

Shoa L Clarke

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5550314/publications.pdf>

Version: 2024-02-01

24
papers

4,770
citations

687363

13
h-index

677142

22
g-index

31
all docs

31
docs citations

31
times ranked

12745
citing authors

#	ARTICLE	IF	CITATIONS
1	GREAT improves functional interpretation of cis-regulatory regions. <i>Nature Biotechnology</i> , 2010, 28, 495-501.	17.5	3,789
2	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021, 600, 675-679.	27.8	353
3	Coding exons function as tissue-specific enhancers of nearby genes. <i>Genome Research</i> , 2012, 22, 1059-1068.	5.5	202
4	Human Developmental Enhancers Conserved between Deuterostomes and Protostomes. <i>PLoS Genetics</i> , 2012, 8, e1002852.	3.5	55
5	Validation of an Integrated Risk Tool, Including Polygenic Risk Score, for Atherosclerotic Cardiovascular Disease in Multiple Ethnicities and Ancestries. <i>American Journal of Cardiology</i> , 2021, 148, 157-164.	1.6	48
6	Mendelian randomization supports bidirectional causality between telomere length and clonal hematopoiesis of indeterminate potential. <i>Science Advances</i> , 2022, 8, eabl6579.	10.3	36
7	The Enhancer Landscape during Early Neocortical Development Reveals Patterns of Dense Regulation and Co-option. <i>PLoS Genetics</i> , 2013, 9, e1003728.	3.5	33
8	Control of pelvic girdle development by genes of the Pbx family and <i>Emx2</i> . <i>Developmental Dynamics</i> , 2011, 240, 1173-1189.	1.8	32
9	PRISM offers a comprehensive genomic approach to transcription factor function prediction. <i>Genome Research</i> , 2013, 23, 889-904.	5.5	32
10	<i>ZEB2</i> Shapes the Epigenetic Landscape of Atherosclerosis. <i>Circulation</i> , 2022, 145, 469-485.	1.6	31
11	Genome-Wide Association Studies of Coronary Artery Disease: Recent Progress and Challenges Ahead. <i>Current Atherosclerosis Reports</i> , 2018, 20, 47.	4.8	24
12	Combining Clinical and Polygenic Risk Improves Stroke Prediction Among Individuals With Atrial Fibrillation. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003168.	3.6	24
13	Associations of Genetically Predicted Lp(a) (Lipoprotein [a]) Levels With Cardiovascular Traits in Individuals of European and African Ancestry. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003354.	3.6	21
14	Cardiorespiratory Fitness, Body Mass Index, and Markers of Insulin Resistance in Apparently Healthy Women and Men. <i>American Journal of Medicine</i> , 2020, 133, 825-830.e2.	1.5	14
15	The Propagation of Racial Disparities in Cardiovascular Genomics Research. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003178.	3.6	14
16	The need for polygenic score reporting standards in evidence-based practice: lipid genetics use case. <i>Current Opinion in Lipidology</i> , 2021, 32, 89-95.	2.7	10
17	Use of Polygenic Risk Scores for Coronary Heart Disease in Ancestrally Diverse Populations. <i>Current Cardiology Reports</i> , 2022, 24, 1169-1177.	2.9	10
18	Erosion of Conserved Binding Sites in Personal Genomes Points to Medical Histories. <i>PLoS Computational Biology</i> , 2016, 12, e1004711.	3.2	7

#	ARTICLE	IF	CITATIONS
19	Using Mendelian randomisation to identify opportunities for type 2 diabetes prevention by repurposing medications used for lipid management. EBioMedicine, 2022, 80, 104038.	6.1	7
20	Coronary Artery Disease Risk of Familial Hypercholesterolemia Genetic Variants Independent of Clinically Observed Longitudinal Cholesterol Exposure. Circulation Genomic and Precision Medicine, 2022, 15, CIRCGEN121003501.	3.6	6
21	Time to Relax the 40-Year Age Threshold for Pharmacologic Cholesterol Lowering. Journal of the American College of Cardiology, 2021, 78, 1965-1967.	2.8	3
22	Abstract 012: Performance of Polygenic Risk Scores for Coronary Artery Disease in the Million Veteran Program. Circulation, 2019, 139, .	1.6	1
23	Editorial commentary: A new era for preventive cardiology. Trends in Cardiovascular Medicine, 2022, 32, 195-197.	4.9	0
24	Abstract 13601: Risk of Coronary Artery Disease Associated With Familial Hypercholesterolemia Genetic Variants is Independent of Historical Low-density Lipoprotein Cholesterol Exposure. Circulation, 2020, 142, .	1.6	0