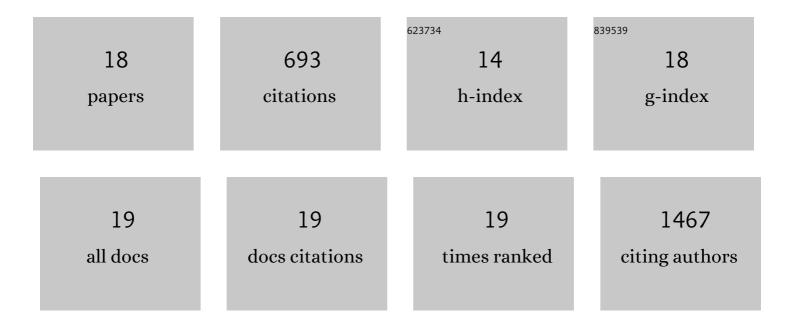
Hassan Foroughi Asl

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genomic profile – a possible diagnostic and prognostic marker in upper tract urothelial carcinoma. BJU International, 2022, 130, 92-101.	2.5	5
2	Cell-free tumour DNA analysis detects copy number alterations in gastro-oesophageal cancer patients. PLoS ONE, 2021, 16, e0245488.	2.5	13
3	Proteogenomics of non-small cell lung cancer reveals molecular subtypes associated with specific therapeutic targets and immune-evasion mechanisms. Nature Cancer, 2021, 2, 1224-1242.	13.2	37
4	Network analysis of coronary artery disease risk genes elucidates disease mechanisms and druggable targets. Scientific Reports, 2018, 8, 3434.	3.3	43
5	Genetic Susceptibility Loci for Cardiovascular Disease and Their Impact on Atherosclerotic Plaques. Circulation Genomic and Precision Medicine, 2018, 11, e002115.	3.6	20
6	Smoking is Associated to DNA Methylation in Atherosclerotic Carotid Lesions. Circulation Genomic and Precision Medicine, 2018, 11, e002030.	3.6	23
7	Poliovirus Receptor–Related 2. Arteriosclerosis, Thrombosis, and Vascular Biology, 2017, 37, 534-542.	2.4	23
8	Additional Candidate Genes for Human Atherosclerotic Disease Identified Through Annotation Based on Chromatin Organization. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	17
9	Network analysis reveals a causal role of mitochondrial gene activity in atherosclerotic lesion formation. Atherosclerosis, 2017, 267, 39-48.	0.8	26
10	Systematic analysis of chromatin interactions at disease associated loci links novel candidate genes to inflammatory bowel disease. Genome Biology, 2016, 17, 247.	8.8	39
11	Human Validation of Genes Associated With a Murine Atherosclerotic Phenotype. Arteriosclerosis, Thrombosis, and Vascular Biology, 2016, 36, 1240-1246.	2.4	44
12	Cross-Tissue Regulatory Gene Networks in Coronary Artery Disease. Cell Systems, 2016, 2, 196-208.	6.2	120
13	Variants in ALOX5, ALOX5AP and LTA4H are not associated with atherosclerotic plaque phenotypes: The Athero-Express Genomics Study. Atherosclerosis, 2015, 239, 528-538.	0.8	22
14	Prediction of Causal Candidate Genes in Coronary Artery Disease Loci. Arteriosclerosis, Thrombosis, and Vascular Biology, 2015, 35, 2207-2217.	2.4	101
15	Expression Quantitative Trait Loci Acting Across Multiple Tissues Are Enriched in Inherited Risk for Coronary Artery Disease. Circulation: Cardiovascular Genetics, 2015, 8, 305-315.	5.1	39
16	Plasma Cholesterol–Induced Lesion Networks Activated before Regression of Early, Mature, and Advanced Atherosclerosis. PLoS Genetics, 2014, 10, e1004201.	3.5	64
17	kruX: matrix-based non-parametric eQTL discovery. BMC Bioinformatics, 2014, 15, 11.	2.6	39
18	Lim Domain Binding 2. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 2068-2077.	2.4	17