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 g-index
147 all docs	147 docs citations	147 times ranked	26518 citing authors

#	Article	IF	Citations
1	Deletion polymorphism in the gene for angiotensin-converting enzyme is a potent risk factor for myocardial infarction. Nature, 1992, 359, 641-644.	13.7	1,880
2	Genomewide Association Analysis of Coronary Artery Disease. New England Journal of Medicine, 2007, 357, 443-453.	13.9	1,865
3	Large-scale association analysis identifies 13 new susceptibility loci for coronary artery disease. Nature Genetics, 2011, 43, 333-338.	9.4	1,685
4	Large-scale association analysis identifies new risk loci for coronary artery disease. Nature Genetics, 2013, 45, 25-33.	9.4	1,439
5	Genome-wide association of early-onset myocardial infarction with single nucleotide polymorphisms and copy number variants. Nature Genetics, 2009, 41, 334-341.	9.4	990
6	Plasma Concentrations and Genetic Variation of Matrix Metalloproteinase 9 and Prognosis of Patients With Cardiovascular Disease. Circulation, 2003, 107, 1579-1585.	1.6	692
7	DNA methylation and body-mass index: a genome-wide analysis. Lancet, The, 2014, 383, 1990-1998.	6.3	686
8	Interleukin-6 receptor pathways in coronary heart disease: a collaborative meta-analysis of 82 studies. Lancet, The, 2012, 379, 1205-1213.	6.3	668
9	Glutathione Peroxidase 1 Activity and Cardiovascular Events in Patients with Coronary Artery Disease. New England Journal of Medicine, 2003, 349, 1605-1613.	13.9	548
10	Genetics and Beyond â€“ The Transcriptome of Human Monocytes and Disease Susceptibility. PLoS ONE, 2010, 5, e10693.	1.1	539
11	Interleukin-18 Is a Strong Predictor of Cardiovascular Death in Stable and Unstable Angina. Circulation, 2002, 106, 24-30.	1.6	534
12	New susceptibility locus for coronary artery disease on chromosome 3q22.3. Nature Genetics, 2009, 41, 280-282.	9.4	440
13	Genome-wide haplotype association study identifies the SLC22A3-LPAL2-LPA gene cluster as a risk locus for coronary artery disease. Nature Genetics, 2009, 41, 283-285.	9.4	427
14	Repeated Replication and a Prospective Meta-Analysis of the Association Between Chromosome 9p21.3 and Coronary Artery Disease. Circulation, 2008, 117, 1675-1684.	1.6	356
15	Asymmetric Dimethylarginine and the Risk of Cardiovascular Events and Death in Patients With Coronary Artery Disease. Circulation Research, 2005, 97, e53-9.	2.0	330
16	Interleukin-18 and the Risk of Coronary Heart Disease in European Men. Circulation, 2003, 108, 2453-2459.	1.6	317
17	A genome-wide association study identifies two loci associated with heart failure due to dilated cardiomyopathy. European Heart Journal, 2011, 32, 1065-1076.	1.0	292
18	Cigarette smoking reduces DNA methylation levels at multiple genomic loci but the effect is partially reversible upon cessation. Epigenetics, 2014, 9, 1382-1396.	1.3	285

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19	A trans-acting locus regulates an anti-viral expression network and type 1 diabetes risk. Nature, 2010, 467, 460-464.	13.7	271
20	Genome-wide association study of intracranial aneurysm identifies three new risk loci. Nature Genetics, 2010, 42, 420-425.	9.4	262
21	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. Circulation, 2006, 114, 201-208.	1.6	236
22	Meta-analysis of 65,734 Individuals Identifies TSPAN15 and SLC44A2 as Two Susceptibility Loci for Venous Thromboembolism. American Journal of Human Genetics, 2015, 96, 532-542.	2.6	222
23	Extensive association analysis between the CETP gene and coronary heart disease phenotypes reveals several putative functional polymorphisms and gene-environment interaction. Genetic Epidemiology, 2000, 19, 64-80.	0.6	205
24	Genetic Analysis of the Interleukin-18 System Highlights the Role of the Interleukin-18 Gene in Cardiovascular Disease. Circulation, 2005, 112, 643-650.	1.6	205
25	Inheritance of coronary artery disease in men: an analysis of the role of the Y chromosome. Lancet, The, 2012, 379, 915-922.	6.3	179
26	The Gln/Arg polymorphism of human paraoxonase (PON 192) is not related to myocardial infarction in the ECTIM Study. Atherosclerosis, 1996, 126, 299-303.	0.4	163
27	Specific haplotypes of the P-selectin gene are associated with myocardial infarction. Human Molecular Genetics, 2002, 11, 2015-2023.	1.4	161
28	Nitric oxide synthase gene polymorphisms, blood pressure and aortic stiffness in normotensive and hypertensive subjects. Journal of Hypertension, 1998, 16, 31-35.	0.3	155
29	Interleukin-6 gene polymorphisms and susceptibility to myocardial infarction: the ECTIM study. Journal of Molecular Medicine, 2001, 79, 300-305.	1.7	155
30	Sequence Diversity in 36 Candidate Genes for Cardiovascular Disorders. American Journal of Human Genetics, 1999, 65, 183-191.	2.6	152
31	Genetic Regulation of Serum Phytosterol Levels and Risk of Coronary Artery Disease. Circulation: Cardiovascular Genetics, 2010, 3, 331-339.	5.1	141
32	A Genome-Wide Association Study Identifies LIPA as a Susceptibility Gene for Coronary Artery Disease. Circulation: Cardiovascular Genetics, 2011, 4, 403-412.	5.1	130
33	The ACE Gene I/D Polymorphism Is Not Associated With the Blood Pressure and Cardiovascular Benefits of ACE Inhibition. Hypertension, 2003, 42, 297-303.	1.3	129
34	Genetics of Venous Thrombosis: Insights from a New Genome Wide Association Study. PLoS ONE, 2011, 6, e25581.	1.1	127
35	Integrating Genome-Wide Genetic Variations and Monocyte Expression Data Reveals Trans-Regulated Gene Modules in Humans. PLoS Genetics, 2011, 7, e1002367.	1.5	126
36	Human CalDAG-GEFI gene (RASGRP2) mutation affects platelet function and causes severe bleeding. Journal of Experimental Medicine, 2014, 211, 1349-1362.	4.2	117

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37	Gene polymorphisms of the renin-angiotensin system in relation to hypertension and parental history of myocardial infarction and stroke. Journal of Hypertension, 1998, 16, 37-44.	0.3	114
38	Genetic Association Study Identifies HSPB7 as a Risk Gene for Idiopathic Dilated Cardiomyopathy. PLoS Genetics, 2010, 6, e1001167.	1.5	110
39	Angiotensin II type 1 receptorâ˜153A/G and 1166A/C gene polymorphisms and increase in aortic stiffness with age in hypertensive subjects. Journal of Hypertension, 2001, 19, 407-413.	0.3	108
40	MORGAM (an international pooling of cardiovascular cohorts). International Journal of Epidemiology, 2004, 34, 21-27.	0.9	105
41	Characterization of a Unique Genetic Variant in the Î²1-adrenoceptor Gene and Evaluation of its Role in Idiopathic Dilated Cardiomyopathy. Journal of Molecular and Cellular Cardiology, 1999, 31, 1025-1032.	0.9	103
42	Glutathione Peroxidase-1 and Homocysteine for Cardiovascular Risk Prediction. Journal of the American College of Cardiology, 2005, 45, 1631-1637.	1.2	103
43	Statistical colocalization of monocyte gene expression and genetic risk variants for type 1 diabetes. Human Molecular Genetics, 2012, 21, 2815-2824.	1.4	103
44	Impact of pathogen burden in patients with coronary artery disease in relation to systemic inflammation and variation in genes encoding cytokines. American Journal of Cardiology, 2003, 92, 515-521.	0.7	100
45	Common variant near the endothelin receptor type A (EDNRA) gene is associated with intracranial aneurysm risk. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 19707-19712.	3.3	100
46	Long-range DNA looping and gene expression analyses identify DEXI as an autoimmune disease candidate gene. Human Molecular Genetics, 2012, 21, 322-333.	1.4	100
47	B-Type Natriuretic Peptide and the Risk of Cardiovascular Events and Death in Patients With Stable Angina. Journal of the American College of Cardiology, 2006, 47, 552-558.	1.2	99
48	Multiple marker approach to risk stratification in patients with stable coronary artery disease. European Heart Journal, 2010, 31, 3024-3031.	1.0	97
49	In-Depth Haplotype Analysis of ABCA1 Gene Polymorphisms in Relation to Plasma ApoA1 Levels and Myocardial Infarction. Arteriosclerosis, Thrombosis, and Vascular Biology, 2004, 24, 775-781.	1.1	96
50	Exome-wide association study reveals novel susceptibility genes to sporadic dilated cardiomyopathy. PLoS ONE, 2017, 12, e0172995.	1.1	92
51	Angiotensin converting enzyme gene polymorphism and ACE inhibition in diabetic nephropathy. Kidney International, 1998, 53, 1002-1006.	2.6	91
52	Platelet-activating factor-acetylhydrolase and PAF-receptor gene haplotypes in relation to future cardiovascular event in patients with coronary artery disease. Human Molecular Genetics, 2004, 13, 1341-1351.	1.4	91
53	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. European Heart Journal, 2005, 26, 241-249.	1.0	90
54	Genome-wide association study for circulating levels of PAI-1 provides novel insights into its regulation. Blood, 2012, 120, 4873-4881.	0.6	90

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55	Common genetic variation of the cholesteryl ester transfer protein gene strongly predicts future cardiovascular death in patients with coronary artery disease. Journal of the American College of Cardiology, 2003, 41, 1983-1989.	1.2	89
56	Genetics of Cardiovascular Diseases. Circulation, 2007, 116, 1714-1724.	1.6	86
57	Lack of Association Between the Trp719Arg Polymorphism in Kinesin-Like Protein-6 and Coronary Artery Disease in 19 Case-Control Studies. Journal of the American College of Cardiology, 2010, 56, 1552-1563.	1.2	84
58	Physical Activity May Modulate Effects ofApoEGenotype on Lipid Profile. Arteriosclerosis, Thrombosis, and Vascular Biology, 2002, 22, 133-140.	1.1	82
59	Progression of diabetic nephropathy in normotensive type 1 diabetic patients. Kidney International, 1999, 56, S101-S105.	2.6	80
60	Cardiovascular risk factors and alcohol consumption in France and Northern Ireland. Atherosclerosis, 1995, 115, 225-232.	0.4	78
61	CORONARY HEART DISEASE IN MIDDLE-AGED FRENCHMEN. Lancet, The, 1980, 315, 1346-1350.	6.3	77
62	The impact of newly identified loci on coronary heart disease, stroke and total mortality in the MORGAM prospective cohorts. Genetic Epidemiology, 2009, 33, 237-246.	0.6	77
63	Lack of association between polymorphisms of eight candidate genes and idiopathic dilated cardiomyopathy. Journal of the American College of Cardiology, 2000, 35, 29-35.	1.2	74
64	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. Diabetes, 2006, 55, 3166-3174.	0.3	74
65	New polymorphisms of the angiotensin II type 1 receptor gene and their associations with myocardial infarction and blood pressure. Journal of Hypertension, 1998, 16, 1443-1447.	0.3	72
66	Bayesian Detection of Expression Quantitative Trait Loci Hot Spots. Genetics, 2011, 189, 1449-1459.	1.2	70
67	Natural genetic variation of the cardiac transcriptome in non-diseased donors and patients with dilated cardiomyopathy. Genome Biology, 2017, 18, 170.	3.8	70
68	Prognostic value of tissue inhibitor of metalloproteinase-1 for cardiovascular death among patients with cardiovascular disease: results from the AtheroGene study. European Heart Journal, 2006, 27, 150-156.	1.0	69
69	Polymorphisms of the Î²2-adrenoceptor (ADRB2) gene and essential hypertension: the ECTIM and PEGASE studies. Journal of Hypertension, 2002, 20, 229-235.	0.3	65
70	Long-Term Renoprotective Effects of Losartan in Diabetic Nephropathy: Interaction with ACE insertion/deletion genotype?. Diabetes Care, 2003, 26, 1501-1506.	4.3	60
71	Polymorphisms of the endothelin-A and -B receptor genes in relation to blood pressure and myocardial infarction The etude cas-tÃ©moins sur l'infarctus du myocarde (ECTIM) study. American Journal of Hypertension, 1999, 12, 304-310.	1.0	59
72	Cytokine Polymorphisms Associated With Carotid Intima-Media Thickness in Stroke Patients. Stroke, 2006, 37, 1691-1696.	1.0	59

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73	GUESS-ing Polygenic Associations with Multiple Phenotypes Using a GPU-Based Evolutionary Stochastic Search Algorithm. PLoS Genetics, 2013, 9, e1003657.	1.5	58
74	A Follow-Up Study of a Genome-wide Association Scan Identifies a Susceptibility Locus for Venous Thrombosis on Chromosome 6p24.1. American Journal of Human Genetics, 2010, 86, 592-595.	2.6	57
75	Sequence polymorphisms in the apolipoprotein(a) gene and their association with lipoprotein(a) levels and myocardial infarction. The ECTIM Study. Atherosclerosis, 1999, 144, 323-333.	0.4	55
76	Heterogeneity of linkage disequilibrium in human genes has implications for association studies of common diseases. Human Molecular Genetics, 2002, 11, 419-429.	1.4	54
77	Genome-Wide Haplotype Analysis of Cis Expression Quantitative Trait Loci in Monocytes. PLoS Genetics, 2013, 9, e1003240.	1.5	53
78	Endothelin gene variants and aortic and cardiac structure in never-treated hypertensives. American Journal of Hypertension, 2001, 14, 755-760.	1.0	50
79	Genome-wide association analysis in dilated cardiomyopathy reveals two new players in systolic heart failure on chromosomes 3p25.1 and 22q11.23. European Heart Journal, 2021, 42, 2000-2011.	1.0	49
80	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. Journal of Hypertension, 2011, 29, 509-519.	0.3	47
81	The angiotensinâ€converting enzyme (ACE) genetic polymorphism: its relationship with plasma ACE level and myocardial infarction. Clinical Genetics, 1994, 46, 94-101.	1.0	46
82	Polymorphism R92Q of the tumour necrosis factor receptor 1 gene is associated with myocardial infarction and carotid intima-media thickness â€“ The ECTIM, AXA, EVA and GENIC Studies. European Journal of Human Genetics, 2004, 12, 213-219.	1.4	45
83	Adverse Associations Between CX3CR1 Polymorphisms and Risk of Cardiovascular or Cerebrovascular Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2005, 25, 847-853.	1.1	44
84	Genome-Wide Association Study for Circulating Tissue Plasminogen Activator Levels and Functional Follow-Up Implicates Endothelial STXBP5 and STX2. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 1093-1101.	1.1	43
85	High expressor paraoxonase PON1 gene promoter polymorphisms are associated with reduced risk of vascular disease in younger coronary patients. Atherosclerosis, 2002, 161, 463-467.	0.4	40
86	Renin-angiotensin-aldosterone system in brain infarction and vascular death. Annals of Neurology, 2005, 58, 131-138.	2.8	40
87	Gene Polymorphisms of the Renin-Angiotensin System and Early Development of Hypertension. American Journal of Hypertension, 2006, 19, 837-842.	1.0	40
88	G/T Substitution in Intron 1 of the UNC13B Gene Is Associated With Increased Risk of Nephropathy in Patients With Type 1 Diabetes. Diabetes, 2008, 57, 2843-2850.	0.3	39
89	Automated Detection of Informative Combined Effects in Genetic Association Studies of Complex Traits. Genome Research, 2003, 13, 1952-1960.	2.4	38
90	Family history, longevity, and risk of coronary heart disease: the PRIME Study. International Journal of Epidemiology, 2003, 32, 71-77.	0.9	37

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91	Runs of Homozygosity: Association with Coronary Artery Disease and Gene Expression in Monocytes and Macrophages. American Journal of Human Genetics, 2015, 97, 228-237.	2.6	37
92	Graphical Modeling of Gene Expression in Monocytes Suggests Molecular Mechanisms Explaining Increased Atherosclerosis in Smokers. PLoS ONE, 2013, 8, e50888.	1.1	36
93	Powerful Identification of Cis-regulatory SNPs in Human Primary Monocytes Using Allele-Specific Gene Expression. PLoS ONE, 2012, 7, e52260.	1.1	36
94	The renin-angiotensin-aldosterone system in cerebral small vessel disease. Journal of Neurology, 2008, 255, 993-1000.	1.8	31
95	Renoprotective effects of losartan in diabetic nephropathy: Interaction with ACE insertion/deletion genotype?. Kidney International, 2002, 62, 192-198.	2.6	30
96	European rational approach for the genetics of diabetic complications EURAGEDIC: patient populations and strategy. Nephrology Dialysis Transplantation, 2007, 23, 161-168.	0.4	30
97	Differential haplotypic expression of the interleukin-18 gene. European Journal of Human Genetics, 2007, 15, 856-863.	1.4	29
98	Four Genetic Loci Influencing Electrocardiographic Indices of Left Ventricular Hypertrophy. Circulation: Cardiovascular Genetics, 2011, 4, 626-635.	5.1	28
99	Polymorphisms of genes of the cardiac calcineurin pathway and cardiac hypertrophy. European Journal of Human Genetics, 2003, 11, 659-664.	1.4	27
100	A meta-analysis of genome-wide association studies identifies ORM1 as a novel gene controlling thrombin generation potential. Blood, 2014, 123, 777-785.	0.6	27
101	Haplotypes of the Caspase-1 Gene, Plasma Caspase-1 Levels, and Cardiovascular Risk. Circulation Research, 2006, 99, 102-108.	2.0	26
102	Identification of two polymorphisms in the promoter of the microsomal triglyceride transfer protein (MTP) gene: lack of association with lipoprotein profiles. Journal of Lipid Research, 1998, 39, 2432-2435.	2.0	26
103	Role of lipid phosphate phosphatase 3 in human aortic endothelial cell function. Cardiovascular Research, 2016, 112, 702-713.	1.8	25
104	SASH1, a new potential link between smoking and atherosclerosis. Atherosclerosis, 2015, 242, 571-579.	0.4	24
105	A polymorphism in the endothelin-A receptor gene is linked to baroreflex sensitivity. Journal of Hypertension, 2005, 23, 2019-2026.	0.3	23
106	Influence of sex and genetic variability on expression of X-linked genes in human monocytes. Genomics, 2011, 98, 320-326.	1.3	23
107	The Choice of the Filtering Method in Microarrays Affects the Inference Regarding Dosage Compensation of the Active X-Chromosome. PLoS ONE, 2011, 6, e23956.	1.1	23
108	Allelic expression mapping across cellular lineages to establish impact of nonâ€coding <scp>SNP</scp>s. Molecular Systems Biology, 2014, 10, 754.	3.2	21

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109	Contribution of Rare and Common Genetic Variants to Plasma Lipid Levels and Carotid Stiffness and Geometry. Circulation: Cardiovascular Genetics, 2015, 8, 628-636.	5.1	21
110	Coronary heart disease and genetics an epidemiologist's view. Trends in Molecular Medicine, 1997, 3, 197-203.	2.6	20
111	Lack of association between complement factor H polymorphisms and coronary artery disease or myocardial infarction. Journal of Molecular Medicine, 2007, 85, 771-775.	1.7	19
112	Molecular genetic analysis of a human insulinâ€like growth factor 1 promoter P1 variation. FASEB Journal, 2009, 23, 1303-1313.	0.2	19
113	Coronary heart disease and polymorphisms in genes affecting lipid metabolism and inflammation. Current Atherosclerosis Reports, 2005, 7, 188-195.	2.0	18
114	Lack of association between polymorphisms of the IL18R1 and IL18RAP genes and cardiovascular risk: the MORGAM Project. BMC Medical Genetics, 2009, 10, 44.	2.1	18
115	No Physical Activity ?? CETP 1b.-629 Interaction Effects on Lipid Profile. Medicine and Science in Sports and Exercise, 2003, 35, 1124-1129.	0.2	17
116	Caution in Interpreting Results from Imputation Analysis When Linkage Disequilibrium Extends over a Large Distance: A Case Study on Venous Thrombosis. PLoS ONE, 2012, 7, e38538.	1.1	17
117	Distribution of apolipoprotein E between apo B- and non apo B-containing lipoproteins according to apo E phenotype. Atherosclerosis, 1997, 131, 257-262.	0.4	16
118	Molecular and functional characterization of polymorphisms in the secreted phospholipase A2 group X gene: relevance to coronary artery disease. Journal of Molecular Medicine, 2009, 87, 723-733.	1.7	16
119	Preservation Analysis of Macrophage Gene Coexpression Between Human and Mouse Identifies PARK2 as a Genetically Controlled Master Regulator of Oxidative Phosphorylation in Humans. G3: Genes, Genomes, Genetics, 2016, 6, 3361-3371.	0.8	15
120	Identification of two polymorphisms in the early growth response protein-1 gene: possible association with lipid variables. Journal of Molecular Medicine, 2000, 78, 81-86.	1.7	14
121	Cytokine phenotype, genotype, and renal outcomes at cardiac surgery. Cytokine, 2013, 61, 275-284.	1.4	14
122	Automated detection of informative combined effects in genetic association studies of complex traits. Genome Research, 2003, 13, 1952-60.	2.4	14
123	Atherosclerosis: From Genetic Polymorphisms to System Genetics. Cardiovascular Toxicology, 2005, 5, 143-152.	1.1	12
124	Haplotypic analysis of tag SNPs of the interleukin-18 gene in relation to cardiovascular disease events: the MORGAM Project. European Journal of Human Genetics, 2008, 16, 1512-1520.	1.4	12
125	Adrenomedullin and Arterial Stiffness. Circulation: Cardiovascular Genetics, 2014, 7, 634-641.	5.1	12
126	SAH gene variants are associated with obesity-related hypertension in Caucasians: the PEGASE Study. Journal of Hypertension, 2007, 25, 557-564.	0.3	11

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127	Polymorphisms in the genes encoding platelet-derived growth factor A and Î± receptor. Journal of Molecular Medicine, 2000, 78, 287-292.	1.7	10
128	Influence of cholesteryl ester transfer protein, peroxisome proliferatorâ€“activated receptor Î±, 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. Metabolism: Clinical and Experimental, 2009, 58, 283-289.	1.5	9
129	Osteopontin gene variation and cardio/cerebrovascular disease phenotypes. Atherosclerosis, 2009, 206, 209-215.	0.4	9
130	Single Nucleotide Polymorphisms with Cis-Regulatory Effects on Long Non-Coding Transcripts in Human Primary Monocytes. PLoS ONE, 2014, 9, e102612.	1.1	9
131	Renin-angiotensin system genes as candidate genes in cardiovascular diseases. Trends in Cardiovascular Medicine, 1993, 3, 250-258.	2.3	8
132	P-selectin gene polymorphisms and risk of coronary heart disease among Tunisians. Journal of Thrombosis and Thrombolysis, 2009, 28, 314-319.	1.0	8
133	Comprehensive Exploration of the Effects of miRNA SNPs on Monocyte Gene Expression. PLoS ONE, 2012, 7, e45863.	1.1	8
134	Identification and Functional Analyses of Molecular Haplotypes of the Human Osteoprotegerin Gene Promoter. Arteriosclerosis, Thrombosis, and Vascular Biology, 2009, 29, 1638-1643.	1.1	7
135	Patterns of Alcohol Consumption and Cardiovascular Risk in Northern Ireland and France. Annals of Epidemiology, 2007, 17, S75-S80.	0.9	6
136	Polymorphisms in 33 inflammatory genes and risk of myocardial infarctionâ€”a system genetics approach. Journal of Molecular Medicine, 2007, 85, 1271-1280.	1.7	6
137	Lack of association of genetic variants in the LRP8 gene with familial and sporadic myocardial infarction. Journal of Molecular Medicine, 2008, 86, 1163-1170.	1.7	6
138	Association of three polymorphisms selected from a genome-wide association study with coronary heart disease in the Tunisian population. Journal of Thrombosis and Thrombolysis, 2010, 29, 114-118.	1.0	6
139	Heritability, Weak Effects, and Rare Variants in Genomewide Association Studies. Clinical Chemistry, 2011, 57, 1263-1266.	1.5	5
140	Molecular investigation of the functional relevance of missense variants of ICAM-1. Pharmacogenetics and Genomics, 2008, 18, 1017-1019.	0.7	3
141	Neutrophil elastase gene variation and coronary heart disease. Pharmacogenetics and Genomics, 2007, 17, 629-637.	0.7	2
142	Coronary heart disease aetiology: associations and causality. Comptes Rendus - Biologies, 2007, 330, 299-305.	0.1	1
143	Effects of Alcohol on Lipids and Lipoprotein Metabolism. Medical Science Symposia Series, 2000, , 11-22.	0.0	0
144	The genetic contribution to the onset of acute coronary heart disease. Developments in Cardiovascular Medicine, 1996, , 111-133.	0.1	0

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145	First case of a humanRASGRP2mutation affecting Rap1 activation in platelets and causing severe bleeding.. Journal of Cell Biology, 2014, 206, 2061OIA111.	2.3	0