

# Marc-Andr Legault

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/922142/marc-andre-legault-publications-by-year.pdf>

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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.

10  
papers

50  
citations

4  
h-index

7  
g-index

17  
ext. papers

80  
ext. citations

6.9  
avg, IF

1.37  
L-index

#	Paper	IF	Citations
10	Genetic meta-analysis of cancer diagnosis following statin use identifies new associations and implicates human leukocyte antigen (HLA) in women. <i>Pharmacogenomics Journal</i> , <b>2021</b> , 21, 446-457	3.5	1
9	Pharmacogenomics of the Efficacy and Safety of Colchicine in COLCOT. <i>Circulation Genomic and Precision Medicine</i> , <b>2021</b> , 14, e003183	5.2	2
8	A genetic model of ivabradine recapitulates results from randomized clinical trials. <i>PLoS ONE</i> , <b>2020</b> , 15, e0236193	3.7	1
7	A genetic model of ivabradine recapitulates results from randomized clinical trials <b>2020</b> , 15, e0236193		
6	A genetic model of ivabradine recapitulates results from randomized clinical trials <b>2020</b> , 15, e0236193		
5	A genetic model of ivabradine recapitulates results from randomized clinical trials <b>2020</b> , 15, e0236193		
4	A genetic model of ivabradine recapitulates results from randomized clinical trials <b>2020</b> , 15, e0236193		
3	genipe: an automated genome-wide imputation pipeline with automatic reporting and statistical tools. <i>Bioinformatics</i> , <b>2016</b> , 32, 3661-3663	7.2	10
2	Comparison of sequencing based CNV discovery methods using monozygotic twin quartets. <i>PLoS ONE</i> , <b>2015</b> , 10, e0122287	3.7	13
1	pyGenClean: efficient tool for genetic data clean up before association testing. <i>Bioinformatics</i> , <b>2013</b> , 29, 1704-5	7.2	10