Nicole L Glazer

List of Publications by Year
in descending order

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90 papers	27,597 citations	20817 60 h-index	40979 93 g-index
95 all docs	95 docs citations	95 times ranked	37893 citing authors

#	Article	IF	Citations
1	Pathologic gene network rewiring implicates PPP1R3A as a central regulator in pressure overload heart failure. Nature Communications, 2019, 10, 2760.	12.8	22
2	Causal Effect of Plasminogen Activator Inhibitor Type 1 on Coronary Heart Disease. Journal of the American Heart Association, 2017, 6, .	3.7	89
3	Large meta-analysis of genome-wide association studies identifies five loci for lean body mass. Nature Communications, 2017, 8, 80.	12.8	147
4	Associations between metabolic dysregulation and circulating biomarkers of fibrosis: the Cardiovascular Health Study. Metabolism: Clinical and Experimental, 2015, 64, 1316-1323.	3.4	6
5	Fibrosis-related biomarkers and large and small vessel disease: The Cardiovascular Health Study. Atherosclerosis, 2015, 239, 539-546.	0.8	18
6	Fibrosis-Related Biomarkers and Incident Cardiovascular Disease in Older Adults. Circulation: Arrhythmia and Electrophysiology, 2014, 7, 583-589.	4.8	29
7	Fibrosis-Related Biomarkers and Risk of Total and Cause-Specific Mortality. American Journal of Epidemiology, 2014, 179, 1331-1339.	3.4	23
8	Circulating fibrosis biomarkers and risk of atrial fibrillation: The Cardiovascular Health Study (CHS). American Heart Journal, 2014, 167, 723-728.e2.	2.7	33
9	Genome-Wide Association Study of <scp>l</scp> -Arginine and Dimethylarginines Reveals Novel Metabolic Pathway for Symmetric Dimethylarginine. Circulation: Cardiovascular Genetics, 2014, 7, 864-872.	5.1	53
10	A Pilot Study Identifying Statin Nonadherence With Visit-to-Visit Variability of Low-Density Lipoprotein Cholesterol. American Journal of Cardiology, 2013, 111, 1437-1442.	1.6	34
11	A Genomeâ€Wide Association Study for Venous Thromboembolism: The Extended Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. Genetic Epidemiology, 2013, 37, 512-521.	1.3	99
12	Genome-Wide Association Study of Cardiac Structure and Systolic Function in African Americans. Circulation: Cardiovascular Genetics, 2013, 6, 37-46.	5.1	46
13	Common Variants in Mendelian Kidney Disease Genes and Their Association with Renal Function. Journal of the American Society of Nephrology: JASN, 2013, 24, 2105-2117.	6.1	33
14	Sustained and Shorter Bouts of Physical Activity Are Related to Cardiovascular Health. Medicine and Science in Sports and Exercise, 2013, 45, 109-115.	0.4	161
15	Genetic variation associated with circulating monocyte count in the eMERGE Network. Human Molecular Genetics, 2013, 22, 2119-2127.	2.9	56
16	Genome-Wide Association Study of Retinopathy in Individuals without Diabetes. PLoS ONE, 2013, 8, e54232.	2.5	22
17	Genetic Loci for Retinal Arteriolar Microcirculation. PLoS ONE, 2013, 8, e65804.	2.5	27
18	Genome-Wide Association and Functional Follow-Up Reveals New Loci for Kidney Function. PLoS Genetics, 2012, 8, e1002584.	3.5	166

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19	Integration of genome-wide association studies with biological knowledge identifies six novel genes related to kidney function. Human Molecular Genetics, 2012, 21, 5329-5343.	2.9	64
20	Cardiometabolic Correlates and Heritability of Fetuin-A, Retinol-Binding Protein 4, and Fatty-Acid Binding Protein 4 in the Framingham Heart Study. Journal of Clinical Endocrinology and Metabolism, 2012, 97, E1943-E1947.	3.6	56
21	No Interactions Between Previously Associated 2-Hour Glucose Gene Variants and Physical Activity or BMI on 2-Hour Glucose Levels. Diabetes, 2012, 61, 1291-1296.	0.6	23
22	Meta-analyses identify 13 loci associated with age at menopause and highlight DNA repair and immune pathways. Nature Genetics, 2012, 44, 260-268.	21.4	303
23	Discovery and Fine Mapping of Serum Protein Loci through Transethnic Meta-analysis. American Journal of Human Genetics, 2012, 91, 744-753.	6.2	69
24	Large-Scale Gene-Centric Meta-analysis across 32 Studies Identifies Multiple Lipid Loci. American Journal of Human Genetics, 2012, 91, 823-838.	6.2	227
25	Transforming growth factor beta-1 and incidence of heart failure in older adults: The Cardiovascular Health Study. Cytokine, 2012, 60, 341-345.	3.2	14
26	Meta-analysis identifies six new susceptibility loci for atrial fibrillation. Nature Genetics, 2012, 44, 670-675.	21.4	533
27	A Genome-Wide Association Search for Type 2 Diabetes Genes in African Americans. PLoS ONE, 2012, 7, e29202.	2.5	197
28	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. Nature Genetics, 2012, 44, 659-669.	21.4	762
29	Multi-Ethnic Analysis of Lipid-Associated Loci: The NHLBI CARe Project. PLoS ONE, 2012, 7, e36473.	2.5	46
30	Gene-Centric Meta-Analysis of Lipid Traits in African, East Asian and Hispanic Populations. PLoS ONE, 2012, 7, e50198.	2.5	40
31	Genetic variation near IRS1 associates with reduced adiposity and an impaired metabolic profile. Nature Genetics, 2011, 43, 753-760.	21.4	289
32	Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. Nature, 2011, 478, 103-109.	27.8	1,855
33	New gene functions in megakaryopoiesis and platelet formation. Nature, 2011, 480, 201-208.	27.8	401
34	Genetic variability within the cholesterol lowering pathway and the effectiveness of statins in reducing the risk of MI. Atherosclerosis, 2011, 217, 458-464.	0.8	38
35	Common Genetic Determinants of Vitamin D Insufficiency: A Genome-Wide Association Study. Obstetrical and Gynecological Survey, 2011, 66, 91-93.	0.4	0
36	Cerivastatin, genetic variants, and the risk of rhabdomyolysis. Pharmacogenetics and Genomics, 2011, 21, 280-288.	1.5	90

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37	Genetic variation associated with plasma von Willebrand factor levels and the risk of incident venous thrombosis. Blood, 2011, 117, 6007-6011.	1.4	97
38	Meta-analysis identifies 29 additional ulcerative colitis risk loci, increasing the number of confirmed associations to 47. Nature Genetics, 2011, 43, 246-252.	21.4	1,201
39	Genome-Wide Association Study for Coronary Artery Calcification With Follow-Up in Myocardial Infarction. Circulation, 2011, 124, 2855-2864.	1.6	269
40	Ascertainment of warfarin and aspirin use by medical record review compared with automated pharmacy data. Pharmacoepidemiology and Drug Safety, 2011, 20, 313-316.	1.9	32
41	Genomeâ€wide association studies of cerebral white matter lesion burden. Annals of Neurology, 2011, 69, 928-939.	5.3	201
42	CUBN Is a Gene Locus for Albuminuria. Journal of the American Society of Nephrology: JASN, 2011, 22, 555-570.	6.1	208
43	A genome-wide association study identifies novel loci associated with circulating IGF-I and IGFBP-3. Human Molecular Genetics, 2011, 20, 1241-1251.	2.9	67
44	A Bivariate Genome-Wide Approach to Metabolic Syndrome. Diabetes, 2011, 60, 1329-1339.	0.6	226
45	Large-Scale Candidate Gene Analysis in Whites and African Americans Identifies IL6R Polymorphism in Relation to Atrial Fibrillation. Circulation: Cardiovascular Genetics, 2011, 4, 557-564.	5.1	74
46	Association of genetic variation with systolic and diastolic blood pressure among African Americans: the Candidate Gene Association Resource study. Human Molecular Genetics, 2011, 20, 2273-2284.	2.9	168
47	Plasma symmetric dimethylarginine reference limits from the Framingham offspring cohort. Clinical Chemistry and Laboratory Medicine, 2011, 49, 1907-10.	2.3	28
48	Reference Intervals for Plasma L-Arginine and the L-Arginine:Asymmetric Dimethylarginine Ratio in the Framingham Offspring Cohort. Journal of Nutrition, 2011, 141, 2186-2190.	2.9	63
49	Copy Number Variation Contributes to Sporadic and Familial Thoracic Aortic Aneurysms and Dissections. Circulation: Cardiovascular Genetics, 2011, 4, 212-213.	5.1	0
50	Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. Nature Genetics, 2011, 43, 1005-1011.	21.4	403
51	Multiple Loci Are Associated with White Blood Cell Phenotypes. PLoS Genetics, 2011, 7, e1002113.	3.5	106
52	Moving Beyond Genome-Wide Association Studies. Circulation: Cardiovascular Genetics, 2011, 4, 91-93.	5.1	2
53	Four Genetic Loci Influencing Electrocardiographic Indices of Left Ventricular Hypertrophy. Circulation: Cardiovascular Genetics, 2011, 4, 626-635.	5.1	28
54	Variation in the ATM Gene May Alter Glycemic Response to Metformin. Circulation: Cardiovascular Genetics, 2011, 4, 210-211.	5.1	5

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55	Diabetes Mellitus, Glycemic Control, and Risk of Atrial Fibrillation. Journal of General Internal Medicine, 2010, 25, 853-858.	2.6	238
56	Genetic predictors of medically refractory ulcerative colitis. Inflammatory Bowel Diseases, 2010, 16, 1830-1840.	1.9	135
57	Hundreds of variants clustered in genomic loci and biological pathways affect human height. Nature, 2010, 467, 832-838.	27.8	1,789
58	Genetic variation in GIPR influences the glucose and insulin responses to an oral glucose challenge. Nature Genetics, 2010, 42, 142-148.	21.4	591
59	Common variants in KCNN3 are associated with lone atrial fibrillation. Nature Genetics, 2010, 42, 240-244.	21.4	438
60	Genome-wide association identifies multiple ulcerative colitis susceptibility loci. Nature Genetics, 2010, 42, 332-337.	21.4	572
61	New loci associated with kidney function and chronic kidney disease. Nature Genetics, 2010, 42, 376-384.	21.4	710
62	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. Nature Genetics, 2010, 42, 949-960.	21.4	836
63	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. Nature Genetics, 2010, 42, 937-948.	21.4	2,634
64	Uromodulin Levels Associate with a Common UMOD Variant and Risk for Incident CKD. Journal of the American Society of Nephrology: JASN, 2010, 21, 337-344.	6.1	146
65	Association of Genome-Wide Variation With the Risk of Incident Heart Failure in Adults of European and African Ancestry. Circulation: Cardiovascular Genetics, 2010, 3, 256-266.	5.1	176
66	Common variants in the calcium-sensing receptor gene are associated with total serum calcium levels. Human Molecular Genetics, 2010, 19, 4296-4303.	2.9	86
67	Large-scale genomic studies reveal central role of ABO in sP-selectin and sICAM-1 levels. Human Molecular Genetics, 2010, 19, 1863-1872.	2.9	233
68	Genomic Variation Associated With Mortality Among Adults of European and African Ancestry With Heart Failure. Circulation: Cardiovascular Genetics, 2010, 3, 248-255.	5.1	80
69	Genome-wide association identifies OBFC1 as a locus involved in human leukocyte telomere biology. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 9293-9298.	7.1	244
70	Common Genetic Variants Associate with Serum Phosphorus Concentration. Journal of the American Society of Nephrology: JASN, 2010, 21, 1223-1232.	6.1	123
71	Genome-Wide Association Studies of Serum Magnesium, Potassium, and Sodium Concentrations Identify Six Loci Influencing Serum Magnesium Levels. PLoS Genetics, 2010, 6, e1001045.	3.5	185
72	Four Novel Loci (19q13, 6q24, 12q24, and 5q14) Influence the Microcirculation In Vivo. PLoS Genetics, 2010, 6, e1001184.	3.5	134

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73	Multiple Genetic Loci Influence Serum Urate Levels and Their Relationship With Gout and Cardiovascular Disease Risk Factors. Circulation: Cardiovascular Genetics, 2010, 3, 523-530.	5.1	285
74	Myocardial infarction and stroke associated with diuretic based two drug antihypertensive regimens: population based case-control study. BMJ: British Medical Journal, 2010, 340, c103-c103.	2.3	26
75	A Meta-analysis of Four Genome-Wide Association Studies of Survival to Age 90 Years or Older: The Cohorts for Heart and Aging Research in Genomic Epidemiology Consortium. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2010, 65A, 478-487.	3.6	117
76	New genetic loci implicated in fasting glucose homeostasis and their impact on type 2 diabetes risk. Nature Genetics, 2010, 42, 105-116.	21.4	1,982
77	Common genetic determinants of vitamin D insufficiency: a genome-wide association study. Lancet, The, 2010, 376, 180-188.	13.7	1,385
78	Genomewide Association Studies of Stroke. New England Journal of Medicine, 2009, 360, 1718-1728.	27.0	420
79	Genetic Variants Associated With Cardiac Structure and Function. JAMA - Journal of the American Medical Association, 2009, 302, 168.	7.4	202
80	NRXN3 Is a Novel Locus for Waist Circumference: A Genome-Wide Association Study from the CHARGE Consortium. PLoS Genetics, 2009, 5, e1000539.	3.5	230
81	Antihypertensive Treatment With ACE Inhibitors or Â-Blockers and Risk of Incident Atrial Fibrillation in a General Hypertensive Population. American Journal of Hypertension, 2009, 22, 538-544.	2.0	44
82	Multiple loci associated with indices of renal function and chronic kidney disease. Nature Genetics, 2009, 41, 712-717.	21.4	553
83	Genome-wide association study of blood pressure and hypertension. Nature Genetics, 2009, 41, 677-687.	21.4	1,224
84	Multiple loci influence erythrocyte phenotypes in the CHARGE Consortium. Nature Genetics, 2009, 41, 1191-1198.	21.4	324
85	Newly Detected Atrial Fibrillation and Compliance With Antithrombotic Guidelines. Archives of Internal Medicine, 2007, 167, 246.	3.8	149
86	Risk of New-Onset Atrial Fibrillation in Relation to Body Mass Index. Archives of Internal Medicine, 2006, 166, 2322.	3.8	258
87	Risk of Myocardial Infarction Attributable to Elevated Levels of Total Cholesterol Among Hypertensives. American Journal of Hypertension, 2005, 18, 759-766.	2.0	10
88	Weight Change and the Risk of Gestational Diabetes in Obese Women. Epidemiology, 2004, 15, 733-737.	2.7	125
89	Microenvironmental VEGF concentration, not total dose, determines a threshold between normal and aberrant angiogenesis. Journal of Clinical Investigation, 2004, 113, 516-527.	8.2	440
90	Angiopoietin-1 protects the adult vasculature against plasma leakage. Nature Medicine, 2000, 6, 460-463.	30.7	1,172