

Lingyao Zeng

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

4,501
citations

567281

15
h-index

642732

23
g-index

25
all docs

25
docs citations

25
times ranked

9195
citing authors

#	ARTICLE	IF	CITATIONS
1	J-shaped association between circulating apoC-III and cardiovascular mortality. <i>European Journal of Preventive Cardiology</i> , 2022, 29, e68-e71.	1.8	2
2	Genetically determined intelligence and coronary artery disease risk. <i>Clinical Research in Cardiology</i> , 2021, 110, 211-219.	3.3	19
3	Common APOC3 variants are associated with circulating ApoC-III and VLDL cholesterol but not with total apolipoprotein B and coronary artery disease. <i>Atherosclerosis</i> , 2020, 311, 84-90.	0.8	9
4	Mendelian randomization analysis does not support causal associations of birth weight with hypertension risk and blood pressure in adulthood. <i>European Journal of Epidemiology</i> , 2020, 35, 685-697.	5.7	9
5	Genetics of educational attainment and coronary risk in Mendelian randomization studies. <i>European Heart Journal</i> , 2020, 41, 894-895.	2.2	5
6	Genetically modulated educational attainment and coronary disease risk. <i>European Heart Journal</i> , 2019, 40, 2413-2420.	2.2	32
7	Genetic Risk Score for Coronary Disease Identifies Predispositions to Cardiovascular and Noncardiovascular Diseases. <i>Journal of the American College of Cardiology</i> , 2019, 73, 2932-2942.	2.8	58
8	Contribution of Gene Regulatory Networks to Heritability of Coronary Artery Disease. <i>Journal of the American College of Cardiology</i> , 2019, 73, 2946-2957.	2.8	45
9	Effects of the coronary artery disease associated LPA and 9p21 loci on risk of aortic valve stenosis. <i>International Journal of Cardiology</i> , 2019, 276, 212-217.	1.7	9
10	Association of the PHACTR1/EDN1 Genetic Locus With Spontaneous Coronary Artery Dissection. <i>Journal of the American College of Cardiology</i> , 2019, 73, 58-66.	2.8	147
11	Multi-ethnic genome-wide association study for atrial fibrillation. <i>Nature Genetics</i> , 2018, 50, 1225-1233.	21.4	552
12	Loss of Cardioprotective Effects at the <i>ADAMTS7</i> Locus as a Result of Gene-Smoking Interactions. <i>Circulation</i> , 2017, 135, 2336-2353.	1.6	51
13	Applications and Limitations of Mouse Models for Understanding Human Atherosclerosis. <i>Cell Metabolism</i> , 2017, 25, 248-261.	16.2	161
14	Genetic and functional interaction of the coronary artery disease risk gene <i>ADAMTS7</i> with LDL-cholesterol. <i>Atherosclerosis</i> , 2017, 263, e34.	0.8	0
15	Association analyses based on false discovery rate implicate new loci for coronary artery disease. <i>Nature Genetics</i> , 2017, 49, 1385-1391.	21.4	571
16	Rheumatoid Arthritis and Coronary Artery Disease: Genetic Analyses Do Not Support a Causal Relation. <i>Journal of Rheumatology</i> , 2017, 44, 4-10.	2.0	9
17	Genomic correlates of glatiramer acetate adverse cardiovascular effects lead to a novel locus mediating coronary risk. <i>PLoS ONE</i> , 2017, 12, e0182999.	2.5	5
18	Coding Variation in <i>ANGPTL4</i> , <i>LPL</i> and <i>SVEP1</i> and the Risk of Coronary Disease. <i>New England Journal of Medicine</i> , 2016, 374, 1134-1144.	27.0	427

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19	Genetics links between transforming growth factor \hat{I}^2 pathway and coronary disease. <i>Atherosclerosis</i> , 2016, 253, 237-246.	0.8	21
20	No Association of Coronary Artery Disease with X-Chromosomal Variants in Comprehensive International Meta-Analysis. <i>Scientific Reports</i> , 2016, 6, 35278.	3.3	25
21	Identifying Novel Gene Variants in Coronary Artery Disease and Shared Genes With Several Cardiovascular Risk Factors. <i>Circulation Research</i> , 2016, 118, 83-94.	4.5	52
22	Genetically Determined Height and Coronary Artery Disease. <i>New England Journal of Medicine</i> , 2015, 372, 1608-1618.	27.0	220
23	A comprehensive 1000 Genomesâ€‘based genome-wide association meta-analysis of coronary artery disease. <i>Nature Genetics</i> , 2015, 47, 1121-1130.	21.4	2,054
24	SyStemCell: A Database Populated with Multiple Levels of Experimental Data from Stem Cell Differentiation Research. <i>PLoS ONE</i> , 2012, 7, e35230.	2.5	13