

Christian Hengstenberg

List of Publications by Year in descending order

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Version: 2024-02-01

341
papers

45,380
citations

5574

82
h-index

2243

201
g-index

361
all docs

361
docs citations

361
times ranked

47038
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	27.8	3,823
2	Biological, clinical and population relevance of 95 loci for blood lipids. <i>Nature</i> , 2010, 466, 707-713.	27.8	3,249
3	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010, 42, 937-948.	21.4	2,634
4	A comprehensive 1000 Genomesâ€‘based genome-wide association meta-analysis of coronary artery disease. <i>Nature Genetics</i> , 2015, 47, 1121-1130.	21.4	2,054
5	Plasma HDL cholesterol and risk of myocardial infarction: a mendelian randomisation study. <i>Lancet, The</i> , 2012, 380, 572-580.	13.7	1,937
6	Genomewide Association Analysis of Coronary Artery Disease. <i>New England Journal of Medicine</i> , 2007, 357, 443-453.	27.0	1,865
7	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	21.4	1,818
8	Hundreds of variants clustered in genomic loci and biological pathways affect human height. <i>Nature</i> , 2010, 467, 832-838.	27.8	1,789
9	Large-scale association analysis identifies 13 new susceptibility loci for coronary artery disease. <i>Nature Genetics</i> , 2011, 43, 333-338.	21.4	1,685
10	Large-scale association analysis identifies new risk loci for coronary artery disease. <i>Nature Genetics</i> , 2013, 45, 25-33.	21.4	1,439
11	Genome-wide association of early-onset myocardial infarction with single nucleotide polymorphisms and copy number variants. <i>Nature Genetics</i> , 2009, 41, 334-341.	21.4	990
12	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. <i>Nature Genetics</i> , 2010, 42, 949-960.	21.4	836
13	Transcatheter Aortic Valve Implantation in Failed Bioprosthetic Surgical Valves. <i>JAMA - Journal of the American Medical Association</i> , 2014, 312, 162.	7.4	762
14	DNA methylation and body-mass index: a genome-wide analysis. <i>Lancet, The</i> , 2014, 383, 1990-1998.	13.7	686
15	Genome-wide association analyses identify 18 new loci associated with serum urate concentrations. <i>Nature Genetics</i> , 2013, 45, 145-154.	21.4	675
16	Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. <i>Nature Genetics</i> , 2013, 45, 501-512.	21.4	578
17	Transcatheter Aortic Valve Replacement for Degenerative Bioprosthetic Surgical Valves. <i>Circulation</i> , 2012, 126, 2335-2344.	1.6	528
18	A genome-wide meta-analysis identifies 22 loci associated with eight hematological parameters in the HaemGen consortium. <i>Nature Genetics</i> , 2009, 41, 1182-1190.	21.4	481

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19	Paclitaxel-Coated Balloon Catheter Versus Paclitaxel-Coated Stent for the Treatment of Coronary In-Stent Restenosis. <i>Circulation</i> , 2009, 119, 2986-2994.	1.6	451
20	New susceptibility locus for coronary artery disease on chromosome 3q22.3. <i>Nature Genetics</i> , 2009, 41, 280-282.	21.4	440
21	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.	21.4	427
22	Coding Variation in <i>ANGPTL4</i> , <i>LPL</i> and <i>SVEP1</i> and the Risk of Coronary Disease. <i>New England Journal of Medicine</i> , 2016, 374, 1134-1144.	27.0	427
23	New gene functions in megakaryopoiesis and platelet formation. <i>Nature</i> , 2011, 480, 201-208.	27.8	401
24	FTO genotype is associated with phenotypic variability of body mass index. <i>Nature</i> , 2012, 490, 267-272.	27.8	383
25	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.	3.5	371
26	ISAR-SAFE: a randomized, double-blind, placebo-controlled trial of 6 vs. 12 months of clopidogrel therapy after drug-eluting stenting. <i>European Heart Journal</i> , 2015, 36, 1252-1263.	2.2	366
27	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
28	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.	3.5	331
29	A comprehensive linkage analysis for myocardial infarction and its related risk factors. <i>Nature Genetics</i> , 2002, 30, 210-214.	21.4	313
30	The impact of low-frequency and rare variants on lipid levels. <i>Nature Genetics</i> , 2015, 47, 589-597.	21.4	310
31	Shared Genetic Susceptibility to Ischemic Stroke and Coronary Artery Disease. <i>Stroke</i> , 2014, 45, 24-36.	2.0	302
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.	2.2	292
33	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.	2.7	285
34	A trans-acting locus regulates an anti-viral expression network and type 1 diabetes risk. <i>Nature</i> , 2010, 467, 460-464.	27.8	271
35	Association of Early Repolarization Pattern on ECG with Risk of Cardiac and All-Cause Mortality: A Population-Based Prospective Cohort Study (MONICA/KORA). <i>PLoS Medicine</i> , 2010, 7, e1000314.	8.4	246
36	Dysfunctional nitric oxide signalling increases risk of myocardial infarction. <i>Nature</i> , 2013, 504, 432-436.	27.8	230

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37	Large-Scale Gene-Centric Meta-analysis across 32 Studies Identifies Multiple Lipid Loci. <i>American Journal of Human Genetics</i> , 2012, 91, 823-838.	6.2	227
38	Genetically Determined Height and Coronary Artery Disease. <i>New England Journal of Medicine</i> , 2015, 372, 1608-1618.	27.0	220
39	Systematic Evaluation of Pleiotropy Identifies 6 Further Loci Associated With Coronary Artery Disease. <i>Journal of the American College of Cardiology</i> , 2017, 69, 823-836.	2.8	214
40	Transcatheter Aortic Valve Replacement in Pure Native Aortic Valve Regurgitation. <i>Journal of the American College of Cardiology</i> , 2017, 70, 2752-2763.	2.8	207
41	Genetic Determinants of Circulating Sphingolipid Concentrations in European Populations. <i>PLoS Genetics</i> , 2009, 5, e1000672.	3.5	184
42	Transcatheter Mitral Valve Replacement for Degenerated Bioprosthetic Valves and Failed Annuloplasty Rings. <i>Journal of the American College of Cardiology</i> , 2017, 70, 1121-1131.	2.8	183
43	Inheritance of coronary artery disease in men: an analysis of the role of the Y chromosome. <i>Lancet</i> , 2012, 379, 915-922.	13.7	179
44	Epigenetics meets metabolomics: an epigenome-wide association study with blood serum metabolic traits. <i>Human Molecular Genetics</i> , 2014, 23, 534-545.	2.9	169
45	Effect of Compensated Renal Dysfunction on Approved Heart Failure Markers. <i>Hypertension</i> , 2005, 46, 118-123.	2.7	162
46	Comparison of Vascular Closure Devices vs Manual Compression After Femoral Artery Puncture. <i>JAMA - Journal of the American Medical Association</i> , 2014, 312, 1981.	7.4	162
47	Design of the Coronary Artery Disease Genome-Wide Replication And Meta-Analysis (CARDIoGRAM) Study. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 475-483.	5.1	159
48	Distinct Heritable Patterns of Angiographic Coronary Artery Disease in Families With Myocardial Infarction. <i>Circulation</i> , 2005, 111, 855-862.	1.6	153
49	Peroxisome Proliferator-Activated Receptor β Gene Regulates Left Ventricular Growth in Response to Exercise and Hypertension. <i>Circulation</i> , 2002, 105, 950-955.	1.6	149
50	Predictors of Permanent Pacemaker Implantations and New-Onset Conduction Abnormalities With the SAPIEN 3 Balloon-Expandable Transcatheter Heart Valve. <i>JACC: Cardiovascular Interventions</i> , 2016, 9, 244-254.	2.9	149
51	A Genome-Wide Association Study Reveals Variants in ARL15 that Influence Adiponectin Levels. <i>PLoS Genetics</i> , 2009, 5, e1000768.	3.5	148
52	Association of the PHACTR1/EDN1 Genetic Locus With Spontaneous Coronary Artery Dissection. <i>Journal of the American College of Cardiology</i> , 2019, 73, 58-66.	2.8	147
53	Edoxaban versus Vitamin K Antagonist for Atrial Fibrillation after TAVR. <i>New England Journal of Medicine</i> , 2021, 385, 2150-2160.	27.0	144
54	Genetic Regulation of Serum Phytosterol Levels and Risk of Coronary Artery Disease. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 331-339.	5.1	141

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55	Genetic variation at chromosome 1p13.3 affects sortilin mRNA expression, cellular LDL-uptake and serum LDL levels which translates to the risk of coronary artery disease. <i>Atherosclerosis</i> , 2010, 208, 183-189.	0.8	141
56	Causal Assessment of Serum Urate Levels in Cardiometabolic Diseases Through a Mendelian Randomization Study. <i>Journal of the American College of Cardiology</i> , 2016, 67, 407-416.	2.8	138
57	Lifelong Reduction of LDL-Cholesterol Related to a Common Variant in the LDL-Receptor Gene Decreases the Risk of Coronary Artery Disease—A Mendelian Randomisation Study. <i>PLoS ONE</i> , 2008, 3, e2986.	2.5	137
58	Safety of the oral factor Xla inhibitor asundexian compared with apixaban in patients with atrial fibrillation (PACIFIC-AF): a multicentre, randomised, double-blind, double-dummy, dose-finding phase 2 study. <i>Lancet</i> , The, 2022, 399, 1383-1390.	13.7	131
59	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
60	N-Terminal Pro-Brain Natriuretic Peptide After Myocardial Infarction. <i>Hypertension</i> , 2002, 39, 99-104.	2.7	128
61	Integrating Genome-Wide Genetic Variations and Monocyte Expression Data Reveals Trans-Regulated Gene Modules in Humans. <i>PLoS Genetics</i> , 2011, 7, e1002367.	3.5	126
62	A Genome-Wide Association Study for Coronary Artery Disease Identifies a Novel Susceptibility Locus in the Major Histocompatibility Complex. <i>Circulation: Cardiovascular Genetics</i> , 2012, 5, 217-225.	5.1	125
63	Genome-wide association study identifies a new locus for coronary artery disease on chromosome 10p11.23. <i>European Heart Journal</i> , 2011, 32, 158-168.	2.2	124
64	Bivalirudin Versus Heparin Anticoagulation in Transcatheter Aortic Valve Replacement. <i>Journal of the American College of Cardiology</i> , 2015, 66, 2860-2868.	2.8	116
65	Lack of Association Between a Polymorphism of the Aldosterone Synthase Gene and Left Ventricular Structure. <i>Circulation</i> , 1999, 99, 2255-2260.	1.6	110
66	Genetic Association Study Identifies HSPB7 as a Risk Gene for Idiopathic Dilated Cardiomyopathy. <i>PLoS Genetics</i> , 2010, 6, e1001167.	3.5	110
67	Circulating Brain-Derived Neurotrophic Factor Concentrations and the Risk of Cardiovascular Disease in the Community. <i>Journal of the American Heart Association</i> , 2015, 4, e001544.	3.7	107
68	Evidence for FHL1 as a novel disease gene for isolated hypertrophic cardiomyopathy. <i>Human Molecular Genetics</i> , 2012, 21, 3237-3254.	2.9	106
69	Conscious Sedation Versus General Anesthesia in Transcatheter Aortic Valve Replacement. <i>JACC: Cardiovascular Interventions</i> , 2018, 11, 567-578.	2.9	102
70	Natural History of Functional Tricuspid Regurgitation. <i>JACC: Cardiovascular Imaging</i> , 2019, 12, 389-397.	5.3	102
71	IL6 Gene Promoter Polymorphisms and Type 2 Diabetes: Joint Analysis of Individual Participants' Data From 21 Studies. <i>Diabetes</i> , 2006, 55, 2915-2921.	0.6	99
72	Neointimal Modification With Scoring Balloon and Efficacy of Drug-Coated Balloon Therapy in Patients With Restenosis in Drug-Eluting Coronary Stents. <i>JACC: Cardiovascular Interventions</i> , 2017, 10, 1332-1340.	2.9	98

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73	Comparison of Self-Expanding Bioprostheses for Transcatheter Aortic Valve Replacement in Patients With Symptomatic Severe Aortic Stenosis. <i>Circulation</i> , 2020, 142, 2431-2442.	1.6	96
74	Real-world experience using the ACURATE neo prosthesis: 30-day outcomes of 1,000 patients enrolled in the SAVI TF registry. <i>EuroIntervention</i> , 2018, 13, e1764-e1770.	3.2	96
75	Association of angiotensin-converting enzyme 2 (ACE2) gene polymorphisms with parameters of left ventricular hypertrophy in men. <i>Journal of Molecular Medicine</i> , 2006, 84, 88-96.	3.9	95
76	Cardiovascular Risk Factors and Estimated Risk for CAD in a Randomized Trial Comparing Calcineurin Inhibitors in Renal Transplantation. <i>American Journal of Transplantation</i> , 2003, 3, 982-987.	4.7	93
77	Exome-wide association study reveals novel susceptibility genes to sporadic dilated cardiomyopathy. <i>PLoS ONE</i> , 2017, 12, e0172995.	2.5	92
78	Incidence and outcomes of emergent cardiac surgery during transfemoral transcatheter aortic valve implantation (TAVI): insights from the European Registry on Emergent Cardiac Surgery during TAVI (EuRECS-TAVI). <i>European Heart Journal</i> , 2018, 39, 676-684.	2.2	91
79	Common Polymorphisms Influencing Serum Uric Acid Levels Contribute to Susceptibility to Gout, but Not to Coronary Artery Disease. <i>PLoS ONE</i> , 2009, 4, e7729.	2.5	90
80	Heritability of Early Repolarization. <i>Circulation: Cardiovascular Genetics</i> , 2011, 4, 134-138.	5.1	89
81	High-Sensitivity Troponin T and Mortality After Elective Percutaneous Coronary Intervention. <i>Journal of the American College of Cardiology</i> , 2016, 68, 2259-2268.	2.8	88
82	A Unifying Concept for the Quantitative Assessment of Secondary Mitral Regurgitation. <i>Journal of the American College of Cardiology</i> , 2019, 73, 2506-2517.	2.8	86
83	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.	2.8	84
84	Multicenter Comparison of Novel Self-Expanding Versus Balloon-Expandable Transcatheter Heart Valves. <i>JACC: Cardiovascular Interventions</i> , 2017, 10, 2078-2087.	2.9	84
85	Light chain and transthyretin cardiac amyloidosis in severe aortic stenosis: prevalence, screening possibilities, and outcome. <i>European Journal of Heart Failure</i> , 2020, 22, 1852-1862.	7.1	82
86	Association of the T8590C Polymorphism of CYP4A11 With Hypertension in the MONICA Augsburg Echocardiographic Substudy. <i>Hypertension</i> , 2005, 46, 766-771.	2.7	80
87	The novel genetic variant predisposing to coronary artery disease in the region of the PSRC1 and CELSR2 genes on chromosome 1 associates with serum cholesterol. <i>Journal of Molecular Medicine</i> , 2008, 86, 1233-1241.	3.9	80
88	Discovery and Fine-Mapping of Glycaemic and Obesity-Related Trait Loci Using High-Density Imputation. <i>PLoS Genetics</i> , 2015, 11, e1005230.	3.5	77
89	Association of Polymorphisms of the Apolipoprotein(a) Gene With Lipoprotein(a) Levels and Myocardial Infarction. <i>Circulation</i> , 2003, 107, 696-701.	1.6	75
90	Association of Common Polymorphisms in GLUT9 Gene with Gout but Not with Coronary Artery Disease in a Large Case-Control Study. <i>PLoS ONE</i> , 2008, 3, e1948.	2.5	75

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91	No association of interleukin-6 gene polymorphism (~ 174 G/C) with myocardial infarction or traditional cardiovascular risk factors. <i>International Journal of Cardiology</i> , 2004, 97, 205-212.	1.7	71
92	Aspirin for primary prevention of cardiovascular disease: a meta-analysis with a particular focus on subgroups. <i>BMC Medicine</i> , 2019, 17, 198.	5.5	71
93	Genetic Variants Within the <i>LPIN1</i> Gene, Encoding Lipin, Are Influencing Phenotypes of the Metabolic Syndrome in Humans. <i>Diabetes</i> , 2008, 57, 209-217.	0.6	70
94	Genetic Predisposition to Higher Blood Pressure Increases Coronary Artery Disease Risk. <i>Hypertension</i> , 2013, 61, 995-1001.	2.7	70
95	Systematic analysis of variants related to familial hypercholesterolemia in families with premature myocardial infarction. <i>European Journal of Human Genetics</i> , 2016, 24, 191-197.	2.8	70
96	The SAVI-TF Registry. <i>JACC: Cardiovascular Interventions</i> , 2018, 11, 1368-1374.	2.9	64
97	Modulation of Genetic Associations with Serum Urate Levels by Body-Mass-Index in Humans. <i>PLoS ONE</i> , 2015, 10, e0119752.	2.5	64
98	Signal transducer of inflammation gp130 modulates atherosclerosis in mice and man. <i>Journal of Experimental Medicine</i> , 2007, 204, 1935-1944.	8.5	63
99	Prosthesis sizing for transcatheter aortic valve implantation – Comparison of three dimensional transesophageal echocardiography with multislice computed tomography. <i>International Journal of Cardiology</i> , 2013, 168, 3431-3438.	1.7	62
100	Edoxaban Versus standard of care and their effects on clinical outcomes in patients having undergone Transcatheter Aortic Valve Implantation in Atrial Fibrillation – Rationale and design of the ENVISAGE-TAVI AF trial. <i>American Heart Journal</i> , 2018, 205, 63-69.	2.7	62
101	Subclinical Leaflet Thrombosis After Transcatheter Aortic Valve Replacement. <i>JACC: Cardiovascular Interventions</i> , 2021, 14, 2643-2656.	2.9	62
102	Meta-analysis of Gene-Level Associations for Rare Variants Based on Single-Variant Statistics. <i>American Journal of Human Genetics</i> , 2013, 93, 236-248.	6.2	60
103	Evolution of outcome and complications in TAVR: a meta-analysis of observational and randomized studies. <i>Scientific Reports</i> , 2020, 10, 15568.	3.3	60
104	Diagnostic and Prognostic Utility of Cardiac Magnetic Resonance Imaging in Aortic Regurgitation. <i>JACC: Cardiovascular Imaging</i> , 2019, 12, 1474-1483.	5.3	59
105	A meta-analysis of genome-wide association studies of the electrocardiographic early repolarization pattern. <i>Heart Rhythm</i> , 2012, 9, 1627-1634.	0.7	58
106	Neutrophil Extracellular Trap Degradation by Differently Polarized Macrophage Subsets. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 2265-2278.	2.4	54
107	Genome-Wide Haplotype Analysis of Cis Expression Quantitative Trait Loci in Monocytes. <i>PLoS Genetics</i> , 2013, 9, e1003240.	3.5	53
108	Point Mutations in Mitochondrial DNA of Patients with Dilated Cardiomyopathy. <i>Journal of Molecular and Cellular Cardiology</i> , 1997, 29, 2699-2709.	1.9	51

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109	Joint analysis of individual participantsâ€™ data from 17 studies on the association of the <i>IL6</i> variant -174G>C with circulating glucose levels, interleukin-6 levels, and body mass index. <i>Annals of Medicine</i> , 2009, 41, 128-138.	3.8	51
110	Early experiences with miniaturized extracorporeal life-support in the catheterization laboratory. <i>European Journal of Cardio-thoracic Surgery</i> , 2012, 42, 858-863.	1.4	51
111	6-Month Outcomes of the TricValveÂ System in Patients With Tricuspid Regurgitation. <i>JACC: Cardiovascular Interventions</i> , 2022, 15, 1366-1377.	2.9	51
112	Outcomes After Transcatheter AorticÂ Valve Replacement Using aÂ NovelÂ Balloon-Expandable TranscatheterÂ HeartÂ Valve. <i>JACC: Cardiovascular Interventions</i> , 2015, 8, 1809-1816.	2.9	50
113	Evaluation of the Aldosterone Synthase (CYP11B2) Gene Polymorphism in Patients With Myocardial Infarction. <i>Hypertension</i> , 2000, 35, 704-709.	2.7	49
114	Familial aggregation of left main coronary artery disease and future risk of coronary events in asymptomatic siblings of affected patients. <i>European Heart Journal</i> , 2007, 28, 2432-2437.	2.2	49
115	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.	2.2	49
116	Functional improvement in heart failure patients treated with beta-blockers is associated with a decline of cytokine levels. <i>International Journal of Cardiology</i> , 2005, 103, 182-186.	1.7	48
117	Familial predisposition of left ventricular hypertrophy. <i>Journal of the American College of Cardiology</i> , 1999, 33, 1685-1691.	2.8	46
118	Generation of Highly Purified Human Cardiomyocytes from Peripheral Blood Mononuclear Cell-Derived Induced Pluripotent Stem Cells. <i>PLoS ONE</i> , 2015, 10, e0126596.	2.5	46
119	Emergency and prophylactic use of miniaturized venoâ€ arterial extracorporeal membrane oxygenation in transcatheter aortic valve implantation. <i>Catheterization and Cardiovascular Interventions</i> , 2013, 82, E542-51.	1.7	45
120	Cerebral Embolism During TranscatheterÂ Aortic Valve Replacement. <i>Journal of the American College of Cardiology</i> , 2016, 68, 589-599.	2.8	45
121	Lack of Association Between the <i>MEF2A</i> Gene and Myocardial Infarction. <i>Circulation</i> , 2008, 117, 185-191.	1.6	44
122	Prognostic impact of sleep duration and sleep efficiency on mortality in patients with chronic heart failure. <i>Sleep Medicine</i> , 2013, 14, 502-509.	1.6	42
123	The lipoprotein subfraction profile: heritability and identification of quantitative trait loci. <i>Journal of Lipid Research</i> , 2008, 49, 715-723.	4.2	41
124	De-Ritis Ratio Improves Long-Term Risk Prediction after Acute Myocardial Infarction. <i>Journal of Clinical Medicine</i> , 2018, 7, 474.	2.4	41
125	Prognostic impact of anemia and iron-deficiency anemia in a contemporary cohort of patients undergoing transcatheter aortic valve implantation. <i>International Journal of Cardiology</i> , 2017, 244, 93-99.	1.7	40
126	Transcatheter Valve SELECTION in Patients With Right Bundle Branch Block and Impact on Pacemaker Implantations. <i>JACC: Cardiovascular Interventions</i> , 2019, 12, 1781-1793.	2.9	38

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127	Tafamidis treatment delays structural and functional changes of the left ventricle in patients with transthyretin amyloid cardiomyopathy. <i>European Heart Journal Cardiovascular Imaging</i> , 2022, 23, 767-780.	1.2	38
128	Impact of three-dimensional transesophageal echocardiography on prosthesis sizing for transcatheter aortic valve implantation. <i>Catheterization and Cardiovascular Interventions</i> , 2012, 80, 956-963.	1.7	37
129	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.	6.2	37
130	A machine learning algorithm supports ultrasound-naïve novices in the acquisition of diagnostic echocardiography loops and provides accurate estimation of LVEF. <i>International Journal of Cardiovascular Imaging</i> , 2021, 37, 577-586.	1.5	37
131	Powerful Identification of Cis-regulatory SNPs in Human Primary Monocytes Using Allele-Specific Gene Expression. <i>PLoS ONE</i> , 2012, 7, e52260.	2.5	36
132	Epistatic interaction between haplotypes of the ghrelin ligand and receptor genes influence susceptibility to myocardial infarction and coronary artery disease. <i>Human Molecular Genetics</i> , 2007, 16, 887-899.	2.9	35
133	Impact of percutaneous closure device type on vascular and bleeding complications after TAVR: A post hoc analysis from the BRAVO randomized trial. <i>Catheterization and Cardiovascular Interventions</i> , 2019, 93, 1374-1381.	1.7	35
134	ST-Segment Elevation Myocardial Infarction Following Transcatheter Aortic Valve Replacement. <i>Journal of the American College of Cardiology</i> , 2021, 77, 2187-2199.	2.8	35
135	Tumor Marker Carbohydrate Antigen 125 Predicts Adverse Outcome After Transcatheter Aortic Valve Implantation. <i>JACC: Cardiovascular Interventions</i> , 2013, 6, 487-496.	2.9	34
136	Cardiac Magnetic Resonance T1 Mapping in Cardiac Amyloidosis. <i>JACC: Cardiovascular Imaging</i> , 2018, 11, 1924-1926.	5.3	34
137	Feature Tracking of Global Longitudinal Strain by Using Cardiovascular MRI Improves Risk Stratification in Heart Failure with Preserved Ejection Fraction. <i>Radiology</i> , 2020, 296, 290-298.	7.3	34
138	Genome-wide meta-analysis of common variant differences between men and women. <i>Human Molecular Genetics</i> , 2012, 21, 4805-4815.	2.9	33
139	Large-Scale Analysis of Determinants, Stability, and Heritability of High-Density Lipoprotein Cholesterol Efflux Capacity. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2017, 37, 1956-1962.	2.4	33
140	Large-Scale Candidate Gene Analysis of HDL Particle Features. <i>PLoS ONE</i> , 2011, 6, e14529.	2.5	32
141	Disproportionate Functional Mitral Regurgitation. <i>JACC: Cardiovascular Imaging</i> , 2019, 12, 2088-2090.	5.3	32
142	Burden, treatment use, and outcome of secondary mitral regurgitation across the spectrum of heart failure: observational cohort study. <i>BMJ</i> , The, 2021, 373, n1421.	6.0	32
143	Common Genetic Variants in <i>ANK2</i> Modulate QT Interval. <i>Circulation: Cardiovascular Genetics</i> , 2008, 1, 93-99.	5.1	29
144	Gender-specific differences in valvular heart disease. <i>Wiener Klinische Wochenschrift</i> , 2020, 132, 61-68.	1.9	29

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145	Four Genetic Loci Influencing Electrocardiographic Indices of Left Ventricular Hypertrophy. Circulation: Cardiovascular Genetics, 2011, 4, 626-635.	5.1	28
146	Genetic associations with lipoprotein subfractions provide information on their biological nature. Human Molecular Genetics, 2012, 21, 1433-1443.	2.9	28
147	Blood urea nitrogen has additive value beyond estimated glomerular filtration rate for prediction of long-term mortality in patients with acute myocardial infarction. European Journal of Internal Medicine, 2019, 59, 84-90.	2.2	28
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