Hooman Allayee

List of Publications by Year in descending order

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95 papers 15,847 citations

50276 46 h-index 92 g-index

98 all docs 98 docs citations 98 times ranked 25150 citing authors

#	Article	IF	Citations
1	Gut flora metabolism of phosphatidylcholine promotes cardiovascular disease. Nature, 2011, 472, 57-63.	27.8	4,238
2	Large-scale association analysis identifies 13 new susceptibility loci for coronary artery disease. Nature Genetics, 2011, 43, 333-338.	21.4	1,685
3	The Collaborative Cross, a community resource for the genetic analysis of complex traits. Nature Genetics, 2004, 36, 1133-1137.	21.4	1,034
4	Trimethylamine-N-Oxide, a Metabolite Associated with Atherosclerosis, Exhibits Complex Genetic and Dietary Regulation. Cell Metabolism, 2013, 17, 49-60.	16.2	794
5	Exome sequencing identifies rare LDLR and APOA5 alleles conferring risk for myocardial infarction. Nature, 2015, 518, 102-106.	27.8	581
6	Arachidonate 5-Lipoxygenase Promoter Genotype, Dietary Arachidonic Acid, and Atherosclerosis. New England Journal of Medicine, 2004, 350, 29-37.	27.0	571
7	An epigenetic clock analysis of race/ethnicity, sex, and coronary heart disease. Genome Biology, 2016, 17, 171.	8.8	535
8	Identification of ADAMTS7 as a novel locus for coronary atherosclerosis and association of ABO with myocardial infarction in the presence of coronary atherosclerosis: two genome-wide association studies. Lancet, The, 2011, 377, 383-392.	13.7	466
9	Relationship of Paraoxonase 1 (PON1) Gene Polymorphisms and Functional Activity With Systemic Oxidative Stress and Cardiovascular Risk. JAMA - Journal of the American Medical Association, 2008, 299, 1265.	7.4	463
10	Identification of 5-Lipoxygenase as a Major Gene Contributing to Atherosclerosis Susceptibility in Mice. Circulation Research, 2002, 91, 120-126.	4.5	387
11	Seventy-five genetic loci influencing the human red blood cell. Nature, 2012, 492, 369-375.	27.8	320
12	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. Nature Genetics, 2015, 47, 1282-1293.	21.4	294
13	Frequency of mononuclear diploid cardiomyocytes underlies natural variation in heart regeneration. Nature Genetics, 2017, 49, 1346-1353.	21.4	252
14	Influence of Leukotriene Pathway Polymorphisms on Response to Montelukast in Asthma. American Journal of Respiratory and Critical Care Medicine, 2006, 173, 379-385.	5.6	225
15	Integrating genotypic and expression data in a segregating mouse population to identify 5-lipoxygenase as a susceptibility gene for obesity and bone traits. Nature Genetics, 2005, 37, 1224-1233.	21.4	210
16	Increased hepatic fat in overweight Hispanic youth influenced by interaction between genetic variation in PNPLA3 and high dietary carbohydrate and sugar consumption. American Journal of Clinical Nutrition, 2010, 92, 1522-1527.	4.7	175
17	Cognitive effects of estradiol after menopause. Neurology, 2016, 87, 699-708.	1.1	162
18	Clinical and Genetic Association of Serum Paraoxonase and Arylesterase Activities With Cardiovascular Risk. Arteriosclerosis, Thrombosis, and Vascular Biology, 2012, 32, 2803-2812.	2.4	153

#	Article	IF	CITATIONS
19	The Hybrid Mouse Diversity Panel: a resource for systems genetics analyses of metabolic and cardiovascular traits. Journal of Lipid Research, 2016, 57, 925-942.	4.2	143
20	Effect of Obesity on Clinical Presentation and Response to Treatment in Asthma. Journal of Asthma, 2006, 43, 553-558.	1.7	142
21	Hybrid mouse diversity panel: a panel of inbred mouse strains suitable for analysis of complex genetic traits. Mammalian Genome, 2012, 23, 680-692.	2.2	134
22	Untargeted metabolomics identifies trimethyllysine, a TMAO-producing nutrient precursor, as a predictor of incident cardiovascular disease risk. JCI Insight, 2018, 3, .	5.0	122
23	Comparative Genome-Wide Association Studies in Mice and Humans for Trimethylamine <i>N</i> -Oxide, a Proatherogenic Metabolite of Choline and <scp>I</scp> -Carnitine. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 1307-1313.	2.4	119
24	Genome-wide analysis highlights contribution of immune system pathways to the genetic architecture of asthma. Nature Communications, 2020, 11 , 1776 .	12.8	119
25	Genome Scan for Blood Pressure in Dutch Dyslipidemic Families Reveals Linkage to a Locus on Chromosome 4p. Hypertension, 2001, 38, 773-778.	2.7	116
26	Genome-wide analysis identifies novel susceptibility loci for myocardial infarction. European Heart Journal, 2021, 42, 919-933.	2.2	113
27	Genome-wide association study and targeted metabolomics identifies sex-specific association of CPS1 with coronary artery disease. Nature Communications, 2016, 7, 10558.	12.8	108
28	Effects of <i>PNPLA3</i> on Liver Fat and Metabolic Profile in Hispanic Children and Adolescents. Diabetes, 2010, 59, 3127-3130.	0.6	100
29	Nutrigenomics, the Microbiome, and Gene-Environment Interactions: New Directions in Cardiovascular Disease Research, Prevention, and Treatment. Circulation: Cardiovascular Genetics, 2016, 9, 291-313.	5.1	99
30	5-Lipoxygenase and atherosclerosis. Current Opinion in Lipidology, 2003, 14, 447-457.	2.7	96
31	Lipoprotein(a) levels and long-term cardiovascular risk in the contemporary era of statin therapy. Journal of Lipid Research, 2010, 51, 3055-3061.	4.2	76
32	Apolipoprotein E4 is associated with improved cognitive function in Amazonian foragerâ€horticulturalists with a high parasite burden. FASEB Journal, 2017, 31, 1508-1515.	0.5	73
33	Genome-Wide Association Study Identifies Nox3 as a Critical Gene for Susceptibility to Noise-Induced Hearing Loss. PLoS Genetics, 2015, 11, e1005094.	3.5	64
34	Toxicity of urban air pollution particulate matter in developing and adult mouse brain: Comparison of total and filter-eluted nanoparticles. Environment International, 2020, 136, 105510.	10.0	64
35	Genetically determined NLRP3 inflammasome activation associates with systemic inflammation and cardiovascular mortality. European Heart Journal, 2021, 42, 1742-1756.	2.2	63
36	Bile acids profile, histopathological indices and genetic variants for non-alcoholic fatty liver disease progression. Metabolism: Clinical and Experimental, 2021, 116, 154457.	3 . 4	62

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37	Costimulation of type-2 innate lymphoid cells by GITR promotes effector function and ameliorates type 2 diabetes. Nature Communications, 2019, 10, 713.	12.8	58
38	The Effect of Montelukast and Low-Dose Theophylline on Cardiovascular Disease Risk Factors in Asthmatics. Chest, 2007, 132, 868-874.	0.8	54
39	Common polymorphisms of ALOX5 and ALOX5AP and risk of coronary artery disease. Human Genetics, 2008, 123, 399-408.	3.8	54
40	Clinical and Genetic Association of Serum Ceruloplasmin With Cardiovascular Risk. Arteriosclerosis, Thrombosis, and Vascular Biology, 2012, 32, 516-522.	2.4	54
41	Polyunsaturated Fatty Acids and Cardiovascular Disease: Implications for Nutrigenetics. Journal of Nutrigenetics and Nutrigenomics, 2009, 2, 140-148.	1.3	53
42	Selected vitamin D metabolic gene variants and risk for autism spectrum disorder in the CHARGE Study. Early Human Development, 2015, 91, 483-489.	1.8	52
43	Ambient Air Pollution Is Associated With the Severity of Coronary Atherosclerosis and Incident Myocardial Infarction in Patients Undergoing Elective Cardiac Evaluation. Journal of the American Heart Association, 2016, 5, .	3.7	51
44	Loss of Cardioprotective Effects at the <i>ADAMTS7</i> locus as a Result of Gene-Smoking Interactions. Circulation, 2017, 135, 2336-2353.	1.6	51
45	Identification of ALOX5 as a gene regulating adiposity and pancreatic function. Diabetologia, 2008, 51, 978-988.	6.3	49
46	Using Mice to Dissect Genetic Factors in Atherosclerosis. Arteriosclerosis, Thrombosis, and Vascular Biology, 2003, 23, 1501-1509.	2.4	48
47	Locus for Elevated Apolipoprotein B Levels on Chromosome 1p31 in Families With Familial Combined Hyperlipidemia. Circulation Research, 2002, 90, 926-931.	4.5	46
48	Nutrigenetic association of the 5-lipoxygenase gene with myocardial infarction. American Journal of Clinical Nutrition, 2008, 88, 934-940.	4.7	45
49	Association of serum HDL-cholesterol and apolipoprotein A1 levels with risk of severe SARS-CoV-2 infection. Journal of Lipid Research, 2021, 62, 100061.	4.2	44
50	Genetic contribution of the leukotriene pathway to coronary artery disease. Human Genetics, 2011, 129, 617-627.	3.8	42
51	Genetic Deficiency of Flavin-Containing Monooxygenase 3 (<i>Fmo3</i>) Protects Against Thrombosis but Has Only a Minor Effect on Plasma Lipid Levels—Brief Report. Arteriosclerosis, Thrombosis, and Vascular Biology, 2019, 39, 1045-1054.	2.4	41
52	Association of a Genetic Risk Score With Prevalent and Incident Myocardial Infarction in Subjects Undergoing Coronary Angiography. Circulation: Cardiovascular Genetics, 2012, 5, 441-449.	5.1	40
53	Contribution of Gut Bacteria to Lipid Levels. Circulation Research, 2015, 117, 750-754.	4.5	40
54	The Genetic Architecture of Coronary Artery Disease: Current Knowledge and Future Opportunities. Current Atherosclerosis Reports, 2017, 19, 6.	4.8	38

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55	Inflammatory Gene Variants in the Tsimane, an Indigenous Bolivian Population with a High Infectious Load. Biodemography and Social Biology, 2011, 57, 33-52.	1.0	37
56	A genome-wide set of congenic mouse strains derived from DBA/2J on a C57BL/6J background. Genomics, 2005, 86, 259-270.	2.9	36
57	Aspirin Hydrolysis in Plasma Is a Variable Function of Butyrylcholinesterase and Platelet-activating Factor Acetylhydrolase 1b2 (PAFAH1b2). Journal of Biological Chemistry, 2013, 288, 11940-11948.	3.4	34
58	ALOX5 gene variants affect eicosanoid production and response to fish oil supplementation. Journal of Lipid Research, 2011, 52, 991-1003.	4.2	31
59	Identification of a Novel Mucin Gene <i>HCG22</i> Associated With Steroid-Induced Ocular Hypertension., 2015, 56, 2737.		28
60	Lower omental tâ€regulatory cell count is associated with higher fasting glucose and lower βâ€cell function in adults with obesity. Obesity, 2016, 24, 1274-1282.	3.0	28
61	APOE4 is associated with elevated blood lipids and lower levels of innate immune biomarkers in a tropical Amerindian subsistence population. ELife, 2021, 10, .	6.0	25
62	The Genetic Architecture of Noise-Induced Hearing Loss: Evidence for a Gene-by-Environment Interaction. G3: Genes, Genomes, Genetics, 2016, 6, 3219-3228.	1.8	24
63	Adult mouse hippocampal transcriptome changes associated with long-term behavioral and metabolic effects of gestational air pollution toxicity. Translational Psychiatry, 2020, 10, 218.	4.8	23
64	Genome-wide and gene-centric analyses of circulating myeloperoxidase levels in the charge and care consortia. Human Molecular Genetics, 2013, 22, 3381-3393.	2.9	22
65	Association of Chromosome 9p21 With Subsequent Coronary Heart Disease Events. Circulation Genomic and Precision Medicine, 2019, 12, e002471.	3.6	22
66	Functional analysis of 5-lipoxygenase promoter repeat variants. Human Molecular Genetics, 2009, 18, 4521-4529.	2.9	21
67	The Genetic Landscape of Hematopoietic Stem Cell Frequency in Mice. Stem Cell Reports, 2015, 5, 125-138.	4.8	21
68	Exposure to Nanoscale Particulate Matter from Gestation to Adulthood Impairs Metabolic Homeostasis in Mice. Scientific Reports, 2019, 9, 1816.	3.3	21
69	Genetic Determinants of Circulating Glycine Levels and Risk of Coronary Artery Disease. Journal of the American Heart Association, 2019, 8, e011922.	3.7	20
70	Apolipoprotein E4 genotype in combination with poor metabolic profile is associated with reduced cognitive performance in healthy postmenopausal women: implications for late onset Alzheimer's disease. Menopause, 2019, 26, 7-15.	2.0	19
71	Subsequent Event Risk in Individuals With Established Coronary Heart Disease. Circulation Genomic and Precision Medicine, 2019, 12, e002470.	3.6	17
72	Arachidonate 5-Lipoxygenase Gene Variants Affect Response to Fish Oil Supplementation by Healthy African Americans. Journal of Nutrition, 2012, 142, 1417-1428.	2.9	16

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73	The Genetic Architecture of Hearing Impairment in Mice: Evidence for Frequency-Specific Genetic Determinants. G3: Genes, Genomes, Genetics, 2015, 5, 2329-2339.	1.8	16
74	Nutrigenetic response to omega-3 fatty acids in obese asthmatics (NOOA): Rationale and methods. Contemporary Clinical Trials, 2013, 34, 326-335.	1.8	15
75	Habitual Diets Rich in Dark-Green Vegetables Are Associated with an Increased Response to ï‰-3 Fatty Acid Supplementation in Americans of African Ancestry. Journal of Nutrition, 2014, 144, 123-131.	2.9	15
76	Association of PLA2G4A with myocardial infarction is modulated by dietary PUFAs. American Journal of Clinical Nutrition, 2012, 95, 959-965.	4.7	14
77	Nonconventional genetic risk factors for cardiovascular disease. Current Atherosclerosis Reports, 2006, 8, 184-192.	4.8	13
78	Genetic and clinical markers of elevated liver fat content in overweight and obese hispanic children. Obesity, 2013, 21, E790-7.	3.0	12
79	Gene-Environment Interactions for Cardiovascular Disease. Current Atherosclerosis Reports, 2021, 23, 75.	4.8	12
80	Association of Factor V Leiden With Subsequent Atherothrombotic Events. Circulation, 2020, 142, 546-555.	1.6	11
81	Using Inbred Mouse Strains to Identify Genes for Complex Diseases. Frontiers in Bioscience - Landmark, 2006, 11, 1216.	3.0	10
82	A GWAS approach identifies Dapp1 as a determinant of air pollution-induced airway hyperreactivity. PLoS Genetics, 2019, 15, e1008528.	3.5	9
83	Dissecting the Genetic Architecture of Cystatin C in Diversity Outbred Mice. G3: Genes, Genomes, Genetics, 2020, 10, 2529-2541.	1.8	9
84	PNPLA3 Genotype, Arachidonic Acid Intake, and Unsaturated Fat Intake Influences Liver Fibrosis in Hispanic Youth with Obesity. Nutrients, 2021, 13, 1621.	4.1	8
85	Clinical Intervention to Reduce Dietary Sugar Does Not Affect Liver Fat in Latino Youth, Regardless of PNPLA3 Genotype: A Randomized Controlled Trial. Journal of Nutrition, 2022, 152, 1655-1665.	2.9	8
86	Allgrove syndrome in a Mexican American family is caused by an ancestral mutation derived from North Africa. Clinical Genetics, 2008, 73, 385-387.	2.0	7
87	CD52-targeted depletion by Alemtuzumab ameliorates allergic airway hyperreactivity and lung inflammation. Mucosal Immunology, 2021, 14, 899-911.	6.0	7
88	Improved Performance of Dynamic Measures of Insulin Response Over Surrogate Indices to Identify Genetic Contributors of Type 2 Diabetes: The GUARDIAN Consortium. Diabetes, 2016, 65, 2072-2080.	0.6	4
89	Near-roadway air pollution, immune cells and adipokines among obese young adults. Environmental Health, 2022, 21, 36.	4.0	4
90	Effect of ApoE4 Genotype on the Association Between Metabolic Phenotype and Subclinical Atherosclerosis in Postmenopausal Women. American Journal of Cardiology, 2019, 124, 1031-1037.	1.6	3

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91	Genome-Wide Association Analysis Identifies Dcc as an Essential Factor in the Innervation of the Peripheral Vestibular System in Inbred Mice. JARO - Journal of the Association for Research in Otolaryngology, 2016, 17, 417-431.	1.8	2
92	Noise Exposure and Distortion Product Otoacoustic Emission Suprathreshold Amplitudes: A Genome-Wide Association Study. Audiology and Neuro-Otology, 2021, 26, 1-9.	1.3	2
93	Genetic evidence for independent causal relationships between metabolic biomarkers and risk of coronary artery diseases. Journal of Lipid Research, 2021, 62, 100064.	4.2	1
94	Non-Conventional Genetic Risk Factors for Cardiovascular Disease. World Review of Nutrition and Dietetics, 2008, 98, 62-76.	0.3	0
95	Effect of Omegaâ€3 fatty acid supplementation and ALOX5 promoter variants on Lipid Profiles in Africanâ€Americans. FASEB Journal, 2009, 23, 724.3.	0.5	0