

# John F Robinson

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

22  
papers

418  
citations

10  
h-index

20  
g-index

22  
ext. papers

537  
ext. citations

4.8  
avg, IF

2.92  
L-index

#	Paper	IF	Citations
22	Enrichment of loss-of-function and copy number variants in ventricular cardiomyopathy genes in atrial fibrillation. <i>Europace</i> , <b>2021</b> , 23, 844-850	3.9	2
21	Simplifying Detection of Copy-Number Variations in Maturity-Onset Diabetes of the Young. <i>Canadian Journal of Diabetes</i> , <b>2021</b> , 45, 71-77	2.1	
20	Association of apolipoprotein E variation with cognitive impairment across multiple neurodegenerative diagnoses. <i>Neurobiology of Aging</i> , <b>2021</b> , 105, 378.e1-378.e9	5.6	1
19	Six years of experience with LipidSeq: clinical and research learnings from a hybrid, targeted sequencing panel for dyslipidemias. <i>BMC Medical Genomics</i> , <b>2020</b> , 13, 23	3.7	23
18	Partial deletions: rare copy-number variants contributing towards severe hypertriglyceridemia. <i>Journal of Lipid Research</i> , <b>2019</b> , 60, 1953-1958	6.3	10
17	Genetic and epigenetic study of an Alzheimer's disease family with monozygotic triplets. <i>Brain</i> , <b>2019</b> , 142, 3375-3381	11.2	6
16	Genetic Variation in the Ontario Neurodegenerative Disease Research Initiative. <i>Canadian Journal of Neurological Sciences</i> , <b>2019</b> , 46, 491-498	1	5
15	Clinical Utility and Practical Considerations of a Coronary Artery Disease Genetic Risk Score. <i>CJC Open</i> , <b>2019</b> , 1, 69-75	2	3
14	Targeted sequencing reveals expanded genetic diversity of human transfer RNAs. <i>RNA Biology</i> , <b>2019</b> , 16, 1574-1585	4.8	9
13	Bioinformatic detection of copy number variation in HNF4A causing maturity onset diabetes of the young. <i>Clinical Genetics</i> , <b>2019</b> , 96, 376-377	4	2
12	Copy Number Variation in GCK in Patients With Maturity-Onset Diabetes of the Young. <i>Journal of Clinical Endocrinology and Metabolism</i> , <b>2019</b> , 104, 3428-3436	5.6	7
11	Ankyrin-B dysfunction predisposes to arrhythmogenic cardiomyopathy and is amenable to therapy. <i>Journal of Clinical Investigation</i> , <b>2019</b> , 129, 3171-3184	15.9	23
10	Targeted Next-generation Sequencing and Bioinformatics Pipeline to Evaluate Genetic Determinants of Constitutional Disease. <i>Journal of Visualized Experiments</i> , <b>2018</b> ,	1.6	14
9	Whole-Genome Duplication of PCSK9 as a Novel Genetic Mechanism for Severe Familial Hypercholesterolemia. <i>Canadian Journal of Cardiology</i> , <b>2018</b> , 34, 1316-1324	3.8	23
8	Polygenic determinants in extremes of high-density lipoprotein cholesterol. <i>Journal of Lipid Research</i> , <b>2017</b> , 58, 2162-2170	6.3	33
7	Polygenic Versus Monogenic Causes of Hypercholesterolemia Ascertained Clinically. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , <b>2016</b> , 36, 2439-2445	9.4	130
6	Proprietary Considerations in the Use of Cardiovascular Genetic Data. <i>Canadian Journal of Cardiology</i> , <b>2016</b> , 32, 1297-1299	3.8	0

5	Whole-genome sequencing in French Canadians from Quebec. <i>Human Genetics</i> , <b>2016</b> , 135, 1213-1221	6.3	13
4	The ONDRISeq panel: custom-designed next-generation sequencing of genes related to neurodegeneration. <i>Npj Genomic Medicine</i> , <b>2016</b> , 1, 16032	6.2	17
3	Exome sequencing identifies NFS1 deficiency in a novel Fe-S cluster disease, infantile mitochondrial complex II/III deficiency. <i>Molecular Genetics &amp; Genomic Medicine</i> , <b>2014</b> , 2, 73-80	2.3	41
2	A novel LIPE nonsense mutation found using exome sequencing in siblings with late-onset familial partial lipodystrophy. <i>Canadian Journal of Cardiology</i> , <b>2014</b> , 30, 1649-54	3.8	50
1	The Ontario Neurodegenerative Disease Research Initiative		6