

Francois Cambien

List of Publications by Year in descending order

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145
papers

21,953
citations

14614

66
h-index

9311

143
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147
all docs

147
docs citations

147
times ranked

26518
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association analysis in dilated cardiomyopathy reveals two new players in systolic heart failure on chromosomes 3p25.1 and 22q11.23. <i>European Heart Journal</i> , 2021, 42, 2000-2011.	1.0	49
2	Natural genetic variation of the cardiac transcriptome in non-diseased donors and patients with dilated cardiomyopathy. <i>Genome Biology</i> , 2017, 18, 170.	3.8	70
3	Exome-wide association study reveals novel susceptibility genes to sporadic dilated cardiomyopathy. <i>PLoS ONE</i> , 2017, 12, e0172995.	1.1	92
4	Preservation Analysis of Macrophage Gene Coexpression Between Human and Mouse Identifies PARK2 as a Genetically Controlled Master Regulator of Oxidative Phosphorylation in Humans. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 3361-3371.	0.8	15
5	Role of lipid phosphate phosphatase 3 in human aortic endothelial cell function. <i>Cardiovascular Research</i> , 2016, 112, 702-713.	1.8	25
6	Contribution of Rare and Common Genetic Variants to Plasma Lipid Levels and Carotid Stiffness and Geometry. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 628-636.	5.1	21
7	Runs of Homozygosity: Association with Coronary Artery Disease and Gene Expression in Monocytes and Macrophages. <i>American Journal of Human Genetics</i> , 2015, 97, 228-237.	2.6	37
8	Meta-analysis of 65,734 Individuals Identifies TSPAN15 and SLC44A2 as Two Susceptibility Loci for Venous Thromboembolism. <i>American Journal of Human Genetics</i> , 2015, 96, 532-542.	2.6	222
9	SASH1, a new potential link between smoking and atherosclerosis. <i>Atherosclerosis</i> , 2015, 242, 571-579.	0.4	24
10	Adrenomedullin and Arterial Stiffness. <i>Circulation: Cardiovascular Genetics</i> , 2014, 7, 634-641.	5.1	12
11	Cigarette smoking reduces DNA methylation levels at multiple genomic loci but the effect is partially reversible upon cessation. <i>Epigenetics</i> , 2014, 9, 1382-1396.	1.3	285
12	Genome-Wide Association Study for Circulating Tissue Plasminogen Activator Levels and Functional Follow-Up Implicates Endothelial <i>STXBP5</i> and <i>STX2</i> . <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, 1093-1101.	1.1	43
13	DNA methylation and body-mass index: a genome-wide analysis. <i>Lancet, The</i> , 2014, 383, 1990-1998.	6.3	686
14	Allelic expression mapping across cellular lineages to establish impact of non-coding <i>scnp>SNP</scnp>s</i> . <i>Molecular Systems Biology</i> , 2014, 10, 754.	3.2	21
15	Human CalDAG-GEFI gene (<i>RASGRP2</i>) mutation affects platelet function and causes severe bleeding. <i>Journal of Experimental Medicine</i> , 2014, 211, 1349-1362.	4.2	117
16	A meta-analysis of genome-wide association studies identifies ORM1 as a novel gene controlling thrombin generation potential. <i>Blood</i> , 2014, 123, 777-785.	0.6	27
17	Single Nucleotide Polymorphisms with Cis-Regulatory Effects on Long Non-Coding Transcripts in Human Primary Monocytes. <i>PLoS ONE</i> , 2014, 9, e102612.	1.1	9
18	First case of a human <i>RASGRP2</i> mutation affecting Rap1 activation in platelets and causing severe bleeding.. <i>Journal of Cell Biology</i> , 2014, 206, 20610IA111.	2.3	0

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19	Large-scale association analysis identifies new risk loci for coronary artery disease. <i>Nature Genetics</i> , 2013, 45, 25-33.	9.4	1,439
20	Cytokine phenotype, genotype, and renal outcomes at cardiac surgery. <i>Cytokine</i> , 2013, 61, 275-284.	1.4	14
21	Genome-Wide Haplotype Analysis of Cis Expression Quantitative Trait Loci in Monocytes. <i>PLoS Genetics</i> , 2013, 9, e1003240.	1.5	53
22	GUESS-ing Polygenic Associations with Multiple Phenotypes Using a GPU-Based Evolutionary Stochastic Search Algorithm. <i>PLoS Genetics</i> , 2013, 9, e1003657.	1.5	58
23	Graphical Modeling of Gene Expression in Monocytes Suggests Molecular Mechanisms Explaining Increased Atherosclerosis in Smokers. <i>PLoS ONE</i> , 2013, 8, e50888.	1.1	36
24	Statistical colocalization of monocyte gene expression and genetic risk variants for type 1 diabetes. <i>Human Molecular Genetics</i> , 2012, 21, 2815-2824.	1.4	103
25	Long-range DNA looping and gene expression analyses identify DEXI as an autoimmune disease candidate gene. <i>Human Molecular Genetics</i> , 2012, 21, 322-333.	1.4	100
26	Inheritance of coronary artery disease in men: an analysis of the role of the Y chromosome. <i>Lancet, The</i> , 2012, 379, 915-922.	6.3	179
27	Interleukin-6 receptor pathways in coronary heart disease: a collaborative meta-analysis of 82 studies. <i>Lancet, The</i> , 2012, 379, 1205-1213.	6.3	668
28	Genome-wide association study for circulating levels of PAI-1 provides novel insights into its regulation. <i>Blood</i> , 2012, 120, 4873-4881.	0.6	90
29	Caution in Interpreting Results from Imputation Analysis When Linkage Disequilibrium Extends over a Large Distance: A Case Study on Venous Thrombosis. <i>PLoS ONE</i> , 2012, 7, e38538.	1.1	17
30	Comprehensive Exploration of the Effects of miRNA SNPs on Monocyte Gene Expression. <i>PLoS ONE</i> , 2012, 7, e45863.	1.1	8
31	Powerful Identification of Cis-regulatory SNPs in Human Primary Monocytes Using Allele-Specific Gene Expression. <i>PLoS ONE</i> , 2012, 7, e52260.	1.1	36
32	A genome-wide association study identifies two loci associated with heart failure due to dilated cardiomyopathy. <i>European Heart Journal</i> , 2011, 32, 1065-1076.	1.0	292
33	Influence of sex and genetic variability on expression of X-linked genes in human monocytes. <i>Genomics</i> , 2011, 98, 320-326.	1.3	23
34	The Choice of the Filtering Method in Microarrays Affects the Inference Regarding Dosage Compensation of the Active X-Chromosome. <i>PLoS ONE</i> , 2011, 6, e23956.	1.1	23
35	A pharmacogenetic analysis of determinants of hypertension and blood pressure response to angiotensin-converting enzyme inhibitor therapy in patients with vascular disease and healthy individuals. <i>Journal of Hypertension</i> , 2011, 29, 509-519.	0.3	47
36	Large-scale association analysis identifies 13 new susceptibility loci for coronary artery disease. <i>Nature Genetics</i> , 2011, 43, 333-338.	9.4	1,685

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37	A Genome-Wide Association Study Identifies <i>LIPA</i> as a Susceptibility Gene for Coronary Artery Disease. <i>Circulation: Cardiovascular Genetics</i> , 2011, 4, 403-412.	5.1	130
38	Common variant near the endothelin receptor type A (<i>EDNRA</i>) gene is associated with intracranial aneurysm risk. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 19707-19712.	3.3	100
39	Bayesian Detection of Expression Quantitative Trait Loci Hot Spots. <i>Genetics</i> , 2011, 189, 1449-1459.	1.2	70
40	Heritability, Weak Effects, and Rare Variants in Genomewide Association Studies. <i>Clinical Chemistry</i> , 2011, 57, 1263-1266.	1.5	5
41	Four Genetic Loci Influencing Electrocardiographic Indices of Left Ventricular Hypertrophy. <i>Circulation: Cardiovascular Genetics</i> , 2011, 4, 626-635.	5.1	28
42	Integrating Genome-Wide Genetic Variations and Monocyte Expression Data Reveals Trans-Regulated Gene Modules in Humans. <i>PLoS Genetics</i> , 2011, 7, e1002367.	1.5	126
43	Genetics of Venous Thrombosis: Insights from a New Genome Wide Association Study. <i>PLoS ONE</i> , 2011, 6, e25581.	1.1	127
44	A Follow-Up Study of a Genome-wide Association Scan Identifies a Susceptibility Locus for Venous Thrombosis on Chromosome 6p24.1. <i>American Journal of Human Genetics</i> , 2010, 86, 592-595.	2.6	57
45	Association of three polymorphisms selected from a genome-wide association study with coronary heart disease in the Tunisian population. <i>Journal of Thrombosis and Thrombolysis</i> , 2010, 29, 114-118.	1.0	6
46	A trans-acting locus regulates an anti-viral expression network and type 1 diabetes risk. <i>Nature</i> , 2010, 467, 460-464.	13.7	271
47	Genome-wide association study of intracranial aneurysm identifies three new risk loci. <i>Nature Genetics</i> , 2010, 42, 420-425.	9.4	262
48	Genetics and Beyond – The Transcriptome of Human Monocytes and Disease Susceptibility. <i>PLoS ONE</i> , 2010, 5, e10693.	1.1	539
49	Multiple marker approach to risk stratification in patients with stable coronary artery disease. <i>European Heart Journal</i> , 2010, 31, 3024-3031.	1.0	97
50	Genetic Association Study Identifies HSPB7 as a Risk Gene for Idiopathic Dilated Cardiomyopathy. <i>PLoS Genetics</i> , 2010, 6, e1001167.	1.5	110
51	Lack of Association Between the Trp719Arg Polymorphism in Kinesin-Like Protein-6 and Coronary Artery Disease in 19 Case-Control Studies. <i>Journal of the American College of Cardiology</i> , 2010, 56, 1552-1563.	1.2	84
52	Genetic Regulation of Serum Phytosterol Levels and Risk of Coronary Artery Disease. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 331-339.	5.1	141
53	Molecular genetic analysis of a human insulin-like growth factor 1 promoter P1 variation. <i>FASEB Journal</i> , 2009, 23, 1303-1313.	0.2	19
54	Identification and Functional Analyses of Molecular Haplotypes of the Human Osteoprotegerin Gene Promoter. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2009, 29, 1638-1643.	1.1	7

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55	Influence of cholesteryl ester transfer protein, peroxisome proliferator-activated receptor α , apolipoprotein E, and apolipoprotein A-I polymorphisms on high-density lipoprotein cholesterol, apolipoprotein A-I, lipoprotein A-I, and lipoprotein A-I:A-II concentrations: the Prospective Epidemiological Study of Myocardial Infarction study. <i>Metabolism: Clinical and Experimental</i> , 2009, 58, 283-289.	1.5	9
56	The impact of newly identified loci on coronary heart disease, stroke and total mortality in the MORGAM prospective cohorts. <i>Genetic Epidemiology</i> , 2009, 33, 237-246.	0.6	77
57	P-selectin gene polymorphisms and risk of coronary heart disease among Tunisians. <i>Journal of Thrombosis and Thrombolysis</i> , 2009, 28, 314-319.	1.0	8
58	Molecular and functional characterization of polymorphisms in the secreted phospholipase A2 group X gene: relevance to coronary artery disease. <i>Journal of Molecular Medicine</i> , 2009, 87, 723-733.	1.7	16
59	Lack of association between polymorphisms of the IL18R1 and IL18RAP genes and cardiovascular risk: the MORGAM Project. <i>BMC Medical Genetics</i> , 2009, 10, 44.	2.1	18
60	New susceptibility locus for coronary artery disease on chromosome 3q22.3. <i>Nature Genetics</i> , 2009, 41, 280-282.	9.4	440
61	Genome-wide haplotype association study identifies the SLC22A3-LPAL2-LPA gene cluster as a risk locus for coronary artery disease. <i>Nature Genetics</i> , 2009, 41, 283-285.	9.4	427
62	Genome-wide association of early-onset myocardial infarction with single nucleotide polymorphisms and copy number variants. <i>Nature Genetics</i> , 2009, 41, 334-341.	9.4	990
63	Osteopontin gene variation and cardio/cerebrovascular disease phenotypes. <i>Atherosclerosis</i> , 2009, 206, 209-215.	0.4	9
64	Lack of association of genetic variants in the LRP8 gene with familial and sporadic myocardial infarction. <i>Journal of Molecular Medicine</i> , 2008, 86, 1163-1170.	1.7	6
65	The renin-angiotensin-aldosterone system in cerebral small vessel disease. <i>Journal of Neurology</i> , 2008, 255, 993-1000.	1.8	31
66	Haplotypic analysis of tag SNPs of the interleukin-18 gene in relation to cardiovascular disease events: the MORGAM Project. <i>European Journal of Human Genetics</i> , 2008, 16, 1512-1520.	1.4	12
67	G/T Substitution in Intron 1 of the UNC13B Gene Is Associated With Increased Risk of Nephropathy in Patients With Type 1 Diabetes. <i>Diabetes</i> , 2008, 57, 2843-2850.	0.3	39
68	Repeated Replication and a Prospective Meta-Analysis of the Association Between Chromosome 9p21.3 and Coronary Artery Disease. <i>Circulation</i> , 2008, 117, 1675-1684.	1.6	356
69	Molecular investigation of the functional relevance of missense variants of ICAM-1. <i>Pharmacogenetics and Genomics</i> , 2008, 18, 1017-1019.	0.7	3
70	Genomewide Association Analysis of Coronary Artery Disease. <i>New England Journal of Medicine</i> , 2007, 357, 443-453.	13.9	1,865
71	European rational approach for the genetics of diabetic complications EURAGEDIC: patient populations and strategy. <i>Nephrology Dialysis Transplantation</i> , 2007, 23, 161-168.	0.4	30
72	Genetics of Cardiovascular Diseases. <i>Circulation</i> , 2007, 116, 1714-1724.	1.6	86

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73	SAH gene variants are associated with obesity-related hypertension in Caucasians: the PEGASE Study. <i>Journal of Hypertension</i> , 2007, 25, 557-564.	0.3	11
74	Neutrophil elastase gene variation and coronary heart disease. <i>Pharmacogenetics and Genomics</i> , 2007, 17, 629-637.	0.7	2
75	Patterns of Alcohol Consumption and Cardiovascular Risk in Northern Ireland and France. <i>Annals of Epidemiology</i> , 2007, 17, S75-S80.	0.9	6
76	Coronary heart disease aetiology: associations and causality. <i>Comptes Rendus - Biologies</i> , 2007, 330, 299-305.	0.1	1
77	Differential haplotypic expression of the interleukin-18 gene. <i>European Journal of Human Genetics</i> , 2007, 15, 856-863.	1.4	29
78	Lack of association between complement factor H polymorphisms and coronary artery disease or myocardial infarction. <i>Journal of Molecular Medicine</i> , 2007, 85, 771-775.	1.7	19
79	Polymorphisms in 33 inflammatory genes and risk of myocardial infarction—a system genetics approach. <i>Journal of Molecular Medicine</i> , 2007, 85, 1271-1280.	1.7	6
80	B-Type Natriuretic Peptide and the Risk of Cardiovascular Events and Death in Patients With Stable Angina. <i>Journal of the American College of Cardiology</i> , 2006, 47, 552-558.	1.2	99
81	Gene Polymorphisms of the Renin-Angiotensin System and Early Development of Hypertension. <i>American Journal of Hypertension</i> , 2006, 19, 837-842.	1.0	40
82	Prognostic value of tissue inhibitor of metalloproteinase-1 for cardiovascular death among patients with cardiovascular disease: results from the AtheroGene study. <i>European Heart Journal</i> , 2006, 27, 150-156.	1.0	69
83	Cytokine Polymorphisms Associated With Carotid Intima-Media Thickness in Stroke Patients. <i>Stroke</i> , 2006, 37, 1691-1696.	1.0	59
84	Analysis of 14 Candidate Genes for Diabetic Nephropathy on Chromosome 3q in European Populations: Strongest Evidence for Association With a Variant in the Promoter Region of the Adiponectin Gene. <i>Diabetes</i> , 2006, 55, 3166-3174.	0.3	74
85	Comparative Impact of Multiple Biomarkers and N-Terminal Pro-Brain Natriuretic Peptide in the Context of Conventional Risk Factors for the Prediction of Recurrent Cardiovascular Events in the Heart Outcomes Prevention Evaluation (HOPE) Study. <i>Circulation</i> , 2006, 114, 201-208.	1.6	236
86	Haplotypes of the Caspase-1 Gene, Plasma Caspase-1 Levels, and Cardiovascular Risk. <i>Circulation Research</i> , 2006, 99, 102-108.	2.0	26
87	A polymorphism in the endothelin-A receptor gene is linked to baroreflex sensitivity. <i>Journal of Hypertension</i> , 2005, 23, 2019-2026.	0.3	23
88	Atherosclerosis: From Genetic Polymorphisms to System Genetics. <i>Cardiovascular Toxicology</i> , 2005, 5, 143-152.	1.1	12
89	Asymmetric Dimethylarginine and the Risk of Cardiovascular Events and Death in Patients With Coronary Artery Disease. <i>Circulation Research</i> , 2005, 97, e53-9.	2.0	330
90	Renin-angiotensin-aldosterone system in brain infarction and vascular death. <i>Annals of Neurology</i> , 2005, 58, 131-138.	2.8	40

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91	Coronary heart disease and polymorphisms in genes affecting lipid metabolism and inflammation. <i>Current Atherosclerosis Reports</i> , 2005, 7, 188-195.	2.0	18
92	Genetic Analysis of the Interleukin-18 System Highlights the Role of the Interleukin-18 Gene in Cardiovascular Disease. <i>Circulation</i> , 2005, 112, 643-650.	1.6	205
93	Analysis of N-terminal-pro-brain natriuretic peptide and C-reactive protein for risk stratification in stable and unstable coronary artery disease: results from the AtheroGene study. <i>European Heart Journal</i> , 2005, 26, 241-249.	1.0	90
94	Adverse Associations Between CX3CR1 Polymorphisms and Risk of Cardiovascular or Cerebrovascular Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2005, 25, 847-853.	1.1	44
95	Glutathione Peroxidase-1 and Homocysteine for Cardiovascular Risk Prediction. <i>Journal of the American College of Cardiology</i> , 2005, 45, 1631-1637.	1.2	103
96	Platelet-activating factor-acetylhydrolase and PAF-receptor gene haplotypes in relation to future cardiovascular event in patients with coronary artery disease. <i>Human Molecular Genetics</i> , 2004, 13, 1341-1351.	1.4	91
97	In-Depth Haplotype Analysis of ABCA1 Gene Polymorphisms in Relation to Plasma ApoA1 Levels and Myocardial Infarction. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2004, 24, 775-781.	1.1	96
98	MORGAM (an international pooling of cardiovascular cohorts). <i>International Journal of Epidemiology</i> , 2004, 34, 21-27.	0.9	105
99	Polymorphism R92Q of the tumour necrosis factor receptor 1 gene is associated with myocardial infarction and carotid intima-media thickness – The ECTIM, AXA, EVA and GENIC Studies. <i>European Journal of Human Genetics</i> , 2004, 12, 213-219.	1.4	45
100	Impact of pathogen burden in patients with coronary artery disease in relation to systemic inflammation and variation in genes encoding cytokines. <i>American Journal of Cardiology</i> , 2003, 92, 515-521.	0.7	100
101	Polymorphisms of genes of the cardiac calcineurin pathway and cardiac hypertrophy. <i>European Journal of Human Genetics</i> , 2003, 11, 659-664.	1.4	27
102	Common genetic variation of the cholesteryl ester transfer protein gene strongly predicts future cardiovascular death in patients with coronary artery disease. <i>Journal of the American College of Cardiology</i> , 2003, 41, 1983-1989.	1.2	89
103	Plasma Concentrations and Genetic Variation of Matrix Metalloproteinase 9 and Prognosis of Patients With Cardiovascular Disease. <i>Circulation</i> , 2003, 107, 1579-1585.	1.6	692
104	Glutathione Peroxidase 1 Activity and Cardiovascular Events in Patients with Coronary Artery Disease. <i>New England Journal of Medicine</i> , 2003, 349, 1605-1613.	13.9	548
105	Long-Term Renoprotective Effects of Losartan in Diabetic Nephropathy: Interaction with ACE insertion/deletion genotype?. <i>Diabetes Care</i> , 2003, 26, 1501-1506.	4.3	60
106	Interleukin-18 and the Risk of Coronary Heart Disease in European Men. <i>Circulation</i> , 2003, 108, 2453-2459.	1.6	317
107	The ACE Gene I/D Polymorphism Is Not Associated With the Blood Pressure and Cardiovascular Benefits of ACE Inhibition. <i>Hypertension</i> , 2003, 42, 297-303.	1.3	129
108	Family history, longevity, and risk of coronary heart disease: the PRIME Study. <i>International Journal of Epidemiology</i> , 2003, 32, 71-77.	0.9	37

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109	No Physical Activity ?? CETP 1b.-629 Interaction Effects on Lipid Profile. <i>Medicine and Science in Sports and Exercise</i> , 2003, 35, 1124-1129.	0.2	17
110	Automated detection of informative combined effects in genetic association studies of complex traits. <i>Genome Research</i> , 2003, 13, 1952-60.	2.4	14
111	Automated Detection of Informative Combined Effects in Genetic Association Studies of Complex Traits. <i>Genome Research</i> , 2003, 13, 1952-1960.	2.4	38
112	Heterogeneity of linkage disequilibrium in human genes has implications for association studies of common diseases. <i>Human Molecular Genetics</i> , 2002, 11, 419-429.	1.4	54
113	Specific haplotypes of the P-selectin gene are associated with myocardial infarction. <i>Human Molecular Genetics</i> , 2002, 11, 2015-2023.	1.4	161
114	Interleukin-18 Is a Strong Predictor of Cardiovascular Death in Stable and Unstable Angina. <i>Circulation</i> , 2002, 106, 24-30.	1.6	534
115	Polymorphisms of the β_2 -adrenoceptor (ADRB2) gene and essential hypertension: the ECTIM and PEGASE studies. <i>Journal of Hypertension</i> , 2002, 20, 229-235.	0.3	65
116	High expressor paraoxonase PON1 gene promoter polymorphisms are associated with reduced risk of vascular disease in younger coronary patients. <i>Atherosclerosis</i> , 2002, 161, 463-467.	0.4	40
117	Physical Activity May Modulate Effects of ApoE Genotype on Lipid Profile. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2002, 22, 133-140.	1.1	82
118	Renoprotective effects of losartan in diabetic nephropathy: Interaction with ACE insertion/deletion genotype?. <i>Kidney International</i> , 2002, 62, 192-198.	2.6	30
119	Endothelin gene variants and aortic and cardiac structure in never-treated hypertensives. <i>American Journal of Hypertension</i> , 2001, 14, 755-760.	1.0	50
120	Angiotensin II type 1 receptor α_1 153A/G and 1166A/C gene polymorphisms and increase in aortic stiffness with age in hypertensive subjects. <i>Journal of Hypertension</i> , 2001, 19, 407-413.	0.3	108
121	Interleukin-6 gene polymorphisms and susceptibility to myocardial infarction: the ECTIM study. <i>Journal of Molecular Medicine</i> , 2001, 79, 300-305.	1.7	155
122	Extensive association analysis between the CETP gene and coronary heart disease phenotypes reveals several putative functional polymorphisms and gene-environment interaction. <i>Genetic Epidemiology</i> , 2000, 19, 64-80.	0.6	205
123	Identification of two polymorphisms in the early growth response protein-1 gene: possible association with lipid variables. <i>Journal of Molecular Medicine</i> , 2000, 78, 81-86.	1.7	14
124	Polymorphisms in the genes encoding platelet-derived growth factor A and α_2 receptor. <i>Journal of Molecular Medicine</i> , 2000, 78, 287-292.	1.7	10
125	Lack of association between polymorphisms of eight candidate genes and idiopathic dilated cardiomyopathy. <i>Journal of the American College of Cardiology</i> , 2000, 35, 29-35.	1.2	74
126	Effects of Alcohol on Lipids and Lipoprotein Metabolism. <i>Medical Science Symposia Series</i> , 2000, , 11-22.	0.0	0

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127	Progression of diabetic nephropathy in normotensive type 1 diabetic patients. <i>Kidney International</i> , 1999, 56, S101-S105.	2.6	80
128	Sequence Diversity in 36 Candidate Genes for Cardiovascular Disorders. <i>American Journal of Human Genetics</i> , 1999, 65, 183-191.	2.6	152
129	Polymorphisms of the endothelin-A and -B receptor genes in relation to blood pressure and myocardial infarction The etude cas-t�moins sur l'infarctus du myocarde (ECTIM) study. <i>American Journal of Hypertension</i> , 1999, 12, 304-310.	1.0	59
130	Sequence polymorphisms in the apolipoprotein(a) gene and their association with lipoprotein(a) levels and myocardial infarction. The ECTIM Study. <i>Atherosclerosis</i> , 1999, 144, 323-333.	0.4	55
131	Characterization of a Unique Genetic Variant in the β 1-adrenoceptor Gene and Evaluation of its Role in Idiopathic Dilated Cardiomyopathy. <i>Journal of Molecular and Cellular Cardiology</i> , 1999, 31, 1025-1032.	0.9	103
132	Angiotensin converting enzyme gene polymorphism and ACE inhibition in diabetic nephropathy. <i>Kidney International</i> , 1998, 53, 1002-1006.	2.6	91
133	Gene polymorphisms of the renin-angiotensin system in relation to hypertension and parental history of myocardial infarction and stroke. <i>Journal of Hypertension</i> , 1998, 16, 37-44.	0.3	114
134	Nitric oxide synthase gene polymorphisms, blood pressure and aortic stiffness in normotensive and hypertensive subjects. <i>Journal of Hypertension</i> , 1998, 16, 31-35.	0.3	155
135	New polymorphisms of the angiotensin II type 1 receptor gene and their associations with myocardial infarction and blood pressure. <i>Journal of Hypertension</i> , 1998, 16, 1443-1447.	0.3	72
136	Identification of two polymorphisms in the promoter of the microsomal triglyceride transfer protein (MTP) gene: lack of association with lipoprotein profiles. <i>Journal of Lipid Research</i> , 1998, 39, 2432-2435.	2.0	26
137	Distribution of apolipoprotein E between apo B- and non apo B-containing lipoproteins according to apo E phenotype. <i>Atherosclerosis</i> , 1997, 131, 257-262.	0.4	16
138	Coronary heart disease and genetics an epidemiologist's view. <i>Trends in Molecular Medicine</i> , 1997, 3, 197-203.	2.6	20
139	The Gln/Arg polymorphism of human paraoxonase (PON 192) is not related to myocardial infarction in the ECTIM Study. <i>Atherosclerosis</i> , 1996, 126, 299-303.	0.4	163
140	The genetic contribution to the onset of acute coronary heart disease. <i>Developments in Cardiovascular Medicine</i> , 1996, , 111-133.	0.1	0
141	Cardiovascular risk factors and alcohol consumption in France and Northern Ireland. <i>Atherosclerosis</i> , 1995, 115, 225-232.	0.4	78
142	The angiotensin-converting enzyme (ACE) genetic polymorphism: its relationship with plasma ACE level and myocardial infarction. <i>Clinical Genetics</i> , 1994, 46, 94-101.	1.0	46
143	Renin-angiotensin system genes as candidate genes in cardiovascular diseases. <i>Trends in Cardiovascular Medicine</i> , 1993, 3, 250-258.	2.3	8
144	Deletion polymorphism in the gene for angiotensin-converting enzyme is a potent risk factor for myocardial infarction. <i>Nature</i> , 1992, 359, 641-644.	13.7	1,880

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145	CORONARY HEART DISEASE IN MIDDLE-AGED FRENCHMEN. <i>Lancet, The</i> , 1980, 315, 1346-1350.	6.3	77