Paul Cheng

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8822492/publications.pdf

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11 papers	1,006 citations	1040056 9 h-index	1372567 10 g-index
13 all docs	13 docs citations	13 times ranked	788 citing authors

#	Article	IF	Citations
1	ZEB2 Shapes the Epigenetic Landscape of Atherosclerosis. Circulation, 2022, 145, 469-485.	1.6	31
2	Axicabtagene Ciloleucel as Second-Line Therapy for Large B-Cell Lymphoma. New England Journal of Medicine, 2022, 386, 640-654.	27.0	586
3	Smad3 regulates smooth muscle cell fate and mediates adverse remodeling and calcification of the atherosclerotic plaque. , 2022, 1, 322-333.		21
4	Embryologic Origin Influences Smooth Muscle Cell Phenotypic Modulation Signatures in Murine Marfan Syndrome Aortic Aneurysm. Arteriosclerosis, Thrombosis, and Vascular Biology, 2022, 42, 1154-1168.	2.4	11
5	Ibrutinib-associated atrial fibrillation treatment with catheter ablation. HeartRhythm Case Reports, 2021, 7, 713-716.	0.4	4
6	Coronary Disease-Associated Gene TCF21 Inhibits Smooth Muscle Cell Differentiation by Blocking the Myocardin-Serum Response Factor Pathway. Circulation Research, 2020, 126, 517-529.	4.5	67
7	Single-Cell Transcriptomic Profiling of Vascular Smooth Muscle Cell Phenotype Modulation in Marfan Syndrome Aortic Aneurysm. Arteriosclerosis, Thrombosis, and Vascular Biology, 2020, 40, 2195-2211.	2.4	126
8	Molecular mechanisms of coronary disease revealed using quantitative trait loci for TCF21 binding, chromatin accessibility, and chromosomal looping. Genome Biology, 2020, 21, 135.	8.8	16
9	Outcomes in Patients With Cardiac Amyloidosis Undergoing Heart Transplantation. JACC: Heart Failure, 2020, 8, 461-468.	4.1	46
10	Environment-Sensing Aryl Hydrocarbon Receptor Inhibits the Chondrogenic Fate of Modulated Smooth Muscle Cells in Atherosclerotic Lesions. Circulation, 2020, 142, 575-590.	1.6	57
11	Coronary artery disease genes SMAD3 and TCF21 promote opposing interactive genetic programs that regulate smooth muscle cell differentiation and disease risk. PLoS Genetics, 2018, 14, e1007681.	3.5	41