

Cristina Barlassina

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

86
papers

4,976
citations

36
h-index

70
g-index

92
ext. papers

5,843
ext. citations

7.6
avg, IF

4.02
L-index

#	Paper	IF	Citations
86	Dissecting the Polygenic Basis of Primary Hypertension: Identification of Key Pathway-Specific Components.. <i>Frontiers in Cardiovascular Medicine</i> , 2022 , 9, 814502	5.4	0
85	A multi-step genomic approach prioritized TBKBP1 gene as relevant for multiple sclerosis susceptibility.. <i>Journal of Neurology</i> , 2022 , 1	5.5	0
84	Transethnic Genome-Wide Association Study Provides Insights in the Genetic Architecture and Heritability of Long QT Syndrome. <i>Circulation</i> , 2020 , 142, 324-338	16.7	27
83	Klotho Gene in Human Salt-Sensitive Hypertension. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2020 , 15, 375-383	6.9	14
82	Transcriptome Analysis of iPSC-Derived Neurons from Rubinstein-Taybi Patients Reveals Deficits in Neuronal Differentiation. <i>Molecular Neurobiology</i> , 2020 , 57, 3685-3701	6.2	6
81	Immune signature drives leukemia escape and relapse after hematopoietic cell transplantation. <i>Nature Medicine</i> , 2019 , 25, 603-611	50.5	136
80	A longitudinal study highlights shared aspects of the transcriptomic response to cardiogenic and septic shock. <i>Critical Care</i> , 2019 , 23, 414	10.8	6
79	gDNA qPCR is statistically more reliable than mRNA analysis in detecting leukemic cells to monitor CML. <i>Cell Death and Disease</i> , 2018 , 9, 349	9.8	3
78	Identification of a transcriptome profile associated with improvement of organ function in septic shock patients after early supportive therapy. <i>Critical Care</i> , 2018 , 22, 312	10.8	5
77	Claudin-14 Gene Polymorphisms and Urine Calcium Excretion. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2018 , 13, 1542-1549	6.9	8
76	Next-generation sequencing of a family with a high penetrance of monoclonal gammopathies for the identification of candidate risk alleles. <i>Cancer</i> , 2017 , 123, 3701-3708	6.4	8
75	Self-renewal and phenotypic conversion are the main physiological responses of macrophages to the endogenous estrogen surge. <i>Scientific Reports</i> , 2017 , 7, 44270	4.9	37
74	PEAR1 is not a major susceptibility gene for cardiovascular disease in a Flemish population. <i>BMC Medical Genetics</i> , 2017 , 18, 45	2.1	10
73	Genome-Wide and Gene-Based Meta-Analyses Identify Novel Loci Influencing Blood Pressure Response to Hydrochlorothiazide. <i>Hypertension</i> , 2017 , 69, 51-59	8.5	25
72	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. <i>Nature Communications</i> , 2016 , 7, 13357	17.4	46
71	ShockOmics: multiscale approach to the identification of molecular biomarkers in acute heart failure induced by shock. <i>Scandinavian Journal of Trauma, Resuscitation and Emergency Medicine</i> , 2016 , 24, 9	3.6	11
70	Interaction between polyphenols intake and PON1 gene variants on markers of cardiovascular disease: a nutrigenetic observational study. <i>Journal of Translational Medicine</i> , 2016 , 14, 186	8.5	29

69	PHACTR1 Is a Genetic Susceptibility Locus for Fibromuscular Dysplasia Supporting Its Complex Genetic Pattern of Inheritance. <i>PLoS Genetics</i> , 2016 , 12, e1006367	6	99
68	Xanthine oxidase gene variants and their association with blood pressure and incident hypertension: a population study. <i>Journal of Hypertension</i> , 2016 , 34, 2147-54	1.9	21
67	Exome sequencing identifies variants in two genes encoding the LIM-proteins NRAP and FHL1 in an Italian patient with BAG3 myofibrillar myopathy. <i>Journal of Muscle Research and Cell Motility</i> , 2016 , 37, 101-15	3.5	17
66	A genome-wide screening and SNPs-to-genes approach to identify novel genetic risk factors associated with frontotemporal dementia. <i>Neurobiology of Aging</i> , 2015 , 36, 2904.e13-26	5.6	34
65	The burden of multiple sclerosis variants in continental Italians and Sardinians. <i>Multiple Sclerosis Journal</i> , 2015 , 21, 1385-95	5	9
64	Pharmacogenomics considerations in the control of hypertension. <i>Pharmacogenomics</i> , 2015 , 16, 1951-64	2.6	9
63	Inactive matrix Gla protein is causally related to adverse health outcomes: a Mendelian randomization study in a Flemish population. <i>Hypertension</i> , 2015 , 65, 463-70	8.5	66
62	Genome-wide association study of kidney function decline in individuals of European descent. <i>Kidney International</i> , 2015 , 87, 1017-29	9.9	83
61	Coronary risk in relation to genetic variation in MEOX2 and TCF15 in a Flemish population. <i>BMC Genetics</i> , 2015 , 16, 116	2.6	10
60	Exome sequencing in seven families and gene-based association studies indicate genetic heterogeneity and suggest possible candidates for fibromuscular dysplasia. <i>Journal of Hypertension</i> , 2015 , 33, 1802-10; discussion 1810	1.9	24
59	TET2 and CSMD1 genes affect SBP response to hydrochlorothiazide in never-treated essential hypertensives. <i>Journal of Hypertension</i> , 2015 , 33, 1301-9	1.9	20
58	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015 , 11, e1005378	6	220
57	Cross-disorder genome-wide analyses suggest a complex genetic relationship between Tourette's syndrome and OCD. <i>American Journal of Psychiatry</i> , 2015 , 172, 82-93	11.9	101
56	An integrated Diet Monitoring Solution for nutrigenomic research. <i>Studies in Health Technology and Informatics</i> , 2015 , 210, 632-6	0.5	2
55	Discovery of new risk loci for IgA nephropathy implicates genes involved in immunity against intestinal pathogens. <i>Nature Genetics</i> , 2014 , 46, 1187-96	36.3	325
54	Genome-wide association study identifies CAMKID variants involved in blood pressure response to losartan: the SOPHIA study. <i>Pharmacogenomics</i> , 2014 , 15, 1643-52	2.6	24
53	Sardinians genetic background explained by runs of homozygosity and genomic regions under positive selection. <i>PLoS ONE</i> , 2014 , 9, e91237	3.7	32
52	Dietary Salt Intake, Blood Pressure, and Genes. <i>Current Nutrition Reports</i> , 2013 , 2, 134-141	6	1

51	Identification of heart rate-associated loci and their effects on cardiac conduction and rhythm disorders. <i>Nature Genetics</i> , 2013 , 45, 621-31	36.3	219
50	Target sequencing, cell experiments, and a population study establish endothelial nitric oxide synthase (eNOS) gene as hypertension susceptibility gene. <i>Hypertension</i> , 2013 , 62, 844-52	8.5	39
49	Sex-stratified genome-wide association studies including 270,000 individuals show sexual dimorphism in genetic loci for anthropometric traits. <i>PLoS Genetics</i> , 2013 , 9, e1003500	6	277
48	Identification By Gene Expression Profiling Of CIITA-Dependent HLA Class II Transcriptional Downregulation As a Novel Mechanism Of Leukemia Immune Escape and Relapse After Allogeneic HSCT. <i>Blood</i> , 2013 , 122, 3748-3748	2.2	
47	Copy-number disorders are a common cause of congenital kidney malformations. <i>American Journal of Human Genetics</i> , 2012 , 91, 987-97	11	161
46	Genomewide association study using a high-density single nucleotide polymorphism array and case-control design identifies a novel essential hypertension susceptibility locus in the promoter region of endothelial NO synthase. <i>Hypertension</i> , 2012 , 59, 248-55	8.5	124
45	An overview of the genetic structure within the Italian population from genome-wide data. <i>PLoS ONE</i> , 2012 , 7, e43759	3.7	44
44	Genes involved in vasoconstriction and vasodilation system affect salt-sensitive hypertension. <i>PLoS ONE</i> , 2011 , 6, e19620	3.7	48
43	Population Stratification Analysis in Genome-Wide Association Studies 2011 , 177-196		
42	Genomic and Transcriptional Immunoediting of Acute Myeloid Leukemia in Response to Allogeneic Hematopoietic Stem Cell Transplantation. <i>Blood</i> , 2011 , 118, 329-329	2.2	
41	alpha- and beta-Adducin polymorphisms affect podocyte proteins and proteinuria in rodents and decline of renal function in human IgA nephropathy. <i>Journal of Molecular Medicine</i> , 2010 , 88, 203-17	5.5	16
40	Loss of mismatched HLA in leukemia after stem-cell transplantation. <i>New England Journal of Medicine</i> , 2009 , 361, 478-88	59.2	337
39	Common genetic variants and haplotypes in renal CLCNKA gene are associated to salt-sensitive hypertension. <i>Human Molecular Genetics</i> , 2007 , 16, 1630-8	5.6	63
38	Gly460Trp alpha-adducin gene polymorphism and endothelial function in untreated hypertensive patients. <i>Journal of Hypertension</i> , 2007 , 25, 2234-9	1.9	8
37	Alpha-adducin and angiotensin-converting enzyme polymorphisms in hypertension: evidence for a joint influence on albuminuria. <i>Journal of Hypertension</i> , 2006 , 24, 931-7	1.9	16
36	The 460Trp allele of alpha-adducin increases carotid intima-media thickness in young adult males. <i>Journal of Hypertension</i> , 2006 , 24, 697-703	1.9	10
35	βAdducin and angiotensin-converting enzyme polymorphisms in hypertension: evidence for a joint influence on albuminuria. <i>Journal of Hypertension</i> , 2006 , 24, 1217	1.9	
34	Role of the adducin family genes in human essential hypertension. <i>Journal of Hypertension</i> , 2005 , 23, 543-9	1.9	41

33	Association between aldosterone synthase (CYP11B2) polymorphism and left ventricular mass in human essential hypertension. <i>Journal of the American College of Cardiology</i> , 2004 , 43, 265-70	15.1	48
32	Endothelial nitric oxide synthase polymorphisms are associated with type 2 diabetes and the insulin resistance syndrome. <i>Diabetes</i> , 2003 , 52, 1270-5	0.9	159
31	Genetic polymorphism of the renin-angiotensin-aldosterone system and arterial hypertension in the Italian population: the GENIPER Project. <i>Journal of Hypertension</i> , 2003 , 21, 1853-60	1.9	46
30	Angiotensinogen gene polymorphism, again?. <i>Journal of Hypertension</i> , 2003 , 21, 1815-8	1.9	2
29	Haematological phenotypes in relation to the C1797T Adducin polymorphism in a Caucasian population. <i>Clinical Science</i> , 2003 , 104, 369	6.5	1
28	Expression analysis of the human adducin gene family and evidence of ADD2 beta4 multiple splicing variants. <i>Biochemical and Biophysical Research Communications</i> , 2003 , 309, 359-67	3.4	14
27	ACE and alpha-adducin polymorphism as markers of individual response to diuretic therapy. <i>Hypertension</i> , 2003 , 41, 398-403	8.5	149
26	Association between hypertension and variation in the alpha- and beta-adducin genes in a white population. <i>Kidney International</i> , 2002 , 62, 2152-9	9.9	54
25	alpha-Adducin 460Trp allele is associated with erythrocyte Na transport rate in North Sardinian primary hypertensives. <i>Hypertension</i> , 2002 , 39, 357-62	8.5	58
24	Genetics of essential hypertension: from families to genes. <i>Journal of the American Society of Nephrology: JASN</i> , 2002 , 13 Suppl 3, S155-64	12.7	40
23	Carotid and femoral intima-media thickness in relation to three candidate genes in a Caucasian population. <i>Journal of Hypertension</i> , 2002 , 20, 1551-61	1.9	50
22	Carotid and femoral artery stiffness in relation to three candidate genes in a white population. <i>Hypertension</i> , 2001 , 38, 1190-7	8.5	77
21	Effects of three candidate genes on prevalence and incidence of hypertension in a Caucasian population. <i>Journal of Hypertension</i> , 2001 , 19, 1349-58	1.9	179
20	Synergistic effect of alpha-adducin and ACE genes causes blood pressure changes with body sodium and volume expansion. <i>Kidney International</i> , 2000 , 57, 1083-90	9.9	64
19	Alpha-adducin polymorphism in hypertensives of South African ancestry. <i>American Journal of Hypertension</i> , 2000 , 13, 719-23	2.3	36
18	The role of alpha-adducin polymorphism in blood pressure and sodium handling regulation may not be excluded by a negative association study. <i>Hypertension</i> , 1999 , 34, 649-54	8.5	139
17	Adducin polymorphism affects renal proximal tubule reabsorption in hypertension. <i>Hypertension</i> , 1999 , 33, 694-7	8.5	108
16	Alpha-adducin polymorphisms and renal sodium handling in essential hypertensive patients. <i>Kidney International</i> , 1998 , 53, 1471-8	9.9	103

15	Adducin in essential hypertension. <i>FEBS Letters</i> , 1998 , 430, 41-4	3.8	26
14	Alpha-adducin gene polymorphism and cardiovascular phenotypes in a general population. <i>Journal of Hypertension</i> , 1997 , 15, 1707-10	1.9	31
13	Genetics of renal mechanisms of primary hypertension: the role of adducin. <i>Journal of Hypertension</i> , 1997 , 15, 1567-71	1.9	13
12	Polymorphisms of alpha-adducin and salt sensitivity in patients with essential hypertension. <i>Lancet, The</i> , 1997 , 349, 1353-7	4.0	473
11	alpha-adducin may control blood pressure both in rats and humans. <i>Clinical and Experimental Pharmacology and Physiology</i> , 1995 , 22, S7-9	3	8
10	Association of the alpha-adducin locus with essential hypertension. <i>Hypertension</i> , 1995 , 25, 320-6	8.5	106
9	Erythrocyte Na ⁺ ,K ⁺ ,Cl ⁻ cotransport and kidney function in essential hypertension. <i>Journal of Hypertension</i> , 1993 , 11, 805-13	1.9	18
8	Genetic models of arterial hypertension--role of tubular ion transport. <i>Pediatric Nephrology</i> , 1993 , 7, 865-70	3.2	3
7	Heritability estimate of erythrocyte Na-K-Cl cotransport in normotensive and hypertensive families. <i>American Journal of Hypertension</i> , 1991 , 4, 725-34	2.3	23
6	Pathogenetic mechanisms in essential hypertension. Analogies between a rat model and the human disease. <i>International Journal of Cardiology</i> , 1989 , 25 Suppl 1, S29-36	3.2	3
5	Membrane Abnormalities in Essential Hypertension:.. <i>Annals of the New York Academy of Sciences</i> , 1986 , 488, 266-275	6.5	1
4	Heritability of Sodium Transport Systems and Hypertension. <i>Annals of the New York Academy of Sciences</i> , 1986 , 488, 576-578	6.5	3
3	Heritability of Sodium Transport Systems and Hypertension. <i>Annals of the New York Academy of Sciences</i> , 1986 , 488, 576-578	6.5	6
2	Renal dysfunction as a possible cause of essential hypertension in predisposed subjects. <i>Kidney International</i> , 1983 , 23, 870-5	9.9	48
1	Transcapillary protein escape in arterial hypertension. <i>Research in Clinic and Laboratory</i> , 1980 , 10, 163-70		2