Aniruddh Pradip Patel

List of Publications by Year
in descending order

21 papers	3,290 citations	777949 13 h-index	843174 20 g-index
23 all docs	23 docs citations	23 times ranked	8689 citing authors

#	Article	IF	Citations
1	Response by Patel and Khera to Letter Regarding Article, â€œQuantifying and Understanding the Higher Risk of Atherosclerotic Cardiovascular Disease Among South Asian Individuals: Results From the UK Biobank Prospective Cohort Studyâ€• Circulation, 2022, 145, e147-e148.	1.6	0
2	Association of the Interaction Between Familial Hypercholesterolemia Variants and Adherence to a Healthy Lifestyle With Risk of Coronary Artery Disease. JAMA Network Open, 2022, 5, e222687.	2.8	17
3	Association of Pathogenic DNA Variants Predisposing to Cardiomyopathy With Cardiovascular Disease Outcomes and All-Cause Mortality. JAMA Cardiology, 2022, 7, 723.	3.0	15
4	Lp(a) (Lipoprotein[a]) Concentrations and Incident Atherosclerotic Cardiovascular Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2021, 41, 465-474.	1.1	104
5	Genetic Predictor to Identify Individuals With High Lipoprotein(a) Concentrations. Circulation Genomic and Precision Medicine, 2021, 14, e003182.	1.6	10
6	Association of premature menopause with incident pulmonary hypertension: A cohort study. PLoS ONE, 2021, 16, e0247398.	1.1	8
7	Lipoprotein(a) and Coronary Artery Disease Risk Without a Family History of Heart Disease. Journal of the American Heart Association, 2021, 10, e017470.	1.6	10
8	Quantifying and Understanding the Higher Risk of Atherosclerotic Cardiovascular Disease Among South Asian Individuals. Circulation, 2021, 144, 410-422.	1.6	72
9	Selection of 51 predictors from 13,782 candidate multimodal features using machine learning improves coronary artery disease prediction. Patterns, 2021, 2, 100364.	3.1	18
10	Validation of a Genome-Wide PolygenicÂScore for Coronary ArteryÂDisease inÂSouth Asians. Journal of the American College of Cardiology, 2020, 76, 703-714.	1.2	76
11	Polygenic background modifies penetrance of monogenic variants for tier 1 genomic conditions. Nature Communications, 2020, 11, 3635.	5.8	277
12	Association of Rare Pathogenic DNA Variants for Familial Hypercholesterolemia, Hereditary Breast and Ovarian Cancer Syndrome, and Lynch Syndrome With Disease Risk in Adults According to Family History. JAMA Network Open, 2020, 3, e203959.	2.8	75
13	Race, socioeconomic deprivation, and hospitalization for COVID-19 in English participants of a national biobank. International Journal for Equity in Health, 2020, 19, 114.	1.5	101
14	Completing the genetic spectrum influencing coronary artery disease: from germline to somatic variation. Cardiovascular Research, 2019, 115, 830-843.	1.8	14
15	A New Murine Model of Clonal Hematopoiesis Investigates JAK2V617F inÂHeartÂFailure. JACC Basic To Translational Science, 2019, 4, 698-700.	1.9	2
16	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. Nature Genetics, 2018, 50, 26-41.	9.4	286
17	Rare and low-frequency coding variants alter human adult height. Nature, 2017, 542, 186-190.	13.7	544
18	Exome-wide association study of plasma lipids in >300,000 individuals. Nature Genetics, 2017, 49, 1758-1766.	9.4	470

#	Article	IF	Citations
19	Targeted exonic sequencing of GWAS loci in the high extremes of the plasma lipids distribution. Atherosclerosis, 2016, 250, 63-68.	0.4	11
20	Association of Low-Frequency and Rare Coding-Sequence Variants with Blood Lipids and Coronary Heart Disease in 56,000 Whites and Blacks. American Journal of Human Genetics, 2014, 94, 223-232.	2.6	287
21	K ⁺ Channel Mutations in Adrenal Aldosterone-Producing Adenomas and Hereditary Hypertension. Science, 2011, 331, 768-772.	6.0	866