

# R Thomas Lumbers

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6739188/publications.pdf>

Version: 2024-02-01

21  
papers

1,730  
citations

840776

11  
h-index

940533

16  
g-index

24  
all docs

24  
docs citations

24  
times ranked

3495  
citing authors

#	ARTICLE	IF	CITATIONS
1	A population-based study of 92 clinically recognized risk factors for heart failure: co-occurrence, prognosis and preventive potential. <i>European Journal of Heart Failure</i> , 2022, 24, 466-480.	7.1	14
2	Therapeutic Targets for Heart Failure Identified Using Proteomics and Mendelian Randomization. <i>Circulation</i> , 2022, 145, 1205-1217.	1.6	50
3	Genetic and environmental determinants of diastolic heart function. , 2022, 1, 361-371.		12
4	Integrating polygenic risk scores in the prediction of type 2 diabetes risk and subtypes in British Pakistanis and Bangladeshis: A population-based cohort study. <i>PLoS Medicine</i> , 2022, 19, e1003981.	8.4	24
5	Estimating the Effect of Reduced Attendance at Emergency Departments for Suspected Cardiac Conditions on Cardiac Mortality During the COVID-19 Pandemic. <i>Circulation: Cardiovascular Quality and Outcomes</i> , 2021, 14, e007085.	2.2	18
6	Transforming and evaluating electronic health record disease phenotyping algorithms using the OMOP common data model: a case study in heart failure. <i>JAMIA Open</i> , 2021, 4, ooab001.	2.0	18
7	Life-Time Covariation of Major Cardiovascular Diseases. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e002963.	3.6	5
8	Evidence-Based Assessment of Genes in Dilated Cardiomyopathy. <i>Circulation</i> , 2021, 144, 7-19.	1.6	213
9	Type 2 Diabetes, Metabolic Traits, and Risk of Heart Failure: A Mendelian Randomization Study. <i>Diabetes Care</i> , 2021, 44, 1699-1705.	8.6	18
10	The genomics of heart failure: design and rationale of the HERMES consortium. <i>ESC Heart Failure</i> , 2021, 8, 5531-5541.	3.1	11
11	Shared genetic pathways contribute to risk of hypertrophic and dilated cardiomyopathies with opposite directions of effect. <i>Nature Genetics</i> , 2021, 53, 128-134.	21.4	155
12	Genome-wide association and Mendelian randomisation analysis provide insights into the pathogenesis of heart failure. <i>Nature Communications</i> , 2020, 11, 163.	12.8	466
13	A genetic model of ivabradine recapitulates results from randomized clinical trials. <i>PLoS ONE</i> , 2020, 15, e0236193.	2.5	3
14	Genetic and functional insights into the fractal structure of the heart. <i>Nature</i> , 2020, 584, 589-594.	27.8	86
15	A genetic model of ivabradine recapitulates results from randomized clinical trials. , 2020, 15, e0236193.		0
16	A genetic model of ivabradine recapitulates results from randomized clinical trials. , 2020, 15, e0236193.		0
17	A genetic model of ivabradine recapitulates results from randomized clinical trials. , 2020, 15, e0236193.		0
18	A genetic model of ivabradine recapitulates results from randomized clinical trials. , 2020, 15, e0236193.		0

#	ARTICLE	IF	CITATIONS
19	Do beta-blockers and inhibitors of the renin-angiotensin aldosterone system improve outcomes in patients with heart failure and left ventricular ejection fraction >40%. Heart, 2019, 105, 1533-1535.	2.9	4
20	A chronological map of 308 physical and mental health conditions from 4 million individuals in the English National Health Service. The Lancet Digital Health, 2019, 1, e63-e77.	12.3	192
21	The druggable genome and support for target identification and validation in drug development. Science Translational Medicine, 2017, 9, .	12.4	437