Laura Andreasen

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/4661242/publications.pdf

Version: 2024-02-01

21	571	840776 11	713466 21
papers	citations	h-index	g-index

21	21	21	1231
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	New population-based exome data are questioning the pathogenicity of previously cardiomyopathy-associated genetic variants. European Journal of Human Genetics, 2013, 21, 918-928.	2.8	200
2	Rare truncating variants in the sarcomeric protein titin associate with familial and early-onset atrial fibrillation. Nature Communications, 2018, 9, 4316.	12.8	93
3	Very early-onset lone atrial fibrillation patients have a high prevalence of rare variants in genes previously associated with atrial fibrillation. Heart Rhythm, 2014, 11, 246-251.	0.7	54
4	Atrial fibrillationâ€"a complex polygenetic disease. European Journal of Human Genetics, 2021, 29, 1051-1060.	2.8	30
5	Genome-wide association study identifies 18 novel loci associated with left atrial volume and function. European Heart Journal, 2021, 42, 4523-4534.	2.2	30
6	Numerous Brugada syndromeâ€"associated genetic variants have no effect on J-point elevation, syncope susceptibility, malignant cardiac arrhythmia, and all-cause mortality. Genetics in Medicine, 2017, 19, 521-528.	2.4	26
7	Verification of threshold for image intensity ratio analyses of late gadolinium enhancement magnetic resonance imaging of left atrial fibrosis in 1.5T scans. International Journal of Cardiovascular Imaging, 2020, 36, 513-520.	1.5	17
8	Loss-of-Function Variants in Cytoskeletal Genes Are Associated with Early-Onset Atrial Fibrillation. Journal of Clinical Medicine, 2020, 9, 372.	2.4	14
9	Brugada syndrome risk loci seem protective against atrial fibrillation. European Journal of Human Genetics, 2014, 22, 1357-1361.	2.8	13
10	Genome-wide association study identifies locus at chromosome 2q32.1 associated with syncope and collapse. Cardiovascular Research, 2020, 116, 138-148.	3.8	13
11	Deep sequencing of atrial fibrillation patients with mitral valve regurgitation shows no evidence of mosaicism but reveals novel rare germline variants. Heart Rhythm, 2017, 14, 1531-1538.	0.7	12
12	Next-generation sequencing of AV nodal reentrant tachycardia patients identifies broad spectrum of variants in ion channel genes. European Journal of Human Genetics, 2018, 26, 660-668.	2.8	12
13	Early-onset atrial fibrillation patients show reduced left ventricular ejection fraction and increased atrial fibrosis. Scientific Reports, 2020, 10, 10039.	3.3	12
14	Genetic Aspects of Lone Atrial Fibrillation: What Do We Know?. Current Pharmaceutical Design, 2014, 21, 667-678.	1.9	12
15	Association of Variants Near the Bradykinin Receptor B2 Gene With Angioedema in Patients Taking ACEÂInhibitors. Journal of the American College of Cardiology, 2021, 78, 696-709.	2.8	10
16	Genetic Modifier of the QTc Interval Associated With Early-Onset Atrial Fibrillation. Canadian Journal of Cardiology, 2013, 29, 1234-1240.	1.7	9
17	Cardiac magnetic resonance systematically overestimates mitral regurgitations by the indirect method. Open Heart, 2020, 7, e001323.	2.3	5
18	Polygenic risk score for ACE-inhibitor-associated cough based on the discovery of new genetic loci. European Heart Journal, 2022, 43, 4707-4718.	2.2	5

#	Article	IF	Citations
19	Brugada Syndrome-Associated Genetic Loci Are Associated With J-Point Elevation and an Increased Risk of Cardiac Arrest. Frontiers in Physiology, 2018, 9, 894.	2.8	2
20	Mendelian randomizationâ€"a powerful tool to study the causal effects of atrial fibrillation on loss of brain volume. BMC Medicine, 2021, 19, 70.	5.5	1
21	Clinical Implications of SCN10A Loss-of-Function Variants in 169â€‰610 Exomes Representing the General Population. Circulation Genomic and Precision Medicine, 2022, 15, CIRCGEN121003574.	3.6	1