

# Zongzhe Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3558995/publications.pdf>

Version: 2024-02-01

29  
papers

404  
citations

759233

12  
h-index

794594

19  
g-index

31  
all docs

31  
docs citations

31  
times ranked

716  
citing authors

#	ARTICLE	IF	CITATIONS
1	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. <i>Journal of the American College of Cardiology</i> , 2014, 64, 267-277.	2.8	55
2	Mutation profiles and clinical characteristics of Chinese males with isolated hypogonadotropic hypogonadism. <i>Fertility and Sterility</i> , 2018, 110, 486-495.e5.	1.0	36
3	Association of TSR1 Variants and Spontaneous Coronary Artery Dissection. <i>Journal of the American College of Cardiology</i> , 2019, 74, 167-176.	2.8	35
4	Effects of Shuanghuanglian oral liquids on patients with COVID-19: a randomized, open-label, parallel-controlled, multicenter clinical trial. <i>Frontiers of Medicine</i> , 2021, 15, 704-717.	3.4	33
5	Targeted resequencing reveals genetic risks in patients with sporadic idiopathic pulmonary fibrosis. <i>Human Mutation</i> , 2018, 39, 1238-1245.	2.5	29
6	ADRB2 polymorphism Arg16Gly modifies the natural outcome of heart failure and dictates therapeutic response to $\beta$ -blockers in patients with heart failure. <i>Cell Discovery</i> , 2018, 4, 57.	6.7	26
7	Novel URAT1 mutations caused acute renal failure after exercise in two Chinese families with renal hypouricemia. <i>Gene</i> , 2013, 512, 97-101.	2.2	22
8	Rapid molecular genetic diagnosis of hypertrophic cardiomyopathy by semiconductor sequencing. <i>Journal of Translational Medicine</i> , 2014, 12, 173.	4.4	21
9	A targeted sequencing approach to find novel pathogenic genes associated with sporadic aortic dissection. <i>Science China Life Sciences</i> , 2018, 61, 1545-1553.	4.9	21
10	Variants of genes encoding collagens and matrix metalloproteinase system increased the risk of aortic dissection. <i>Science China Life Sciences</i> , 2017, 60, 57-65.	4.9	18
11	MYH7B variants cause hypertrophic cardiomyopathy by activating the CaMK-signaling pathway. <i>Science China Life Sciences</i> , 2020, 63, 1347-1362.	4.9	17
12	A novel nonsense mutation in LMNA gene identified by Exome Sequencing in an atrial fibrillation family. <i>European Journal of Medical Genetics</i> , 2016, 59, 396-400.	1.3	16
13	Genetic arrhythmias complicating patients with dilated cardiomyopathy. <i>Heart Rhythm</i> , 2020, 17, 305-312.	0.7	11
14	A PLN nonsense variant causes severe dilated cardiomyopathy in a novel autosomal recessive inheritance mode. <i>International Journal of Cardiology</i> , 2019, 279, 122-125.	1.7	10
15	RBM20 Is a Candidate Gene for Hypertrophic Cardiomyopathy. <i>Canadian Journal of Cardiology</i> , 2021, 37, 1751-1759.	1.7	10
16	Next-generation sequencing identifies novel mutations in the FBN1 gene for two Chinese families with Marfan syndrome. <i>Molecular Medicine Reports</i> , 2016, 14, 151-158.	2.4	6
17	<i>COL5A1</i> Variants Cause Aortic Dissection by Activating TGF $\beta$ Signaling Pathway. <i>Journal of the American Heart Association</i> , 2021, 10, e019276.	3.7	6
18	Aortic root aortopathy in bicuspid aortic valve associated with high genetic risk. <i>BMC Cardiovascular Disorders</i> , 2021, 21, 413.	1.7	6

#	ARTICLE	IF	CITATIONS
19	Distal myopathy induced arrhythmogenic right ventricular cardiomyopathy in a pedigree carrying novel DSG2 null variant. <i>International Journal of Cardiology</i> , 2020, 298, 25-31.	1.7	5
20	A Novel Gain-of-Function <i>KCNQ3</i> Variant Associated with Brugada Syndrome. <i>Cardiology</i> , 2020, 145, 623-632.	1.4	4
21	Whole-exome sequencing identifies a de novo PDE3A variant causing autosomal dominant hypertension with brachydactyly type E syndrome: a case report. <i>BMC Medical Genetics</i> , 2020, 21, 144.	2.1	3
22	Targeted resequencing showing novel common and rare genetic variants increases the risk of asthma in the Chinese Han population. <i>Journal of Clinical Laboratory Analysis</i> , 2021, 35, e23813.	2.1	3
23	Identification of COL3A1 variants associated with sporadic thoracic aortic dissection: a case-control study. <i>Frontiers of Medicine</i> , 2021, 15, 438-447.	3.4	3
24	Case Report: A Novel LAMP2 Splice-Altering Mutation Causes Cardiac-Only Danon Disease. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 763240.	2.4	3
25	Rare Variants and Polymorphisms of FBN1 Gene May Increase the Risk of Non-Syndromic Aortic Dissection. <i>Frontiers in Genetics</i> , 2022, 13, 778806.	2.3	2
26	The coexistence of a novel WNK1 variant and a copy number variation causes hereditary sensory and autonomic neuropathy type IIA. <i>BMC Medical Genetics</i> , 2019, 20, 91.	2.1	1
27	An isolated hypogonadotropic hypogonadism male with a novel de novo FGFR1 mutation fathered a normal son evidenced by prenatal genetic diagnosis. <i>Andrologia</i> , 2020, 52, e13821.	2.1	1
28	A nonsynonymous polymorphism (rs117179004, T392M) of hyaluronidase 1 (HYAL1) is associated with increased risk of idiopathic pulmonary fibrosis in Southern Han Chinese. <i>Journal of Clinical Laboratory Analysis</i> , 2021, 35, e23782.	2.1	0
29	Aortic Coarctation Associated With Hypertrophic Cardiomyopathy in a Woman With Hypertension and Syncope: A Case Report With 8-Year Follow-Up. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 818884.	2.4	0