

# Daniel Quiat

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4418577/publications.pdf>

Version: 2024-02-01

12  
papers

1,425  
citations

1306789

7  
h-index

1199166

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

2621  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Family of microRNAs Encoded by Myosin Genes Governs Myosin Expression and Muscle Performance. <i>Developmental Cell</i> , 2009, 17, 662-673.	3.1	865
2	MicroRNAs in cardiovascular disease: from pathogenesis to prevention and treatment. <i>Journal of Clinical Investigation</i> , 2013, 123, 11-18.	3.9	260
3	Concerted regulation of myofiber-specific gene expression and muscle performance by the transcriptional repressor Sox6. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 10196-10201.	3.3	122
4	Genomic frontiers in congenital heart disease. <i>Nature Reviews Cardiology</i> , 2022, 19, 26-42.	6.1	93
5	Association of Damaging Variants in Genes With Increased Cancer Risk Among Patients With Congenital Heart Disease. <i>JAMA Cardiology</i> , 2021, 6, 457.	3.0	34
6	Discordant clinical features of identical hypertrophic cardiomyopathy twins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	19
7	High-Throughput Screening of Kawasaki Disease Sera for Antiviral Antibodies. <i>Journal of Infectious Diseases</i> , 2020, 222, 1853-1857.	1.9	9
8	Genome-Wide De Novo Variants in Congenital Heart Disease Are Not Associated With Maternal Diabetes or Obesity. <i>Circulation Genomic and Precision Medicine</i> , 2022, 15, CIRCGEN121003500.	1.6	8
9	Neither cardiac mitochondrial DNA variation nor copy number contribute to congenital heart disease risk. <i>American Journal of Human Genetics</i> , 2022, 109, 961-966.	2.6	5
10	Population Prevalence of Premature Truncating Variants in Plakophilin-2 and Association With Arrhythmogenic Right Ventricular Cardiomyopathy: A UK Biobank Analysis. <i>Circulation Genomic and Precision Medicine</i> , 2022, 15, 101161CIRCGEN121003507.	1.6	5
11	An ancient founder mutation located between <i>ROBO1</i> and <i>ROBO2</i> is responsible for increased microtia risk in Amerindigenous populations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2203928119.	3.3	4
12	Left Ventricular Outflow Tract Gradient Is Associated With Coronary Artery Obstruction in Children With Williams-Beuren Syndrome. <i>Journal of Cardiothoracic and Vascular Anesthesia</i> , 2021, 35, 3677-3680.	0.6	1