

Folkert Asselbergs

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

499
papers

22,071
citations

65
h-index

138
g-index

601
ext. papers

28,902
ext. citations

7.7
avg, IF

6.44
L-index

#	Paper	IF	Citations
499	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015 , 518, 197-206	50.4	2687
498	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014 , 46, 1173-86	36.3	1339
497	The interleukin-6 receptor as a target for prevention of coronary heart disease: a mendelian randomisation analysis. <i>Lancet, The</i> , 2012 , 379, 1214-24	40	658
496	Interleukin-6 receptor pathways in coronary heart disease: a collaborative meta-analysis of 82 studies. <i>Lancet, The</i> , 2012 , 379, 1205-13	40	522
495	Worsening renal function and prognosis in heart failure: systematic review and meta-analysis. <i>Journal of Cardiac Failure</i> , 2007 , 13, 599-608	3.3	444
494	Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. <i>Nature Genetics</i> , 2013 , 45, 501-12	36.3	437
493	Mendelian randomization of blood lipids for coronary heart disease. <i>European Heart Journal</i> , 2015 , 36, 539-50	9.5	417
492	Rare and low-frequency coding variants alter human adult height. <i>Nature</i> , 2017 , 542, 186-190	50.4	412
491	HMG-coenzyme A reductase inhibition, type 2 diabetes, and bodyweight: evidence from genetic analysis and randomised trials. <i>Lancet, The</i> , 2015 , 385, 351-61	40	409
490	Association between alcohol and cardiovascular disease: Mendelian randomisation analysis based on individual participant data. <i>BMJ, The</i> , 2014 , 349, g4164	5.9	406
489	Bioinformatics challenges for genome-wide association studies. <i>Bioinformatics</i> , 2010 , 26, 445-55	7.2	401
488	Effects of fosinopril and pravastatin on cardiovascular events in subjects with microalbuminuria. <i>Circulation</i> , 2004 , 110, 2809-16	16.7	401
487	Blood pressure-lowering treatment based on cardiovascular risk: a meta-analysis of individual patient data. <i>Lancet, The</i> , 2014 , 384, 591-598	40	376
486	Coding Variation in ANGPTL4, LPL, and SVEP1 and the Risk of Coronary Disease. <i>New England Journal of Medicine</i> , 2016 , 374, 1134-44	59.2	325
485	FTO genotype is associated with phenotypic variability of body mass index. <i>Nature</i> , 2012 , 490, 267-72	50.4	304
484	Seventy-five genetic loci influencing the human red blood cell. <i>Nature</i> , 2012 , 492, 369-75	50.4	257
483	Common variants in 22 loci are associated with QRS duration and cardiac ventricular conduction. <i>Nature Genetics</i> , 2010 , 42, 1068-76	36.3	249

482	PCSK9 genetic variants and risk of type 2 diabetes: a mendelian randomisation study. <i>Lancet Diabetes and Endocrinology</i> , 2017 , 5, 97-105	18.1	225
481	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. <i>Nature Genetics</i> , 2015 , 47, 1282-1293	36.3	223
480	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	6	220
479	A Genotype-Guided Strategy for Oral P2Y Inhibitors in Primary PCI. <i>New England Journal of Medicine</i> , 2019 , 381, 1621-1631	59.2	219
478	Large-scale gene-centric meta-analysis across 39 studies identifies type 2 diabetes loci. <i>American Journal of Human Genetics</i> , 2012 , 90, 410-25	11	214
477	The Translational Landscape of the Human Heart. <i>Cell</i> , 2019 , 178, 242-260.e29	56.2	210
476	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. <i>Nature Genetics</i> , 2014 , 46, 826-36	36.3	199
475	Large-scale gene-centric meta-analysis across 32 studies identifies multiple lipid loci. <i>American Journal of Human Genetics</i> , 2012 , 91, 823-38	11	189
474	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. <i>Nature Genetics</i> , 2018 , 50, 26-41	36.3	186
473	Trans-ancestry meta-analyses identify rare and common variants associated with blood pressure and hypertension. <i>Nature Genetics</i> , 2016 , 48, 1151-1161	36.3	181
472	Do men and women respond differently to blood pressure-lowering treatment? Results of prospectively designed overviews of randomized trials. <i>European Heart Journal</i> , 2008 , 29, 2669-80	9.5	172
471	Association of Lipid Fractions With Risks for Coronary Artery Disease and Diabetes. <i>JAMA Cardiology</i> , 2016 , 1, 692-9	16.2	168
470	Sodium intake affects urinary albumin excretion especially in overweight subjects. <i>Journal of Internal Medicine</i> , 2004 , 256, 324-30	10.8	159
469	Genetics, Clinical Features, and Long-Term Outcome of Noncompaction Cardiomyopathy. <i>Journal of the American College of Cardiology</i> , 2018 , 71, 711-722	15.1	158
468	Causal effects of body mass index on cardiometabolic traits and events: a Mendelian randomization analysis. <i>American Journal of Human Genetics</i> , 2014 , 94, 198-208	11	156
467	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	15.1	146
466	Genome-wide association and Mendelian randomisation analysis provide insights into the pathogenesis of heart failure. <i>Nature Communications</i> , 2020 , 11, 163	17.4	140
465	Dominant missense mutations in ABCC9 cause Cantú syndrome. <i>Nature Genetics</i> , 2012 , 44, 793-6	36.3	139

464	Gene-centric meta-analysis in 87,736 individuals of European ancestry identifies multiple blood-pressure-related loci. <i>American Journal of Human Genetics</i> , 2014 , 94, 349-60	11	131
463	Directional dominance on stature and cognition in diverse human populations. <i>Nature</i> , 2015 , 523, 459-463	50.4	119
462	Loci influencing blood pressure identified using a cardiovascular gene-centric array. <i>Human Molecular Genetics</i> , 2013 , 22, 1663-78	5.6	119
461	Correction of human phospholamban R14del mutation associated with cardiomyopathy using targeted nucleases and combination therapy. <i>Nature Communications</i> , 2015 , 6, 6955	17.4	119
460	Genetic analysis in 418 index patients with idiopathic dilated cardiomyopathy: overview of 10 years' experience. <i>European Journal of Heart Failure</i> , 2013 , 15, 628-36	12.3	117
459	Shadows of complexity: what biological networks reveal about epistasis and pleiotropy. <i>BioEssays</i> , 2009 , 31, 220-7	4.1	117
458	Interleukin-6 receptor pathways in abdominal aortic aneurysm. <i>European Heart Journal</i> , 2013 , 34, 3707-16	16.5	111
457	Big data from electronic health records for early and late translational cardiovascular research: challenges and potential. <i>European Heart Journal</i> , 2018 , 39, 1481-1495	9.5	106
456	A new prediction model for ventricular arrhythmias in arrhythmogenic right ventricular cardiomyopathy. <i>European Heart Journal</i> , 2019 , 40, 1850-1858	9.5	104
455	Meta-analysis of Dense Genecentric Association Studies Reveals Common and Uncommon Variants Associated with Height. <i>American Journal of Human Genetics</i> , 2011 , 88, 6-18	11	103
454	Pharmacological blood pressure lowering for primary and secondary prevention of cardiovascular disease across different levels of blood pressure: an individual participant-level data meta-analysis. <i>Lancet, The</i> , 2021 , 397, 1625-1636	40	101
453	Microanatomy of the Human Atherosclerotic Plaque by Single-Cell Transcriptomics. <i>Circulation Research</i> , 2020 , 127, 1437-1455	15.7	96
452	Effect of statins on atrial fibrillation: collaborative meta-analysis of published and unpublished evidence from randomised controlled trials. <i>BMJ, The</i> , 2011 , 342, d1250	5.9	95
451	Acceleration of cardiovascular disease by a dysfunctional prostacyclin receptor mutation: potential implications for cyclooxygenase-2 inhibition. <i>Circulation Research</i> , 2008 , 102, 986-93	15.7	95
450	Secretory phospholipase A(2)-IIA and cardiovascular disease: a mendelian randomization study. <i>Journal of the American College of Cardiology</i> , 2013 , 62, 1966-1976	15.1	91
449	Carotid Intima-Media Thickness Progression as Surrogate Marker for Cardiovascular Risk: Meta-Analysis of 119 Clinical Trials Involving 100 667 Patients. <i>Circulation</i> , 2020 , 142, 621-642	16.7	88
448	Association between chromosome 9p21 variants and the ankle-brachial index identified by a meta-analysis of 21 genome-wide association studies. <i>Circulation: Cardiovascular Genetics</i> , 2012 , 5, 100-12		84
447	Race/Ethnic Differences in the Associations of the Framingham Risk Factors with Carotid IMT and Cardiovascular Events. <i>PLoS ONE</i> , 2015 , 10, e0132321	3.7	83

446	A Mendelian Randomization Study of Circulating Uric Acid and Type 2 Diabetes. <i>Diabetes</i> , 2015 , 64, 3028-36	3.6	79
445	Cost-effectiveness of screening for albuminuria with subsequent fosinopril treatment to prevent cardiovascular events: A pharmaco-economic analysis linked to the prevention of renal and vascular endstage disease (PREVEND) study and the prevention of renal and vascular endstage disease intervention trial (PREVEND-IT). <i>Clinical Therapeutics</i> , 2006 , 28, 432-44	3.5	79
444	Causal Effects of Body Mass Index on Cardiometabolic Traits and Events: A Mendelian Randomization Analysis. <i>American Journal of Human Genetics</i> , 2014 , 94, 312	11	78
443	Meta-analysis of Dense Genecentric Association Studies Reveals Common and Uncommon Variants Associated with Height. <i>American Journal of Human Genetics</i> , 2012 , 90, 1116-1117	11	78
442	The effect of statins on urinary albumin excretion and glomerular filtration rate: results from both a randomized clinical trial and an observational cohort study. <i>Nephrology Dialysis Transplantation</i> , 2006 , 21, 3106-14	4.3	77
441	52 Genetic Loci Influencing Myocardial Mass. <i>Journal of the American College of Cardiology</i> , 2016 , 68, 1435-1448	15.1	76
440	Separating the mechanism-based and off-target actions of cholesteryl ester transfer protein inhibitors with CETP gene polymorphisms. <i>Circulation</i> , 2010 , 121, 52-62	16.7	76
439	Gene-centric meta-analyses of 108 912 individuals confirm known body mass index loci and reveal three novel signals. <i>Human Molecular Genetics</i> , 2013 , 22, 184-201	5.6	73
438	Relations between lipoprotein(a) concentrations, LPA genetic variants, and the risk of mortality in patients with established coronary heart disease: a molecular and genetic association study. <i>Lancet Diabetes and Endocrinology</i> , 2017 , 5, 534-543	18.1	69
437	Myeloperoxidase polymorphism related to cardiovascular events in coronary artery disease. <i>American Journal of Medicine</i> , 2004 , 116, 429-30	2.4	69
436	C-reactive protein and microalbuminuria are associated with atrial fibrillation. <i>International Journal of Cardiology</i> , 2005 , 98, 73-7	3.2	68
435	Prognostic burden of heart failure recorded in primary care, acute hospital admissions, or both: a population-based linked electronic health record cohort study in 2.1 million people. <i>European Journal of Heart Failure</i> , 2017 , 19, 1119-1127	12.3	66
434	Causal Effect of Plasminogen Activator Inhibitor Type 1 on Coronary Heart Disease. <i>Journal of the American Heart Association</i> , 2017 , 6,	6	65
433	Adult height, coronary heart disease and stroke: a multi-locus Mendelian randomization meta-analysis. <i>International Journal of Epidemiology</i> , 2016 , 45, 1927-1937	7.8	65
432	Platelet-reactivity tests identify patients at risk of secondary cardiovascular events: a systematic review and meta-analysis. <i>Journal of Thrombosis and Haemostasis</i> , 2014 , 12, 736-47	15.4	65
431	Genome-wide association study for circulating levels of PAI-1 provides novel insights into its regulation. <i>Blood</i> , 2012 , 120, 4873-81	2.2	65
430	Cystatin C and Cardiovascular Disease: A Mendelian Randomization Study. <i>Journal of the American College of Cardiology</i> , 2016 , 68, 934-45	15.1	65
429	A Simple and Computationally Efficient Approach to Multifactor Dimensionality Reduction Analysis of Gene-Gene Interactions for Quantitative Traits. <i>PLoS ONE</i> , 2013 , 8, e66545	3.7	63

428	The Cardiomyopathy Registry of the EURObservational Research Programme of the European Society of Cardiology: baseline data and contemporary management of adult patients with cardiomyopathies. <i>European Heart Journal</i> , 2018 , 39, 1784-1793	9.5	60
427	CYP2C19 genotype-guided antiplatelet therapy in ST-segment elevation myocardial infarction patients-Rationale and design of the Patient Outcome after primary PCI (POPular) Genetics study. <i>American Heart Journal</i> , 2014 , 168, 16-22.e1	4.9	59
426	Dilated Cardiomyopathy Due to BLC2-Associated Athanogene 3 '(BAG3) Mutations. <i>Journal of the American College of Cardiology</i> , 2018 , 72, 2471-2481	15.1	53
425	Pulmonary artery pressure-guided therapy in ambulatory patients with symptomatic heart failure: the CardioMEMS European Monitoring Study for Heart Failure (MEMS-HF). <i>European Journal of Heart Failure</i> , 2020 , 22, 1891-1901	12.3	52
424	Metabolomics Profile in Depression: A Pooled Analysis of 230 Metabolic Markers in 5283 Cases With Depression and 10,145 Controls. <i>Biological Psychiatry</i> , 2020 , 87, 409-418	7.9	51
423	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. <i>Nature Communications</i> , 2017 , 8, 15805	17.4	50
422	Annotation of loci from genome-wide association studies using tissue-specific quantitative interaction proteomics. <i>Nature Methods</i> , 2014 , 11, 868-74	21.6	50
421	Common genetic variation near the phospholamban gene is associated with cardiac repolarisation: meta-analysis of three genome-wide association studies. <i>PLoS ONE</i> , 2009 , 4, e6138	3.7	50
420	Cardiac complications in patients hospitalised with COVID-19. <i>European Heart Journal: Acute Cardiovascular Care</i> , 2020 , 9, 817-823	4.3	50
419	Meta-analysis of gene-level associations for rare variants based on single-variant statistics. <i>American Journal of Human Genetics</i> , 2013 , 93, 236-48	11	49
418	Truncating Titin (TTN) Variants in Chemotherapy-Induced Cardiomyopathy. <i>Journal of Cardiac Failure</i> , 2017 , 23, 476-479	3.3	48
417	Genetics of coronary artery disease: genome-wide association studies and beyond. <i>Atherosclerosis</i> , 2012 , 225, 1-10	3.1	48
416	Association of renal function with cardiac calcifications in older adults: the cardiovascular health study. <i>Nephrology Dialysis Transplantation</i> , 2009 , 24, 834-40	4.3	46
415	Vascular endothelial growth factor: the link between cardiovascular risk factors and microalbuminuria?. <i>International Journal of Cardiology</i> , 2004 , 93, 211-5	3.2	45
414	Meta-analysis of up to 622,409 individuals identifies 40 novel smoking behaviour associated genetic loci. <i>Molecular Psychiatry</i> , 2020 , 25, 2392-2409	15.1	45
413	Chemotherapy-Related Cardiac Dysfunction: A Systematic Review of Genetic Variants Modulating Individual Risk. <i>Circulation Genomic and Precision Medicine</i> , 2018 , 11, e001753	5.2	44
412	Pharmacogenetics of ACE inhibitor-induced angioedema and cough: a systematic review and meta-analysis. <i>Pharmacogenomics</i> , 2013 , 14, 249-60	2.6	44
411	Design and Implementation of the International Genetics and Translational Research in Transplantation Network. <i>Transplantation</i> , 2015 , 99, 2401-12	1.8	44

410	High prevalence of microalbuminuria in chronic heart failure patients. <i>Journal of Cardiac Failure</i> , 2005 , 11, 602-6	3.3	44
409	C-reactive protein and microalbuminuria differ in their associations with various domains of vascular disease. <i>Atherosclerosis</i> , 2004 , 172, 107-14	3.1	44
408	Genetic Association of Lipids and Lipid Drug Targets With Abdominal Aortic Aneurysm: A Meta-analysis. <i>JAMA Cardiology</i> , 2018 , 3, 26-33	16.2	44
407	Gender gap in acute coronary heart disease: Myth or reality?. <i>World Journal of Cardiology</i> , 2012 , 4, 36-47	2.1	43
406	Predicting arrhythmic risk in arrhythmogenic right ventricular cardiomyopathy: A systematic review and meta-analysis. <i>Heart Rhythm</i> , 2018 , 15, 1097-1107	6.7	42
405	European Cardiomyopathy Pilot Registry: EURObservational Research Programme of the European Society of Cardiology. <i>European Heart Journal</i> , 2016 , 37, 164-73	9.5	42
404	Serially measured circulating microRNAs and adverse clinical outcomes in patients with acute heart failure. <i>European Journal of Heart Failure</i> , 2018 , 20, 89-96	12.3	41
403	Associations of autozygosity with a broad range of human phenotypes. <i>Nature Communications</i> , 2019 , 10, 4957	17.4	40
402	A systematic analysis of genetic dilated cardiomyopathy reveals numerous ubiquitously expressed and muscle-specific genes. <i>European Journal of Heart Failure</i> , 2015 , 17, 484-93	12.3	40
401	Distinct fibrosis pattern in desmosomal and phospholamban mutation carriers in hereditary cardiomyopathies. <i>Heart Rhythm</i> , 2017 , 14, 1024-1032	6.7	39
400	PR interval genome-wide association meta-analysis identifies 50 loci associated with atrial and atrioventricular electrical activity. <i>Nature Communications</i> , 2018 , 9, 2904	17.4	39
399	Genetic variants at chromosome 9p21 and risk of first versus subsequent coronary heart disease events: a systematic review and meta-analysis. <i>Journal of the American College of Cardiology</i> , 2014 , 63, 2234-45	15.1	39
398	Impact of statins in microalbuminuric subjects with the metabolic syndrome: a substudy of the PREVEND Intervention Trial. <i>European Heart Journal</i> , 2005 , 26, 1314-20	9.5	39
397	Characteristic adaptations of the extracellular matrix in dilated cardiomyopathy. <i>International Journal of Cardiology</i> , 2016 , 220, 634-46	3.2	38
396	Concept and design of a genome-wide association genotyping array tailored for transplantation-specific studies. <i>Genome Medicine</i> , 2015 , 7, 90	14.4	38
395	Gene-centric meta-analysis of lipid traits in African, East Asian and Hispanic populations. <i>PLoS ONE</i> , 2012 , 7, e50198	3.7	37
394	The Prognostic Value of Right Ventricular Deformation Imaging in Early Arrhythmogenic Right Ventricular Cardiomyopathy. <i>JACC: Cardiovascular Imaging</i> , 2019 , 12, 446-455	8.4	37
393	Long-term effects of fosinopril and pravastatin on cardiovascular events in subjects with microalbuminuria: Ten years of follow-up of Prevention of Renal and Vascular End-stage Disease Intervention Trial (PREVEND IT). <i>American Heart Journal</i> , 2011 , 161, 1171-8	4.9	36

392	Gender-specific correlations of plasminogen activator inhibitor-1 and tissue plasminogen activator levels with cardiovascular disease-related traits. <i>Journal of Thrombosis and Haemostasis</i> , 2007 , 5, 313-20	15.4	36
391	Fatty acid oxidation flux predicts the clinical severity of VLCAD deficiency. <i>Genetics in Medicine</i> , 2015 , 17, 989-94	8.1	35
390	Effects of blood pressure lowering on cardiovascular risk according to baseline body-mass index: a meta-analysis of randomised trials. <i>Lancet, The</i> , 2015 , 385, 867-74	4.0	35
389	Effects of fosinopril and pravastatin on carotid intima-media thickness in subjects with increased albuminuria. <i>Stroke</i> , 2005 , 36, 649-53	6.7	35
388	Exome Chip Meta-analysis Fine Maps Causal Variants and Elucidates the Genetic Architecture of Rare Coding Variants in Smoking and Alcohol Use. <i>Biological Psychiatry</i> , 2019 , 85, 946-955	7.9	35
387	Shared genetic pathways contribute to risk of hypertrophic and dilated cardiomyopathies with opposite directions of effect. <i>Nature Genetics</i> , 2021 , 53, 128-134	36.3	35
386	Genetic drug target validation using Mendelian randomisation. <i>Nature Communications</i> , 2020 , 11, 3255	17.4	34
385	Genome-wide association studies identify genetic loci for low von Willebrand factor levels. <i>European Journal of Human Genetics</i> , 2016 , 24, 1035-40	5.3	34
384	Cardiac amyloidosis: the need for early diagnosis. <i>Netherlands Heart Journal</i> , 2019 , 27, 525-536	2.2	34
383	Loss of Y Chromosome in Blood Is Associated With Major Cardiovascular Events During Follow-Up in Men After Carotid Endarterectomy. <i>Circulation: Cardiovascular Genetics</i> , 2017 , 10, e001544		34
382	Correlates of endothelial function and their relationship with inflammation in patients with familial hypercholesterolaemia. <i>Clinical Science</i> , 2003 , 104, 627-32	6.5	34
381	Meta-analysis of rare and common exome chip variants identifies S1PR4 and other loci influencing blood cell traits. <i>Nature Genetics</i> , 2016 , 48, 867-76	36.3	34
380	Human Validation of Genes Associated With a Murine Atherosclerotic Phenotype. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016 , 36, 1240-6	9.4	34
379	New Blood Pressure-Associated Loci Identified in Meta-Analyses of 475 000 Individuals. <i>Circulation: Cardiovascular Genetics</i> , 2017 , 10,		33
378	Genome-wide association study for circulating tissue plasminogen activator levels and functional follow-up implicates endothelial STXBP5 and STX2. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014 , 34, 1093-101	9.4	33
377	A role for CETP TaqIB polymorphism in determining susceptibility to atrial fibrillation: a nested case control study. <i>BMC Medical Genetics</i> , 2006 , 7, 39	2.1	33
376	Prognostic value of myeloperoxidase in patients with chest pain. <i>New England Journal of Medicine</i> , 2004 , 350, 516-8; author reply 516-8	59.2	33
375	CAPACITY-COVID: a European Registry to determine the role of cardiovascular disease in the COVID-19 pandemic. <i>European Heart Journal</i> , 2020 , 41, 1795-1796	9.5	32

374	Network analysis of coronary artery disease risk genes elucidates disease mechanisms and druggable targets. <i>Scientific Reports</i> , 2018 , 8, 3434	4.9	31
373	The ENCODE project and perspectives on pathways. <i>Genetic Epidemiology</i> , 2014 , 38, 275-80	2.6	31
372	A comprehensive evaluation of the genetic architecture of sudden cardiac arrest. <i>European Heart Journal</i> , 2018 , 39, 3961-3969	9.5	31
371	Variants associating with uterine leiomyoma highlight genetic background shared by various cancers and hormone-related traits. <i>Nature Communications</i> , 2018 , 9, 3636	17.4	31
370	A mutation update for the FLNC gene in myopathies and cardiomyopathies. <i>Human Mutation</i> , 2020 , 41, 1091-1111	4.7	30
369	Interaction between dietary fat intake and the cholesterol ester transfer protein TaqIB polymorphism in relation to HDL-cholesterol concentrations among US diabetic men. <i>American Journal of Clinical Nutrition</i> , 2007 , 86, 1524-9	7	30
368	Clinical impact of vasomotor function assessment and the role of ACE-inhibitors and statins. <i>Vascular Pharmacology</i> , 2005 , 42, 125-40	5.9	30
367	Cardiovascular Disease Risk Factors in Ghana during the Rural-to-Urban Transition: A Cross-Sectional Study. <i>PLoS ONE</i> , 2016 , 11, e0162753	3.7	30
366	Genome-wide association study on plasma levels of midregional-proadrenomedullin and C-terminal-pro-endothelin-1. <i>Hypertension</i> , 2013 , 61, 602-8	8.5	28
365	Systematic analysis of chromatin interactions at disease associated loci links novel candidate genes to inflammatory bowel disease. <i>Genome Biology</i> , 2016 , 17, 247	18.3	28
364	Risk factors for incident heart failure in age- and sex-specific strata: a population-based cohort using linked electronic health records. <i>European Journal of Heart Failure</i> , 2019 , 21, 1197-1206	12.3	28
363	Adverse Drug Reactions to Guideline-Recommended Heart Failure Drugs in Women: A Systematic Review of the Literature. <i>JACC: Heart Failure</i> , 2019 , 7, 258-266	7.9	27
362	The impact of susceptibility loci for coronary artery disease on other vascular domains and recurrence risk. <i>European Heart Journal</i> , 2013 , 34, 2896-904	9.5	27
361	Incremental value of a genetic risk score for the prediction of new vascular events in patients with clinically manifest vascular disease. <i>Atherosclerosis</i> , 2015 , 239, 451-8	3.1	27
360	Epistatic effects of polymorphisms in genes from the renin-angiotensin, bradykinin, and fibrinolytic systems on plasma t-PA and PAI-1 levels. <i>Genomics</i> , 2007 , 89, 362-9	4.3	27
359	Invasive versus non-invasive management of older patients with non-ST elevation myocardial infarction (SENIOR-NSTEMI): a cohort study based on routine clinical data. <i>Lancet, The</i> , 2020 , 396, 623-634	4.0	27
358	Age-stratified and blood-pressure-stratified effects of blood-pressure-lowering pharmacotherapy for the prevention of cardiovascular disease and death: an individual participant-level data meta-analysis. <i>Lancet, The</i> , 2021 , 398, 1053-1064	4.0	27
357	Mild renal dysfunction is associated with electrocardiographic left ventricular hypertrophy. <i>American Journal of Hypertension</i> , 2005 , 18, 342-7	2.3	26

356	The gender-specific role of polymorphisms from the fibrinolytic, renin-angiotensin, and bradykinin systems in determining plasma t-PA and PAI-1 levels. <i>Thrombosis and Haemostasis</i> , 2006 , 96, 471-477	7	26
355	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. <i>Nature Genetics</i> , 2020 , 52, 1314-1332	36.3	26
354	Genetic variability in the absorption of dietary sterols affects the risk of coronary artery disease. <i>European Heart Journal</i> , 2020 , 41, 2618-2628	9.5	26
353	Intronic Polymorphisms in the CDKN2B-AS1 Gene Are Strongly Associated with the Risk of Myocardial Infarction and Coronary Artery Disease in the Saudi Population. <i>International Journal of Molecular Sciences</i> , 2016 , 17, 395	6.3	26
352	The relation between systemic inflammation and incident cancer in patients with stable cardiovascular disease: a cohort study. <i>European Heart Journal</i> , 2019 , 40, 3901-3909	9.5	25
351	Long-term outcome in men and women after CABG; results from the IMAGINE trial. <i>Atherosclerosis</i> , 2015 , 241, 284-8	3.1	25
350	Exome-chip meta-analysis identifies novel loci associated with cardiac conduction, including ADAMTS6. <i>Genome Biology</i> , 2018 , 19, 87	18.3	25
349	Sex matters to the heart: A special issue dedicated to the impact of sex related differences of cardiovascular diseases. <i>Atherosclerosis</i> , 2015 , 241, 205-7	3.1	25
348	Effects of lymphotoxin-alpha gene and galectin-2 gene polymorphisms on inflammatory biomarkers, cellular adhesion molecules and risk of coronary heart disease. <i>Clinical Science</i> , 2007 , 112, 291-8	6.5	25
347	A systematic review and meta-analysis of 130,000 individuals shows smoking does not modify the association of APOE genotype on risk of coronary heart disease. <i>Atherosclerosis</i> , 2014 , 237, 5-12	3.1	24
346	Gender differences in health-related quality of life in patients undergoing coronary angiography. <i>Open Heart</i> , 2015 , 2, e000231	3	24
345	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021 ,	50.4	24
344	Genome-wide associations for benign prostatic hyperplasia reveal a genetic correlation with serum levels of PSA. <i>Nature Communications</i> , 2018 , 9, 4568	17.4	24
343	Determinants of angiotensin-converting enzyme inhibitor (ACEI) intolerance and angioedema in the UK Clinical Practice Research Datalink. <i>British Journal of Clinical Pharmacology</i> , 2016 , 82, 1647-1659	3.8	23
342	High resolution systematic digital histological quantification of cardiac fibrosis and adipose tissue in phospholamban p.Arg14del mutation associated cardiomyopathy. <i>PLoS ONE</i> , 2014 , 9, e94820	3.7	23
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91	Real-time imputation of missing predictor values in clinical practice. <i>European Heart Journal Digital Health</i> , 2021 , 2, 154-164	2.3	2
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65	Simultaneous pulmonary, cerebral and coronary emboli. <i>International Journal of Cardiology</i> , 2012 , 157, e18-20	3.2	1
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61	Transcriptomic-based clustering of advanced atherosclerotic plaques identifies subgroups of plaques with differential underlying biology that associate with clinical presentation		1
60	Discovering patterns of pleiotropy in genome-wide association studies		1
59	Genome wide association analysis in dilated cardiomyopathy reveals two new key players in systolic heart failure on chromosome 3p25.1 and 22q11.23		1
58	Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9		1
57	Age at menarche and heart failure risk: The EPIC-NL study. <i>Maturitas</i> , 2020 , 131, 34-39	5	1
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42	Comparing clinical performance of current implantable cardioverter-defibrillator implantation recommendations in arrhythmogenic right ventricular cardiomyopathy. <i>Europace</i> , 2021 ,	3.9	1
41	Sex-dependent gene regulation of human atherosclerotic plaques by DNA methylation and transcriptome integration points to smooth muscle cell involvement in women		1
40	Trends for Readmission and Mortality After Heart Failure Hospitalisation in Malaysia, 2007 to 2016.. <i>Global Heart</i> , 2022 , 17, 20	2.9	1
39	Electrocardiogram-based mortality prediction in patients with COVID-19 using machine learning.. <i>Netherlands Heart Journal</i> , 2022 , 1	2.2	1
38	High-frequency metabolite profiling and the incidence of recurrent cardiac events in patients with post-acute coronary syndrome. <i>Biomarkers</i> , 2020 , 25, 235-240	2.6	0
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35	The role of cognitive and brain reserve in memory decline and atrophy rate in mid and late-life: The SMART-MR study.. <i>Cortex</i> , 2022 , 148, 204-214	3.8	0
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31	Echocardiographic Deformation Imaging for Early Detection of Genetic Cardiomyopathies: JACC Review Topic of the Week.. <i>Journal of the American College of Cardiology</i> , 2022 , 79, 594-608	15.1	○
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29	Temporal trends in heart failure medication prescription in a population-based cohort study. <i>BMJ Open</i> , 2021 , 11, e043290	3	○
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26	A novel risk model for predicting potentially life-threatening arrhythmias in non-ischemic dilated cardiomyopathy (DCM-SVA risk). <i>International Journal of Cardiology</i> , 2021 , 339, 75-82	3.2	○
25	Elucidating mechanisms of genetic cross-disease associations at the PROCR vascular disease locus.. <i>Nature Communications</i> , 2022 , 13, 1222	17.4	○
24	Learning from individualised variation for evidence generation within a learning health system.. <i>British Journal of Anaesthesia</i> , 2022 ,	5.4	○
23	The benefit of vaccination against COVID-19 outweighs the potential risk of myocarditis and pericarditis.. <i>Netherlands Heart Journal</i> , 2022 , 30, 190	2.2	○
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