Lena Refsgaard

List of Publications by Year
in descending order

19 papers	1,474 citations	758635 12 h-index	794141 19 g-index
19 all docs	19 docs citations	19 times ranked	3266 citing authors

#	Article	IF	Citations
1	Whole-Exome Sequencing Implicates Neuronal Calcium Channel with Familial Atrial Fibrillation. Frontiers in Genetics, 2022, 13, 806429.	1.1	1
2	Early-onset atrial fibrillation patients show reduced left ventricular ejection fraction and increased atrial fibrosis. Scientific Reports, 2020, 10, 10039.	1.6	12
3	A Novel Loss-of-Function Variant in the Chloride Ion Channel Gene Clcn2 Associates with Atrial Fibrillation. Scientific Reports, 2020, 10, 1453.	1.6	10
4	Genome-wide Study of Atrial Fibrillation Identifies Seven Risk Loci and Highlights Biological Pathways and Regulatory Elements Involved in Cardiac Development. American Journal of Human Genetics, 2018, 102, 103-115.	2.6	86
5	Rare truncating variants in the sarcomeric protein titin associate with familial and early-onset atrial fibrillation. Nature Communications, 2018, 9, 4316.	5.8	93
6	Multi-ethnic genome-wide association study for atrial fibrillation. Nature Genetics, 2018, 50, 1225-1233.	9.4	552
7	Generation of induced pluripotent stem cells (iPSC) from an atrial fibrillation patient carrying a PITX2 p.M200V mutation. Stem Cell Research, 2017, 24, 8-11.	0.3	7
8	Generation of induced pluripotent stem cells (iPSC) from an atrial fibrillation patient carrying a KCNA5 p.D322H mutation. Stem Cell Research, 2017, 24, 29-32.	0.3	3
9	I_{Ks} Gainâ€•and Lossâ€ofâ€Function in Earlyâ€Onset Lone Atrial Fibrillation. Journal of Cardiovascular Electrophysiology, 2015, 26, 715-723.	0.8	28
10	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.	0.3	54
11	Brugada syndrome risk loci seem protective against atrial fibrillation. European Journal of Human Genetics, 2014, 22, 1357-1361.	1.4	13
12	New population-based exome data are questioning the pathogenicity of previously cardiomyopathy-associated genetic variants. European Journal of Human Genetics, 2013, 21, 918-928.	1.4	200
13	Mutations in Genes Encoding Cardiac Ion Channels Previously Associated With Sudden Infant Death Syndrome (SIDS) Are Present With High Frequency in New Exome Data. Canadian Journal of Cardiology, 2013, 29, 1104-1109.	0.8	58
14	High prevalence of genetic variants previously associated with Brugada syndrome in new exome data. Clinical Genetics, 2013, 84, 489-495.	1.0	102
15	A novel KCND3 gain-of-function mutation associated with early-onset of persistent lone atrial fibrillation. Cardiovascular Research, 2013, 98, 488-495.	1.8	104
16	Screening of the Ito Regulatory Subunit Klf15 in Patients with Early-Onset Lone Atrial Fibrillation. Frontiers in Genetics, 2013, 4, 88.	1.1	5
17	High prevalence of genetic variants previously associated with LQT syndrome in new exome data. European Journal of Human Genetics, 2012, 20, 905-908.	1.4	121
18	Mutation analysis of the candidate genes SCN1B-4B, FHL1, and LMNA in patients with arrhythmogenic right ventricular cardiomyopathy. Applied & Translational Genomics, 2012, 1, 44-46.	2.1	3

#	Article	IF	Citations
19	Sodium Current and Potassium Transient Outward Current Genes in Brugada Syndrome: Screening and Bioinformatics. Canadian Journal of Cardiology, 2012, 28, 196-200.	0.8	22