Zongzhe Li

List of Publications by Year in descending order

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759233 794594 29 404 12 19 citations h-index g-index papers 31 31 31 716 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Rare Variants and Polymorphisms of FBN1 Gene May Increase the Risk of Non-Syndromic Aortic Dissection. Frontiers in Genetics, 2022, 13, 778806.	2.3	2
2	Effects of Shuanghuanglian oral liquids on patients with COVID-19: a randomized, open-label, parallel-controlled, multicenter clinical trial. Frontiers of Medicine, 2021, 15, 704-717.	3.4	33
3	A nonsynonymous polymorphism (rs117179004, T392M) of hyaluronidase 1 (HYAL1) is associated with increased risk of idiopathic pulmonary fibrosis in Southern Han Chinese. Journal of Clinical Laboratory Analysis, 2021, 35, e23782.	2.1	O
4	Targeted resequencing showing novel common and rare genetic variants increases the risk of asthma in the Chinese Han population. Journal of Clinical Laboratory Analysis, 2021, 35, e23813.	2.1	3
5	Identification of COL3A1 variants associated with sporadic thoracic aortic dissection: a case-control study. Frontiers of Medicine, 2021, 15, 438-447.	3.4	3
6	<i>COL5A1</i> Variants Cause Aortic Dissection by Activating TGFâ€Î²â€Signaling Pathway. Journal of the American Heart Association, 2021, 10, e019276.	3.7	6
7	RBM20 Is a Candidate Gene for Hypertrophic Cardiomyopathy. Canadian Journal of Cardiology, 2021, 37, 1751-1759.	1.7	10
8	Aortic root aortopathy in bicuspid aortic valve associated with high genetic risk. BMC Cardiovascular Disorders, 2021, 21, 413.	1.7	6
9	Case Report: A Novel LAMP2 Splice-Altering Mutation Causes Cardiac-Only Danon Disease. Frontiers in Cardiovascular Medicine, 2021, 8, 763240.	2.4	3
10	Aortic Coarctation Associated With Hypertrophic Cardiomyopathy in a Woman With Hypertension and Syncope: A Case Report With 8-Year Follow-Up. Frontiers in Cardiovascular Medicine, 2021, 8, 818884.	2.4	0
11	Genetic arrhythmias complicating patients with dilated cardiomyopathy. Heart Rhythm, 2020, 17, 305-312.	0.7	11
12	Distal myopathy induced arrhythmogenic right ventricular cardiomyopathy in a pedigree carrying novel DSG2 null variant. International Journal of Cardiology, 2020, 298, 25-31.	1.7	5
13	An isolated hypogonadotropic hypogonadism male with a novel de novo FGFR1 mutation fathered a normal son evidenced by prenatal genetic diagnosis. Andrologia, 2020, 52, e13821.	2.1	1
14	A Novel Gain-of-Function <i>KCND3</i> Variant Associated with Brugada Syndrome. Cardiology, 2020, 145, 623-632.	1.4	4
15	MYH7B variants cause hypertrophic cardiomyopathy by activating the CaMK-signaling pathway. Science China Life Sciences, 2020, 63, 1347-1362.	4.9	17
16	Whole-exome sequencing identifies a de novo PDE3A variant causing autosomal dominant hypertension with brachydactyly type E syndrome: a case report. BMC Medical Genetics, 2020, 21, 144.	2.1	3
17	Association of TSR1 Variants and Spontaneous Coronary Artery Dissection. Journal of the American College of Cardiology, 2019, 74, 167-176.	2.8	35
18	The coexistence of a novel WNK1 variant and a copy number variation causes hereditary sensory and autonomic neuropathy type IIA. BMC Medical Genetics, 2019, 20, 91.	2.1	1

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#	Article	lF	CITATION
19	A PLN nonsense variant causes severe dilated cardiomyopathy in a novel autosomal recessive inheritance mode. International Journal of Cardiology, 2019, 279, 122-125.	1.7	10
20	A targeted sequencing approach to find novel pathogenic genes associated with sporadic aortic dissection. Science China Life Sciences, 2018, 61, 1545-1553.	4.9	21
21	ADRB2 polymorphism Arg16Gly modifies the natural outcome of heart failure and dictates therapeutic response to \hat{l}^2 -blockers in patients with heart failure. Cell Discovery, 2018, 4, 57.	6.7	26
22	Mutation profiles and clinical characteristics of Chinese males with isolated hypogonadotropic hypogonadism. Fertility and Sterility, 2018, 110, 486-495.e5.	1.0	36
23	Targeted resequencing reveals genetic risks in patients with sporadic idiopathic pulmonary fibrosis. Human Mutation, 2018, 39, 1238-1245.	2.5	29
24	Variants of genes encoding collagens and matrix metalloproteinase system increased the risk of aortic dissection. Science China Life Sciences, 2017, 60, 57-65.	4.9	18
25	A novel nonsense mutation in LMNA gene identified by Exome Sequencing in an atrial fibrillation family. European Journal of Medical Genetics, 2016, 59, 396-400.	1.3	16
26	Next-generation sequencing identifies novel mutations in the FBN1 gene for two Chinese families with Marfan syndrome. Molecular Medicine Reports, 2016, 14, 151-158.	2.4	6
27	A Functional Variant in APOA5/A4/C3/A1 Gene Cluster Contributes to Elevated Triglycerides and Severity of CAD by Interfering With MicroRNA 3201 Binding Efficiency. Journal of the American College of Cardiology, 2014, 64, 267-277.	2.8	55
28	Rapid molecular genetic diagnosis of hypertrophic cardiomyopathy by semiconductor sequencing. Journal of Translational Medicine, 2014, 12, 173.	4.4	21
29	Novel URAT1 mutations caused acute renal failure after exercise in two Chinese families with renal hypouricemia. Gene, 2013, 512, 97-101.	2.2	22