

Sarah Bowdin

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

2,226
citations

516710

16
h-index

752698

20
g-index

21
all docs

21
docs citations

21
times ranked

4994
citing authors

#	ARTICLE	IF	CITATIONS
1	The Cardiac Genome Clinic: implementing genome sequencing in pediatric heart disease. <i>Genetics in Medicine</i> , 2020, 22, 1015-1024.	2.4	51
2	The Personal Genome Project Canada: findings from whole genome sequences of the inaugural 56 participants. <i>Cmaj</i> , 2018, 190, E126-E136.	2.0	57
3	Periodic reanalysis of whole-genome sequencing data enhances the diagnostic advantage over standard clinical genetic testing. <i>European Journal of Human Genetics</i> , 2018, 26, 740-744.	2.8	88
4	Genome-wide sequencing expands the phenotypic spectrum of EP300 variants. <i>European Journal of Medical Genetics</i> , 2018, 61, 125-129.	1.3	8
5	Improved diagnostic yield compared with targeted gene sequencing panels suggests a role for whole-genome sequencing as a first-tier genetic test. <i>Genetics in Medicine</i> , 2018, 20, 435-443.	2.4	404
6	Reappraisal of Reported Genes for Sudden Arrhythmic Death. <i>Circulation</i> , 2018, 138, 1195-1205.	1.6	271
7	Use of Clinical Exome Sequencing in Isolated Congenital Heart Disease. <i>Circulation: Cardiovascular Genetics</i> , 2017, 10, .	5.1	13
8	Care and cost consequences of pediatric whole genome sequencing compared to chromosome microarray. <i>European Journal of Human Genetics</i> , 2017, 25, 1303-1312.	2.8	32
9	Genome sequencing as a platform for pharmacogenetic genotyping: a pediatric cohort study. <i>Npj Genomic Medicine</i> , 2017, 2, 19.	3.8	41
10	Whole-genome sequencing expands diagnostic utility and improves clinical management in paediatric medicine. <i>Npj Genomic Medicine</i> , 2016, 1, .	3.8	295
11	Recommendations for the integration of genomics into clinical practice. <i>Genetics in Medicine</i> , 2016, 18, 1075-1084.	2.4	125
12	<i>FGFR3</i> associated craniosynostosis syndromes and gastrointestinal defects. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 3215-3221.	1.2	13
13	Exome sequencing identifies rare variants in multiple genes in atrioventricular septal defect. <i>Genetics in Medicine</i> , 2016, 18, 189-198.	2.4	39
14	MKS1 regulates ciliary INPP5E levels in Joubert syndrome. <i>Journal of Medical Genetics</i> , 2016, 53, 62-72.	3.2	48
15	MG-132...Diagnostic utility of whole genome sequencing in paediatric medicine. <i>Journal of Medical Genetics</i> , 2015, 52, A12.1-A12.	3.2	1
16	Novel MASP1 mutations are associated with an expanded phenotype in 3MC1 syndrome. <i>Orphanet Journal of Rare Diseases</i> , 2015, 10, 128.	2.7	46
17	MG-130...Utilising whole exome sequencing to identify causative variants in genetically heterogeneous disorders. <i>Journal of Medical Genetics</i> , 2015, 52, A11.2-A11.	3.2	0
18	Heterozygous mutations in <i>ERF</i> cause syndromic craniosynostosis with multiple suture involvement. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 2544-2547.	1.2	21

#	ARTICLE	IF	CITATIONS
19	Loeysâ€Dietz syndrome: a primer for diagnosis and management. <i>Genetics in Medicine</i> , 2014, 16, 576-587.	2.4	435
20	The Genome Clinic: A Multidisciplinary Approach to Assessing the Opportunities and Challenges of Integrating Genomic Analysis into Clinical Care. <i>Human Mutation</i> , 2014, 35, 513-519.	2.5	31
21	PhenoTips: Patient Phenotyping Software for Clinical and Research Use. <i>Human Mutation</i> , 2013, 34, 1057-1065.	2.5	207