Heather Mason-Suares

List of Publications by Year in descending order

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516710 552781 1,579 28 16 26 citations g-index h-index papers 31 31 31 4231 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Combined X-linked familial exudative vitreoretinopathy and retinopathy of prematurity phenotype in an infant with mosaic turner syndrome with ring X chromosome. Ophthalmic Genetics, 2023, 44, 198-203.	1.2	O
2	RASopathies., 2021,, 389-398.		0
3	Considerations for whole exome sequencing unique to prenatal care. Human Genetics, 2020, 139, 1149-1159.	3.8	18
4	Association of Rare Pathogenic DNA Variants for Familial Hypercholesterolemia, Hereditary Breast and Ovarian Cancer Syndrome, and Lynch Syndrome With Disease Risk in Adults According to Family History. JAMA Network Open, 2020, 3, e203959.	5.9	75
5	An assessment of the role of vinculin loss of function variants in inherited cardiomyopathy. Human Mutation, 2020, 41, 1577-1587.	2.5	10
6	A Role for Chromosomal Microarray Testing in the Workup of Male Infertility. Journal of Molecular Diagnostics, 2020, 22, 1189-1198.	2.8	4
7	Expanding the Noonan spectrum/RASopathy NGS panel: Benefits of adding <i>NF1</i> and <i>SPRED1</i> Molecular Genetics & amp; Genomic Medicine, 2020, 8, e1180.	1.2	9
8	Rare Genetic Variants Associated With Sudden Cardiac Death in Adults. Journal of the American College of Cardiology, 2019, 74, 2623-2634.	2.8	27
9	Considerations for clinical curation, classification, and reporting of low-penetrance and low effect size variants associated with disease risk. Genetics in Medicine, 2019, 21, 2765-2773.	2.4	20
10	Polygenic Prediction of Weight and Obesity Trajectories from Birth to Adulthood. Cell, 2019, 177, 587-596.e9.	28.9	516
11	ClinGen's RASopathy Expert Panel consensus methods for variant interpretation. Genetics in Medicine, 2018, 20, 1334-1345.	2.4	126
12	Analysis of intragenic USH2A copy number variation unveils broad spectrum of unique and recurrent variants. European Journal of Medical Genetics, 2018, 61, 621-626.	1.3	9
13	When ultrasound anomalies are present: An estimation of the frequency of chromosome abnormalities not detected by cellâ€free DNA aneuploidy screens. Prenatal Diagnosis, 2018, 38, 250-257.	2.3	13
14	Early cancer diagnoses through BRCA1/2 screening of unselected adult biobank participants. Genetics in Medicine, 2018, 20, 554-558.	2.4	46
15	NGS testing for cardiomyopathy: Utility of adding RASopathy-associated genes. Human Mutation, 2018, 39, 954-958.	2.5	11
16	Exome Sequencing–Based Screening for <i>BRCA1/2</i> Expected Pathogenic Variants Among Adult Biobank Participants. JAMA Network Open, 2018, 1, e182140.	5.9	163
17	Assessing the gene–disease association of 19 genes with the RASopathies using the ClinGen gene curation framework. Human Mutation, 2018, 39, 1485-1493.	2.5	66
	A Model for Genome-First Care: Returning Secondary Genomic Findings to Participants and Their Healthcare Providers in a Large Research Cohort. American Journal of Human Genetics, 2018, 103,	6.2	130

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19	Juvenile myelomonocytic leukemia-associated variants are associated with neo-natal lethal Noonan syndrome. European Journal of Human Genetics, 2017, 25, 509-511.	2.8	18
20	The current state of clinical interpretation of sequence variants. Current Opinion in Genetics and Development, 2017, 42, 33-39.	3.3	77
21	Electronic health record phenotype in subjects with genetic variants associated with arrhythmogenic right ventricular cardiomyopathy: a study of 30,716 subjects with exome sequencing. Genetics in Medicine, 2017, 19, 1245-1252.	2.4	43
22	Elucidation of MRAS-mediated Noonan syndrome with cardiac hypertrophy. JCI Insight, 2017, 2, e91225.	5.0	66
23	Training the Future Leaders in Personalized Medicine. Journal of Personalized Medicine, 2016, 6, 1.	2.5	21
24	Detecting Copy Number Variation via Next Generation Technology. Current Genetic Medicine Reports, 2016, 4, 74-85.	1.9	20
25	Retrospective study of prenatal ultrasound findings in newborns with a Noonan spectrum disorder. Prenatal Diagnosis, 2016, 36, 418-423.	2.3	25
26	Targeted Droplet-Digital PCR as a Tool for Novel Deletion Discovery at the DFNB1 Locus. Human Mutation, 2016, 37, 119-126.	2.5	37
27	Polycomb silencing of the Drosophila 4E-BP gene regulates imaginal disc cell growth. Developmental Biology, 2013, 380, 111-124.	2.0	3
28	Density matters: comparison of array platforms for detection of copy-number variation and copy-neutral abnormalities. Genetics in Medicine, 2013, 15, 706-712.	2.4	24