

# Per Eriksson

## List of Publications by Year in descending order

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Version: 2024-02-01

135  
papers

12,259  
citations

87888

38  
h-index

29157

104  
g-index

140  
all docs

140  
docs citations

140  
times ranked

22266  
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	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196.	27.8	1,328
3	Allele-specific increase in basal transcription of the plasminogen-activator inhibitor 1 gene is associated with myocardial infarction.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1995, 92, 1851-1855.	7.1	733
4	Association between alcohol and cardiovascular disease: Mendelian randomisation analysis based on individual participant data. <i>BMJ, The</i> , 2014, 349, g4164-g4164.	6.0	528
5	Progression of Coronary Atherosclerosis Is Associated with a Common Genetic Variant of the Human Stromelysin-1 Promoter Which Results in Reduced Gene Expression. <i>Journal of Biological Chemistry</i> , 1996, 271, 13055-13060.	3.4	437
6	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. <i>Nature Genetics</i> , 2016, 48, 1171-1184.	21.4	362
7	A Common Functional Polymorphism (C->A Substitution at Position -863) in the Promoter Region of the Tumour Necrosis Factor- $\alpha$ (TNF- $\alpha$ ) Gene Associated With Reduced Circulating Levels of TNF- $\alpha$ . <i>Human Molecular Genetics</i> , 1999, 8, 1443-1449.	2.9	307
8	Positional identification of TNFSF4, encoding OX40 ligand, as a gene that influences atherosclerosis susceptibility. <i>Nature Genetics</i> , 2005, 37, 365-372.	21.4	264
9	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.	6.2	222
10	Adenosine-to-inosine RNA editing controls cathepsin S expression in atherosclerosis by enabling HuR-mediated post-transcriptional regulation. <i>Nature Medicine</i> , 2016, 22, 1140-1150.	30.7	222
11	NLRP3 Inflammasome Expression and Activation in Human Atherosclerosis. <i>Journal of the American Heart Association</i> , 2016, 5, .	3.7	220
12	Mapping of 79 loci for 83 plasma protein biomarkers in cardiovascular disease. <i>PLoS Genetics</i> , 2017, 13, e1006706.	3.5	194
13	miR-24 limits aortic vascular inflammation and murine abdominal aneurysm development. <i>Nature Communications</i> , 2014, 5, 5214.	12.8	187
14	H19 Induces Abdominal Aortic Aneurysm Development and Progression. <i>Circulation</i> , 2018, 138, 1551-1568.	1.6	169
15	Influences of matrix metalloproteinase-3 gene variation on extent of coronary atherosclerosis and risk of myocardial infarction. <i>Journal of the American College of Cardiology</i> , 2003, 41, 2130-2137.	2.8	132
16	A Common Functional Polymorphism in the Promoter Region of the Microsomal Triglyceride Transfer Protein Gene Influences Plasma LDL Levels. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 1998, 18, 756-761.	2.4	130
17	Secretory Phospholipase A2-IIA and Cardiovascular Disease. <i>Journal of the American College of Cardiology</i> , 2013, 62, 1966-1976.	2.8	115
18	Serum matrix metalloproteinase-3 concentration is influenced by MMP-3 -1612 5A/6A promoter genotype and associated with myocardial infarction. <i>Journal of Internal Medicine</i> , 2005, 258, 411-419.	6.0	113

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19	Association of Genetic Risk Variants With Expression of Proximal Genes Identifies Novel Susceptibility Genes for Cardiovascular Disease. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 365-373.	5.1	103
20	ROBO4 variants predispose individuals to bicuspid aortic valve and thoracic aortic aneurysm. <i>Nature Genetics</i> , 2019, 51, 42-50.	21.4	101
21	MicroRNA-210 Enhances Fibrous Cap Stability in Advanced Atherosclerotic Lesions. <i>Circulation Research</i> , 2017, 120, 633-644.	4.5	98
22	Upregulation of the 5-Lipoxygenase Pathway in Human Aortic Valves Correlates With Severity of Stenosis and Leads to Leukotriene-Induced Effects on Valvular Myofibroblasts. <i>Circulation</i> , 2011, 123, 1316-1325.	1.6	92
23	Genome-wide analysis yields new loci associating with aortic valve stenosis. <i>Nature Communications</i> , 2018, 9, 987.	12.8	91
24	Protein-altering and regulatory genetic variants near GATA4 implicated in bicuspid aortic valve. <i>Nature Communications</i> , 2017, 8, 15481.	12.8	90
25	Human Evidence That the Cystatin C Gene Is Implicated in Focal Progression of Coronary Artery Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2004, 24, 551-557.	2.4	88
26	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. <i>Nature Communications</i> , 2021, 12, 24.	12.8	87
27	Unraveling Divergent Gene Expression Profiles in Bicuspid and Tricuspid Aortic Valve Patients with Thoracic Aortic Dilatation: The ASAP Study. <i>Molecular Medicine</i> , 2011, 17, 1365-1373.	4.4	81
28	Biomechanical Properties of the Thoracic Aneurysmal Wall: Differences Between Bicuspid Aortic Valve and Tricuspid Aortic Valve Patients. <i>Annals of Thoracic Surgery</i> , 2014, 98, 65-71.	1.3	78
29	Genetic Variants in LRP1 and ULK4 Are Associated with Acute Aortic Dissections. <i>American Journal of Human Genetics</i> , 2016, 99, 762-769.	6.2	73
30	Phenotypic Modulation of Smooth Muscle Cells in Atherosclerosis Is Associated With Downregulation of <i>LMOD1</i> , <i>SYNPO2</i> , <i>PDLIM7</i> , <i>PLN</i> , and <i>SYNM</i> . <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, 1947-1961.	2.4	64
31	Impaired Splicing of Fibronectin Is Associated With Thoracic Aortic Aneurysm Formation in Patients With Bicuspid Aortic Valve. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2011, 31, 691-697.	2.4	48
32	Functional Analysis of a Novel Genome-Wide Association Study Signal in <i>SMAD3</i> That Confers Protection From Coronary Artery Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, 972-983.	2.4	48
33	Identifying the susceptibility genes for coronary artery disease: from hyperbole through doubt to cautious optimism. <i>Journal of Internal Medicine</i> , 2008, 263, 538-552.	6.0	47
34	Identification of the <i>BCAR1-CFDP1-TMEM170A</i> Locus as a Determinant of Carotid Intima-Media Thickness and Coronary Artery Disease Risk. <i>Circulation: Cardiovascular Genetics</i> , 2012, 5, 656-665.	5.1	47
35	Prevention of radiotherapy-induced arterial inflammation by interleukin-1 blockade. <i>European Heart Journal</i> , 2019, 40, 2495-2503.	2.2	44
36	Perilipin 5 is protective in the ischemic heart. <i>International Journal of Cardiology</i> , 2016, 219, 446-454.	1.7	43

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37	Association of TERC and OBFC1 Haplotypes with Mean Leukocyte Telomere Length and Risk for Coronary Heart Disease. <i>PLoS ONE</i> , 2013, 8, e83122.	2.5	42
38	Allele-specific MMP-3 transcription under in vivo conditions. <i>Biochemical and Biophysical Research Communications</i> , 2006, 348, 1150-1156.	2.1	39
39	Relationship between $\beta$ -2 adrenoceptor gene haplotypes and adipocyte lipolysis in women. <i>International Journal of Obesity</i> , 2004, 28, 185-190.	3.4	38
40	Human genetic evidence that OX40 is implicated in myocardial infarction. <i>Biochemical and Biophysical Research Communications</i> , 2006, 339, 1001-1006.	2.1	38
41	PCSK6 Is a Key Protease in the Control of Smooth Muscle Cell Function in Vascular Remodeling. <i>Circulation Research</i> , 2020, 126, 571-585.	4.5	38
42	ATG16L1 Expression in Carotid Atherosclerotic Plaques Is Associated With Plaque Vulnerability. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, 1226-1235.	2.4	37
43	Genetic approach to the role of cysteine proteases in the expansion of abdominal aortic aneurysms. <i>British Journal of Surgery</i> , 2003, 91, 86-89.	0.3	36
44	Increased Arterial Blood Pressure and Vascular Remodeling in Mice Lacking Salt-Inducible Kinase 1 (SIK1). <i>Circulation Research</i> , 2015, 116, 642-652.	4.5	36
45	Mesenchymal state of intimal cells may explain higher propensity to ascending aortic aneurysm in bicuspid aortic valves. <i>Scientific Reports</i> , 2016, 6, 35712.	3.3	36
46	Common Genetic Determinants of Lung Function, Subclinical Atherosclerosis and Risk of Coronary Artery Disease. <i>PLoS ONE</i> , 2014, 9, e104082.	2.5	36
47	Aneurysm Development in Patients With a Bicuspid Aortic Valve Is Not Associated With Transforming Growth Factor- $\beta$ Activation. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, 973-980.	2.4	35
48	Genome-wide association study with additional genetic and post-transcriptional analyses reveals novel regulators of plasma factor XI levels. <i>Human Molecular Genetics</i> , 2017, 26, ddw401.	2.9	35
49	Integrative studies implicate matrix metalloproteinase-12 as a culprit gene for large artery atherosclerotic stroke. <i>Journal of Internal Medicine</i> , 2017, 282, 429-444.	6.0	34
50	Imatinib treatment attenuates growth and inflammation of angiotensin II induced abdominal aortic aneurysm. <i>Atherosclerosis</i> , 2016, 249, 101-109.	0.8	33
51	Iron alters valvular interstitial cell function and is associated with calcification in aortic stenosis. <i>European Heart Journal</i> , 2016, 37, 3532-3535.	2.2	32
52	Effect of macrophage differentiation and exposure to mildly oxidized LDL on the proteolytic repertoire of THP-1 monocytes. <i>Journal of Lipid Research</i> , 2004, 45, 1768-1776.	4.2	30
53	TRIF adaptor signaling is important in abdominal aortic aneurysm formation. <i>Atherosclerosis</i> , 2015, 241, 561-568.	0.8	30
54	Genotype-phenotype relationships in an investigation of the role of proteases in abdominal aortic aneurysm expansion. <i>British Journal of Surgery</i> , 2005, 92, 1372-1376.	0.3	29

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55	The mirâ€200 family regulates key pathogenic events in ascending aortas of individuals with bicuspid aortic valves. <i>Journal of Internal Medicine</i> , 2019, 285, 102-114.	6.0	29
56	Elastic Properties of the Descending Aorta in Patients with a Bicuspid or Tricuspid Aortic Valve and Aortic Valvular Disease. <i>Journal of the American Society of Echocardiography</i> , 2014, 27, 393-404.	2.8	25
57	Altered DNA methylation indicates an oscillatory flow mediated epithelial-to-mesenchymal transition signature in ascending aorta of patients with bicuspid aortic valve. <i>Scientific Reports</i> , 2018, 8, 2777.	3.3	25
58	A serum 25-hydroxyvitamin D concentration-associated genetic variant in DHCR7 interacts with type 2 diabetes status to influence subclinical atherosclerosis (measured by carotid intimaâ€“media) Tj ETQ0 0 0 rgBT /Overlock 104f 50 617	10.4	104
59	Differential expression of sex hormone receptors in abdominal aortic aneurysms. <i>Maturitas</i> , 2017, 96, 39-44.	2.4	24
60	Copy number variation analysis in bicuspid aortic valve-related aortopathy identifies TBX20 as a contributing gene. <i>European Journal of Human Genetics</i> , 2019, 27, 1033-1043.	2.8	24
61	Genetic Variants of Tumor Necrosis Factor Superfamily, Member 4 (TNFSF4), and Risk of Incident Atherothrombosis and Venous Thromboembolism. <i>Clinical Chemistry</i> , 2008, 54, 833-840.	3.2	23
62	A Common Polymorphism in the Promoter Region of the TNFSF4 Gene Is Associated with Lower Allele-Specific Expression and Risk of Myocardial Infarction. <i>PLoS ONE</i> , 2011, 6, e17652.	2.5	22
63	Allele-specific chromatin remodeling of the tumor necrosis factor-Î± promoter. <i>Biochemical and Biophysical Research Communications</i> , 2006, 351, 777-783.	2.1	20
64	Hydrogen peroxide induces mRNA for tumour necrosis factor Î± in human endothelial cells. <i>Free Radical Research</i> , 1999, 31, 503-512.	3.3	19
65	Aortic valve type and calcification as assessed by transthoracic and transoesophageal echocardiography. <i>Clinical Physiology and Functional Imaging</i> , 2015, 35, 306-313.	1.2	19
66	AllelicImbalance: an R/ bioconductor package for detecting, managing, and visualizing allele expression imbalance data from RNA sequencing. <i>BMC Bioinformatics</i> , 2015, 16, 194.	2.6	19
67	The composition of collagen in the aneurysm wall of men and women. <i>Journal of Vascular Surgery</i> , 2017, 66, 579-585.e1.	1.1	19
68	Notch, BMP and WNT/Î²-catenin network is impaired in endothelial cells of the patients with thoracic aortic aneurysm. <i>Atherosclerosis Supplements</i> , 2018, 35, e6-e13.	1.2	19
69	Integrated Human Evaluation of the Lysophosphatidic Acid Pathway as a Novel Therapeutic Target in Atherosclerosis. <i>Molecular Therapy - Methods and Clinical Development</i> , 2018, 10, 17-28.	4.1	18
70	Subclinical atherosclerosis and its progression are modulated by <i>PLIN2</i> through a feedâ€“forward loop between LXR and autophagy. <i>Journal of Internal Medicine</i> , 2019, 286, 660-675.	6.0	18
71	Tunica-Specific Transcriptome of Abdominal Aortic Aneurysm and the Effect of Intraluminal Thrombus, Smoking, and Diameter Growth Rate. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 2700-2713.	2.4	18
72	Lack of Salt-Inducible Kinase 2 (SIK2) Prevents the Development of Cardiac Hypertrophy in Response to Chronic High-Salt Intake. <i>PLoS ONE</i> , 2014, 9, e95771.	2.5	16

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73	Cysteinyl leukotriene receptor 1 antagonism prevents experimental abdominal aortic aneurysm. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 1907-1912.	7.1	16
74	DNA methylation age is associated with an altered hemostatic profile in a multiethnic meta-analysis. <i>Blood</i> , 2018, 132, 1842-1850.	1.4	16
75	Upregulated Autophagy in Calcific Aortic Valve Stenosis Confers Protection of Valvular Interstitial Cells. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1486.	4.1	16
76	Isochromosome 17 in a Patient with a Myeloproliferative Disorder Terminating in Eosinophilic Leukemia. <i>Acta Medica Scandinavica</i> , 1979, 206, 321-325.	0.0	15
77	Differences in Elastin and Elastolytic Enzymes between Men and Women with Abdominal Aortic Aneurysm. <i>Aorta</i> , 2014, 2, 179-185.	0.5	15
78	Elevated Adiponectin Levels Suppress Perivascular and Aortic Inflammation and Prevent AngII-induced Advanced Abdominal Aortic Aneurysms. <i>Scientific Reports</i> , 2016, 6, 31414.	3.3	15
79	Relative survival after aortic valve surgery in patients with bicuspid aortic valves. <i>Heart</i> , 2021, 107, 1167-1172.	2.9	15
80	Ascending aortic dilatation is rarely associated with coronary artery disease regardless of aortic valve morphology. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2014, 148, 2973-2980.e1.	0.8	14
81	Endothelial/Epithelial Mesenchymal Transition in Ascending Aortas of Patients With Bicuspid Aortic Valve. <i>Frontiers in Cardiovascular Medicine</i> , 2019, 6, 182.	2.4	14
82	Altered Protein Composition of Subcutaneous Adipose Tissue in Chronic Kidney Disease. <i>Kidney International Reports</i> , 2017, 2, 1208-1218.	0.8	13
83	High-Resolution Regulatory Maps Connect Vascular Risk Variants to Disease-Related Pathways. <i>Circulation Genomic and Precision Medicine</i> , 2019, 12, e002353.	3.6	13
84	TLR7 Expression Is Associated with M2 Macrophage Subset in Calcific Aortic Valve Stenosis. <i>Cells</i> , 2020, 9, 1710.	4.1	13
85	Preylcysteine oxidase 1, an emerging player in atherosclerosis. <i>Communications Biology</i> , 2021, 4, 1109.	4.4	13
86	Plaque Evaluation by Ultrasound and Transcriptomics Reveals BCLAF1 as a Regulator of Smooth Muscle Cell Lipid Transdifferentiation in Atherosclerosis. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2022, 42, 659-676.	2.4	12
87	Association Between Thoracic Aortic Disease and Inguinal Hernia. <i>Journal of the American Heart Association</i> , 2014, 3, .	3.7	11
88	ClusterSignificance: a bioconductor package facilitating statistical analysis of class cluster separations in dimensionality reduced data. <i>Bioinformatics</i> , 2017, 33, 3126-3128.	4.1	11
89	FADS1 (Fatty Acid Desaturase 1) Genotype Associates With Aortic Valve FADS mRNA Expression, Fatty Acid Content and Calcification. <i>Circulation Genomic and Precision Medicine</i> , 2020, 13, e002710.	3.6	11
90	The glucocorticoid receptor acts as an antirepressor in receptor-dependent in vitro transcription. <i>FEBS Journal</i> , 1993, 215, 505-511.	0.2	10

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91	Vessel wall morphology is equivalent for different artery types and localizations of advanced human aneurysms. <i>Histochemistry and Cell Biology</i> , 2017, 148, 425-433.	1.7	10
92	Transcriptomic profiling of experimental arterial injury reveals new mechanisms and temporal dynamics in vascular healing response. <i>JVS Vascular Science</i> , 2020, 1, 13-27.	1.1	10
93	Sex hormones in men with abdominal aortic aneurysm. <i>Journal of Vascular Surgery</i> , 2021, 74, 2023-2029.	1.1	10
94	Ceramides are associated with inflammatory processes in human mediastinal adipose tissue. <i>Nutrition, Metabolism and Cardiovascular Diseases</i> , 2014, 24, 124-131.	2.6	9
95	Effects of the coronary artery disease associated LPA and 9p21 loci on risk of aortic valve stenosis. <i>International Journal of Cardiology</i> , 2019, 276, 212-217.	1.7	9
96	Promoter anchored interaction landscape of THP-1 macrophages captures early immune response processes. <i>Cellular Immunology</i> , 2020, 355, 104148.	3.0	9
97	Neutrophil to lymphocyte ratio is not related to carotid atherosclerosis progression and cardiovascular events in the primary prevention of cardiovascular disease: Results from the IMPROVE study. <i>BioFactors</i> , 2021, , .	5.4	9
98	Neutrophil Elastase-Derived Fibrin Degradation Products Indicate Presence of Abdominal Aortic Aneurysms and Correlate with Intraluminal Thrombus Volume. <i>Thrombosis and Haemostasis</i> , 2018, 118, 329-339.	3.4	8
99	&lt;p&gt;Molecular Imaging of Inflammation in a Mouse Model of Atherosclerosis Using a Zirconium-89-Labeled Probe&lt;p&gt;. <i>International Journal of Nanomedicine</i> , 2020, Volume 15, 6137-6152.	6.7	8
100	The overlap of genetic susceptibility to schizophrenia and cardiometabolic disease can be used to identify metabolically different groups of individuals. <i>Scientific Reports</i> , 2021, 11, 632.	3.3	8
101	The glucocorticoid receptor in homodimeric and monomeric form visualised by electron microscopy. <i>Journal of Structural Biology</i> , 1991, 107, 48-55.	2.8	7
102	Aneurysm Development in Patients With Bicuspid Aortic Valve (BAV): Possible Connection to Repair Deficiency?. <i>Aorta</i> , 2013, 1, 13-22.	0.5	7
103	Elevated circulating fasting glucagon-like peptide-1 in surgical patients with aortic valve disease and diabetes. <i>Diabetology and Metabolic Syndrome</i> , 2017, 9, 79.	2.7	7
104	New candidate genes for ST â€elevation myocardial infarction. <i>Journal of Internal Medicine</i> , 2020, 287, 66-77.	6.0	7
105	Functional Analysis of the Coronary Heart Disease Risk Locus on Chromosome 21q22. <i>Disease Markers</i> , 2017, 2017, 1-10.	1.3	6
106	A Genome Wide Association Study on plasma FV levels identified PLXDC2 as a new modifier of the coagulation process. <i>Journal of Thrombosis and Haemostasis</i> , 2019, 17, 1808-1814.	3.8	6
107	Interleukin 6 trans-signalling and the risk of future cardiovascular events in men and women. <i>Open Heart</i> , 2021, 8, e001694.	2.3	6
108	Intima-media thickness of the descending aorta in patients with bicuspid aortic valve. <i>IJC Heart and Vasculature</i> , 2016, 11, 74-79.	1.1	5

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109	Genetic Variants Associated with Non-Alcoholic Fatty Liver Disease Do Not Associate with Measures of Sub-Clinical Atherosclerosis: Results from the IMPROVE Study. <i>Genes</i> , 2020, 11, 1243.	2.4	5
110	Arginase 1 is upregulated at admission in patients with ST-elevation myocardial infarction. <i>Journal of Internal Medicine</i> , 2021, 290, 1061-1070.	6.0	5
111	The tyrosine kinase inhibitor nilotinib targets discoidin domain receptor 2 in calcific aortic valve stenosis.. <i>British Journal of Pharmacology</i> , 0, , .	5.4	5
112	Reply to "Lack of support for association between common variation in TNFSF4 and myocardial infarction in a German population" Nature Genetics, 2008, 40, 1387-1388.	21.4	4
113	Pre- and postoperative left atrial and ventricular volumetric and deformation analyses in severe aortic regurgitation. <i>Cardiovascular Ultrasound</i> , 2021, 19, 14.	1.6	4
114	Cardiac expression of the microsomal triglyceride transport protein protects the heart function during ischemia. <i>Journal of Molecular and Cellular Cardiology</i> , 2019, 137, 1-8.	1.9	3
115	Sex Differences in Aortopathy and Valve Diseases Among Patients Undergoing Cardiac Surgical Procedure. <i>Annals of Thoracic Surgery</i> , 2022, , .	1.3	3
116	A novel anti-inflammatory role links the CARS2 locus to protection from coronary artery disease. <i>Atherosclerosis</i> , 2022, 348, 8-15.	0.8	3
117	Auxilin is a novel susceptibility gene for congenital heart block which directly impacts fetal heart function. <i>Annals of the Rheumatic Diseases</i> , 2022, 81, 1151-1161.	0.9	3
118	P329Regulation of LTBP expression as a modulator of TGFb availability in patients with BAV. <i>Cardiovascular Research</i> , 2018, 114, S84-S84.	3.8	0
119	P3674New candidate genes for plaque rupture in myocardial infarction. <i>European Heart Journal</i> , 2018, 39, .	2.2	0
120	P1217Zirconium-89 labelled probe for molecular imaging of inflammation in experimental atherosclerosis. <i>European Heart Journal</i> , 2019, 40, .	2.2	0
121	Comparison of quantitative trait loci methods: Total expression and allelic imbalance method in brain RNA-seq. <i>PLoS ONE</i> , 2019, 14, e0217765.	2.5	0
122	Abstract 397: Analysis of Cell Phenotype in Relation to TGFÎ² Treatment of Aortic Smooth Muscle Cells and Myofibroblasts Isolated from Aortas and Valves of Thoracic Aortic Aneurysm Patients with a Tricuspid or a Bicuspid Valve. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2012, 32, .	2.4	0
123	Abstract 65: Interleukin-6 Signaling and Abdominal Aortic Aneurysm. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2012, 32, .	2.4	0
124	Abstract 284: microRNAs are Novel Plasma Biomarkers for Diagnosis and Prognosis of Abdominal Aortic Aneurysm Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2013, 33, .	2.4	0
125	Abstract 539: MicroRNA-27b Regulates Salt-Inducible Kinase 1 (SIK1) in Vascular Fibrosis. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2013, 33, .	2.4	0
126	Abstract 277: Downregulation of PDGF-D is Associated with Increased Collagen Production in Abdominal Aortic Aneurysm. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2013, 33, .	2.4	0

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127	Abstract 267: CD93: A Novel Myocardial Infarction- Associated Protein with Glucose Regulatory Properties in Humans and Mice. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, .	2.4	0
128	Abstract 318: Matrix Metalloproteinase 12 is Causally Implicated in Cardiovascular Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, .	2.4	0
129	Abstract 636: Accelerated Atherosclerosis in the Context of Rheumatoid Arthritis. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, .	2.4	0
130	Abstract 19: Role of Adenosine-to-Inosine RNA Editing of <i>Alu</i> Elements in Human Vascular Inflammatory Diseases. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, .	2.4	0
131	Abstract 173: Proprotein Convertase Subtilisin/Kexin Type 6 is a Key Protease in the Control of Smooth Muscle Cell Function in Vascular Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, .	2.4	0
132	Abstract 552: End Stage Human Aneurysm Disease in Different Arterial Positions is Similar, Aneurysm Induction in Mouse Models is Not. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, .	2.4	0
133	Abstract P112: Prediction of the Development of Aortopathy in Patients With Bicuspid Aortic Valves. <i>Circulation</i> , 2019, 139, .	1.6	0
134	Abstract 467: PCSK6 Is Upregulated in Vascular Diseases Characterized by Inflammation and Smooth Muscle Cell Proliferation. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, .	2.4	0
135	Abstract 367: <i>Pcsk6</i> Is a Key Protease Modulating Smooth Muscle Cell Activation in Vascular Remodeling and Plaque Vulnerability. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, .	2.4	0