

Min-Lee Yang

List of Publications by Year in descending order

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18
papers

894
citations

623734

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839539

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docs citations

19
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2688
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular genetic evaluation of pediatric renovascular hypertension due to renal artery stenosis and abdominal aortic coarctation in neurofibromatosis type 1. <i>Human Molecular Genetics</i> , 2022, 31, 334-346.	2.9	2
2	Spontaneous coronary artery dissection is infrequent in individuals with heritable thoracic aortic disease despite partially shared genetic susceptibility. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 1448-1456.	1.2	2
3	Rare loss-of-function mutations of <i>PTGIR</i> are enriched in fibromuscular dysplasia. <i>Cardiovascular Research</i> , 2021, 117, 1154-1165.	3.8	20
4	An Asian-specific <i>MPL</i> genetic variant alters JAK-STAT signaling and influences platelet count in the population. <i>Human Molecular Genetics</i> , 2021, 30, 836-842.	2.9	4
5	Epidemiologic and Genetic Associations of Erythropoietin With Blood Pressure, Hypertension, and Coronary Artery Disease. <i>Hypertension</i> , 2021, 78, 1555-1566.	2.7	1
6	Genetic investigation of fibromuscular dysplasia identifies risk loci and shared genetics with common cardiovascular diseases. <i>Nature Communications</i> , 2021, 12, 6031.	12.8	34
7	Common variants in signaling transcription-factor-binding sites drive phenotypic variability in red blood cell traits. <i>Nature Genetics</i> , 2020, 52, 1333-1345.	21.4	24
8	Chromosome 1q21.2 and additional loci influence risk of spontaneous coronary artery dissection and myocardial infarction. <i>Nature Communications</i> , 2020, 11, 4432.	12.8	60
9	A Novel Recurrent <i>COL5A1</i> Genetic Variant Is Associated With a Dysplasia-Associated Arterial Disease Exhibiting Dissections and Fibromuscular Dysplasia. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 2686-2699.	2.4	30
10	Genome-wide Trans-ethnic Meta-analysis Identifies Seven Genetic Loci Influencing Erythrocyte Traits and a Role for RBPMS in Erythropoiesis. <i>American Journal of Human Genetics</i> , 2017, 100, 51-63.	6.2	45
11	Loss-of-Function Mutations in <i>YY1AP1</i> Lead to Grange Syndrome and a Fibromuscular Dysplasia-Like Vascular Disease. <i>American Journal of Human Genetics</i> , 2017, 100, 21-30.	6.2	54
12	<i>PHACTR1</i> Is a Genetic Susceptibility Locus for Fibromuscular Dysplasia Supporting Its Complex Genetic Pattern of Inheritance. <i>PLoS Genetics</i> , 2016, 12, e1006367.	3.5	146
13	Exome-wide association analysis reveals novel coding sequence variants associated with lipid traits in Chinese. <i>Nature Communications</i> , 2015, 6, 10206.	12.8	86
14	Clinical and biochemical profiles suggest fibromuscular dysplasia is a systemic disease with altered <i>TGFβ2</i> expression and connective tissue features. <i>FASEB Journal</i> , 2014, 28, 3313-3324.	0.5	68
15	Effects of Long-Term Averaging of Quantitative Blood Pressure Traits on the Detection of Genetic Associations. <i>American Journal of Human Genetics</i> , 2014, 95, 49-65.	6.2	73
16	Genomic Estimates of Aneuploid Content in Glioblastoma Multiforme and Improved Classification. <i>Clinical Cancer Research</i> , 2012, 18, 5595-5605.	7.0	34
17	A gender-specific association of CNV at 6p21.3 with NPC susceptibility. <i>Human Molecular Genetics</i> , 2011, 20, 2889-2896.	2.9	44
18	Genome-wide Association Study Reveals Multiple Nasopharyngeal Carcinoma-Associated Loci within the HLA Region at Chromosome 6p21.3. <i>American Journal of Human Genetics</i> , 2009, 85, 194-203.	6.2	166