Javad Jabbari

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/10826609/publications.pdf

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19	999	687363 13	794594 19
papers	citations	h-index	g-index

19	19	19	1910
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Exome data clouds the pathogenicity of genetic variants in Pulmonary Arterial Hypertension. Molecular Genetics & Genomic Medicine, 2018, 6, 835-844.	1.2	3
2	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
3	Stability of Circulating Blood-Based MicroRNAs â€“ Pre-Analytic Methodological Considerations. PLoS ONE, 2017, 12, e0167969.	2.5	247
4	A Common Variant in SCN5A and the Risk of Ventricular Fibrillation Caused by First ST-Segment Elevation Myocardial Infarction. PLoS ONE, 2017, 12, e0170193.	2.5	17
5	Association of common genetic variants related to atrial fibrillation and the risk of ventricular fibrillation in the setting of first ST-elevation myocardial infarction. BMC Medical Genetics, 2017, 18, 138.	2.1	2
6	The pathogenicity of genetic variants previously associated with left ventricular nonâ€compaction. Molecular Genetics & Genomic Medicine, 2016, 4, 135-142.	1.2	11
7	Role of common and rare variants in SCN10A: results from the Brugada syndrome QRS locus gene discovery collaborative study. Cardiovascular Research, 2015, 106, 520-529.	3.8	108
8	Common and Rare Variants in SCN10A Modulate the Risk of Atrial Fibrillation. Circulation: Cardiovascular Genetics, 2015, 8, 64-73.	5.1	50
9	Incidence and Risk Factors of Ventricular Fibrillation Before Primary Angioplasty in Patients With First STâ€Elevation Myocardial Infarction: A Nationwide Study in Denmark. Journal of the American Heart Association, 2015, 4, e001399.	3.7	91
10	Common and Rare Variants in SCN10A Modulate the Risk of Atrial Fibrillation. Circulation: Cardiovascular Genetics, 2015, 8, 64-73.	5.1	59
11	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
12	New population-based exome data question the pathogenicity of some genetic variants previously associated with Marfan syndrome. BMC Genetics, 2014, 15, 74.	2.7	15
13	Brugada syndrome risk loci seem protective against atrial fibrillation. European Journal of Human Genetics, 2014, 22, 1357-1361.	2.8	13
14	Rare Variants in GJA5 Are Associated With Early-Onset Lone Atrial Fibrillation. Canadian Journal of Cardiology, 2013, 29, 111-116.	1.7	46
15	New Exome Data Question the Pathogenicity of Genetic Variants Previously Associated With Catecholaminergic Polymorphic Ventricular Tachycardia. Circulation: Cardiovascular Genetics, 2013, 6, 481-489.	5.1	74
16	Genetic Loci on Chromosomes 4q25, 7p31, and 12p12 Are Associated With Onset of Lone Atrial Fibrillation Before the Age of 40 Years. Canadian Journal of Cardiology, 2012, 28, 191-195.	1.7	50
17	Mutations in the potassium channel subunit KCNE1 are associated with early-onset familial atrial fibrillation. BMC Medical Genetics, 2012, 13, 24.	2.1	79
18	Common Polymorphisms in KNCJ5 Are Associated with Early-Onset Lone Atrial Fibrillation in Caucasians. Cardiology, 2011, 118, 116-120.	1.4	24

#	Article	IF	Citations
19	Screening of KCNN3 in patients with early-onset lone atrial fibrillation. Europace, 2011, 13, 963-967.	1.7	44