics in Medicine</i> , 2014, 16, 751-758.	2.4	191
4	A patient with vertebral, cognitive and behavioural abnormalities and a de novo deletion of NRXN1. <i>Journal of Medical Genetics</i> , 2007, 45, 239-243.	3.2	123
5	ACTB Loss-of-Function Mutations Result in a Pleiotropic Developmental Disorder. <i>American Journal of Human Genetics</i> , 2017, 101, 1021-1033.	6.2	83
6	Novel deletions of 14q11.2 associated with developmental delay, cognitive impairment and similar minor anomalies in three children. <i>Journal of Medical Genetics</i> , 2007, 44, 556-561.	3.2	68
7	Epigenetic Impacts on Neurodevelopment: Pathophysiological Mechanisms and Genetic Modes of Action. <i>Pediatric Research</i> , 2011, 69, 92R-100R.	2.3	62
8	Osteopoikilosis, short stature and mental retardation as key features of a new microdeletion syndrome on 12q14. <i>Journal of Medical Genetics</i> , 2007, 44, 264-268.	3.2	58
9	Duplications of the critical Rubinstein-Taybi deletion region on chromosome 16p13.3 cause a novel recognisable syndrome. <i>Journal of Medical Genetics</i> , 2010, 47, 155-161.	3.2	47
10	The impact of array genomic hybridization on mental retardation research: a review of current technologies and their clinical utility. <i>Clinical Genetics</i> , 2007, 72, 271-287.	2.0	43
11	MN1 C-terminal truncation syndrome is a novel neurodevelopmental and craniofacial disorder with partial rhombencephalosynapsis. <i>Brain</i> , 2020, 143, 55-68.	7.6	38
12	Single exon-resolution targeted chromosomal microarray analysis of known and candidate intellectual disability genes. <i>European Journal of Human Genetics</i> , 2014, 22, 792-800.	2.8	35
13	A distinct neurodevelopmental syndrome with intellectual disability, autism spectrum disorder, characteristic facies, and macrocephaly is caused by defects in CHD8. <i>Journal of Human Genetics</i> , 2019, 64, 271-280.	2.3	35
14	Detection of pathogenic copy number variants in children with idiopathic intellectual disability using 500 K SNP array genomic hybridization. <i>BMC Genomics</i> , 2009, 10, 526.	2.8	30
15	A characteristic syndrome associated with microduplication of 8q12, inclusive of CHD7. <i>European Journal of Medical Genetics</i> , 2009, 52, 436-439.	1.3	21
16	Use of Affymetrix Arrays in the Diagnosis of Gene Copy Number Variation. <i>Current Protocols in Human Genetics</i> , 2015, 85, 8.13.1-8.13.13.	3.5	15
17	Comprehensive whole genome sequence analyses yields novel genetic and structural insights for Intellectual Disability. <i>BMC Genomics</i> , 2017, 18, 403.	2.8	15
18	A novel de novo 1.1 Mb duplication of 17q21.33 associated with cognitive impairment and other anomalies. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 1257-1262.	1.2	14

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19	Intragenic CNVs for epigenetic regulatory genes in intellectual disability: Survey identifies pathogenic and benign single exon changes. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 2916-2926.	1.2	14
20	De novo pathogenic <i>DNM1L</i> variant in a patient diagnosed with atypical hereditary sensory and autonomic neuropathy. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e00961.	1.2	12
21	Maternal Prenatal Exposures in Pregnancy and Autism Spectrum Disorder: An Insight into the Epigenetics of Drugs and Diet as Key Environmental Influences. <i>Advances in Neurobiology</i> , 2020, 24, 143-162.	1.8	10
22	Discovering a new part of the phenotypic spectrum of Coffin-Siris syndrome in a fetal cohort. <i>Genetics in Medicine</i> , 2022, 24, 1753-1760.	2.4	6
23	Life's history chronicle for a patient with the recently described chromosome 4q21 microdeletion syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 2606-2609.	1.2	5
24	Chromodomain helicase DNA-binding proteins and neurodevelopmental disorders. , 0, , .		5
25	Recent Major Transcriptomics and Epitranscriptomics Contributions toward Personalized and Precision Medicine. <i>Journal of Personalized Medicine</i> , 2022, 12, 199.	2.5	3
26	First Whole Transcriptome RNAseq on CHD8 Haploinsufficient Patient and Meta-Analyses Across Cellular Models Uncover Likely Key Pathophysiological Target Genes. <i>Cureus</i> , 2020, 12, e11571.	0.5	2
27	Understanding environmental epigenomics in autism spectrum disorder: an interview with Farah R Zahir. <i>Epigenomics</i> , 2021, 13, 1341-1345.	2.1	0
28	The Need for Precision Therapies as Determined by Genetic Signature for Cystic Fibrosis. <i>Journal of Personalized Medicine</i> , 2021, 11, 1353.	2.5	0