

# Svati Shah

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4069047/publications.pdf>

Version: 2024-02-01

11  
papers

737  
citations

1307594

7  
h-index

1281871

11  
g-index

11  
all docs

11  
docs citations

11  
times ranked

2208  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association study reveals novel genetic loci: a new polygenic risk score for mitral valve prolapse. <i>European Heart Journal</i> , 2022, 43, 1668-1680.	2.2	25
2	Comprehensive nutrient consumption estimation and metabolic profiling during ketogenic diet and relationship with myocardial glucose uptake on FDG-PET. <i>European Heart Journal Cardiovascular Imaging</i> , 2022, 23, 1690-1697.	1.2	4
3	Sexual Dimorphism in Cardiovascular Biomarkers: Clinical and Research Implications. <i>Circulation Research</i> , 2022, 130, 578-592.	4.5	13
4	Phenome-wide Association Study of Severe COVID-19 Genetic Risk Variants. <i>Journal of the American Heart Association</i> , 2022, 11, e024004.	3.7	10
5	Insulin Growth Factor Phenotypes in Heart Failure With Preserved Ejection Fraction, an INSPIRE Registry and CATHGEN Study. <i>Journal of Cardiac Failure</i> , 2022, 28, 935-946.	1.7	2
6	Association of obesity with cardiovascular outcomes in patients with type 2 diabetes and cardiovascular disease: Insights from TECOS. <i>American Heart Journal</i> , 2020, 219, 47-57.	2.7	45
7	The Project Baseline Health Study: a step towards a broader mission to map human health. <i>Npj Digital Medicine</i> , 2020, 3, 84.	10.9	38
8	Pilot study of myocardial ischemia-induced metabolomic changes in emergency department patients undergoing stress testing. <i>PLoS ONE</i> , 2019, 14, e0211762.	2.5	7
9	Multi-ethnic genome-wide association study for atrial fibrillation. <i>Nature Genetics</i> , 2018, 50, 1225-1233.	21.4	552
10	Short-term effects of air temperature on plasma metabolite concentrations in patients undergoing cardiac catheterization. <i>Environmental Research</i> , 2016, 151, 224-232.	7.5	5
11	Association of exome sequences with plasma C-reactive protein levels in >9000 participants. <i>Human Molecular Genetics</i> , 2015, 24, 559-571.	2.9	36