

Eddie Ip

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4678091/publications.pdf>

Version: 2024-02-01

10
papers

392
citations

1162367

8
h-index

1372195

10
g-index

10
all docs

10
docs citations

10
times ranked

1009
citing authors

#	ARTICLE	IF	CITATIONS
1	NAD Deficiency, Congenital Malformations, and Niacin Supplementation. <i>New England Journal of Medicine</i> , 2017, 377, 544-552.	13.9	177
2	A Screening Approach to Identify Clinically Actionable Variants Causing Congenital Heart Disease in Exome Data. <i>Circulation Genomic and Precision Medicine</i> , 2018, 11, e001978.	1.6	65
3	Identification of clinically actionable variants from genome sequencing of families with congenital heart disease. <i>Genetics in Medicine</i> , 2019, 21, 1111-1120.	1.1	54
4	Functional genomics and gene-environment interaction highlight the complexity of congenital heart disease caused by Notch pathway variants. <i>Human Molecular Genetics</i> , 2020, 29, 566-579.	1.4	32
5	Genetic burden and associations with adverse neurodevelopment in neonates with congenital heart disease. <i>American Heart Journal</i> , 2018, 201, 33-39.	1.2	19
6	Heterozygous loss of <i>WBP11</i> function causes multiple congenital defects in humans and mice. <i>Human Molecular Genetics</i> , 2021, 29, 3662-3678.	1.4	14
7	VPOT: A Customizable Variant Prioritization Ordering Tool for Annotated Variants. <i>Genomics, Proteomics and Bioinformatics</i> , 2019, 17, 540-545.	3.0	10
8	Whole genome sequencing in transposition of the great arteries and associations with clinically relevant heart, brain and laterality genes. <i>American Heart Journal</i> , 2022, 244, 1-13.	1.2	10
9	Spliceogen: an integrative, scalable tool for the discovery of splice-altering variants. <i>Bioinformatics</i> , 2019, 35, 4405-4407.	1.8	7
10	CHDgene: A Curated Database for Congenital Heart Disease Genes. <i>Circulation Genomic and Precision Medicine</i> , 2022, 15, 101161CIRCGEN121003539.	1.6	4