

Lasse Folkersen

List of Publications by Year in descending order

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Version: 2024-02-01

113
papers

15,674
citations

76196

40
h-index

34900

98
g-index

132
all docs

132
docs citations

132
times ranked

28244
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. | 13.7 | 3,823 |
| 2 | Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186. | 9.4 | 1,818 |
| 3 | Large-scale association analysis identifies new risk loci for coronary artery disease. <i>Nature Genetics</i> , 2013, 45, 25-33. | 9.4 | 1,439 |
| 4 | New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196. | 13.7 | 1,328 |
| 5 | Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. <i>Nature Genetics</i> , 2012, 44, 991-1005. | 9.4 | 746 |
| 6 | Association between alcohol and cardiovascular disease: Mendelian randomisation analysis based on individual participant data. <i>BMJ</i> , The, 2014, 349, g4164-g4164. | 3.0 | 528 |
| 7 | CD49a Expression Defines Tissue-Resident CD8 + T Cells Poised for Cytotoxic Function in Human Skin. <i>Immunity</i> , 2017, 46, 287-300. | 6.6 | 465 |
| 8 | 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. | 9.4 | 362 |
| 9 | Genome-Wide Association Identifies Nine Common Variants Associated With Fasting Proinsulin Levels and Provides New Insights Into the Pathophysiology of Type 2 Diabetes. <i>Diabetes</i> , 2011, 60, 2624-2634. | 0.3 | 335 |
| 10 | Genomic and drug target evaluation of 90 cardiovascular proteins in 30,931 individuals. <i>Nature Metabolism</i> , 2020, 2, 1135-1148. | 5.1 | 327 |
| 11 | Meta-analysis of 65,734 Individuals Identifies TSPAN15 and SLC44A2 as Two Susceptibility Loci for Venous Thromboembolism. <i>American Journal of Human Genetics</i> , 2015, 96, 532-542. | 2.6 | 222 |
| 12 | NLRP3 Inflammasome Expression and Activation in Human Atherosclerosis. <i>Journal of the American Heart Association</i> , 2016, 5, . | 1.6 | 220 |
| 13 | Mapping of 79 loci for 83 plasma protein biomarkers in cardiovascular disease. <i>PLoS Genetics</i> , 2017, 13, e1006706. | 1.5 | 194 |
| 14 | Abdominal Aortic Aneurysm Is Associated with a Variant in Low-Density Lipoprotein Receptor-Related Protein 1. <i>American Journal of Human Genetics</i> , 2011, 89, 619-627. | 2.6 | 185 |
| 15 | Genome-wide association study identifies a sequence variant within the DAB2IP gene conferring susceptibility to abdominal aortic aneurysm. <i>Nature Genetics</i> , 2010, 42, 692-697. | 9.4 | 181 |
| 16 | Meta-Analysis of Genome-Wide Association Studies for Abdominal Aortic Aneurysm Identifies Four New Disease-Specific Risk Loci. <i>Circulation Research</i> , 2017, 120, 341-353. | 2.0 | 166 |
| 17 | Relationship between CAD Risk Genotype in the Chromosome 9p21 Locus and Gene Expression. Identification of Eight New ANRIL Splice Variants. <i>PLoS ONE</i> , 2009, 4, e7677. | 1.1 | 145 |
| 18 | Interleukin-6 receptor pathways in abdominal aortic aneurysm. <i>European Heart Journal</i> , 2013, 34, 3707-3716. | 1.0 | 143 |

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|----|---|------|-----------|
| 19 | Multiethnic Meta-Analysis of Genome-Wide Association Studies in >100 000 Subjects Identifies 23 Fibrinogen-Associated Loci but No Strong Evidence of a Causal Association Between Circulating Fibrinogen and Cardiovascular Disease. <i>Circulation</i> , 2013, 128, 1310-1324. | 1.6 | 128 |
| 20 | Prediction of Ischemic Events on the Basis of Transcriptomic and Genomic Profiling in Patients Undergoing Carotid Endarterectomy. <i>Molecular Medicine</i> , 2012, 18, 669-675. | 1.9 | 118 |
| 21 | Secretory Phospholipase A2-IIA and Cardiovascular Disease. <i>Journal of the American College of Cardiology</i> , 2013, 62, 1966-1976. | 1.2 | 115 |
| 22 | Gene expression signatures, pathways and networks in carotid atherosclerosis. <i>Journal of Internal Medicine</i> , 2016, 279, 293-308. | 2.7 | 114 |
| 23 | 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 |
| 24 | Genome-wide association analysis of self-reported events in 6135 individuals and 252 827 controls identifies 8 loci associated with thrombosis. <i>Human Molecular Genetics</i> , 2016, 25, 1867-1874. | 1.4 | 103 |
| 25 | Evaluation of polygenic prediction methodology within a reference-standardized framework. <i>PLoS Genetics</i> , 2021, 17, e1009021. | 1.5 | 99 |
| 26 | Toll-Like Receptor 7 Protects From Atherosclerosis by Constraining Inflammatory Macrophage Activation. <i>Circulation</i> , 2012, 126, 952-962. | 1.6 | 92 |
| 27 | Genome-wide analysis yields new loci associating with aortic valve stenosis. <i>Nature Communications</i> , 2018, 9, 987. | 5.8 | 91 |
| 28 | Genome-wide association study for circulating levels of PAI-1 provides novel insights into its regulation. <i>Blood</i> , 2012, 120, 4873-4881. | 0.6 | 90 |
| 29 | Protein-altering and regulatory genetic variants near GATA4 implicated in bicuspid aortic valve. <i>Nature Communications</i> , 2017, 8, 15481. | 5.8 | 90 |
| 30 | Causal Effect of Plasminogen Activator Inhibitor Type 1 on Coronary Heart Disease. <i>Journal of the American Heart Association</i> , 2017, 6, . | 1.6 | 89 |
| 31 | Profiling of Atherosclerotic Lesions by Gene and Tissue Microarrays Reveals PCSK6 as a Novel Protease in Unstable Carotid Atherosclerosis. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2013, 33, 2432-2443. | 1.1 | 84 |
| 32 | 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. | 1.9 | 81 |
| 33 | Human Disease Variation in the Light of Population Genomics. <i>Cell</i> , 2019, 177, 115-131. | 13.5 | 75 |
| 34 | GLP-1 Induces Barrier Protective Expression in Brunner's Glands and Regulates Colonic Inflammation. <i>Inflammatory Bowel Diseases</i> , 2016, 22, 2078-2097. | 0.9 | 62 |
| 35 | H1N1 vaccination in Sjögren's syndrome triggers polyclonal B cell activation and promotes autoantibody production. <i>Annals of the Rheumatic Diseases</i> , 2017, 76, 1755-1763. | 0.5 | 51 |
| 36 | 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. | 1.1 | 48 |

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|----|--|-----|-----------|
| 37 | 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. | 1.1 | 48 |
| 38 | 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 |
| 39 | Impute.me: An Open-Source, Non-profit Tool for Using Data From Direct-to-Consumer Genetic Testing to Calculate and Interpret Polygenic Risk Scores. <i>Frontiers in Genetics</i> , 2020, 11, 578. | 1.1 | 47 |
| 40 | High plasma adiponectin concentration is associated with all-cause mortality in patients with carotid atherosclerosis. <i>Atherosclerosis</i> , 2012, 225, 491-496. | 0.4 | 43 |
| 41 | Genome-Wide Association Study for Circulating Tissue Plasminogen Activator Levels and Functional Follow-Up Implicates Endothelial <i>STXBP5</i> and <i>STX2</i> . <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, 1093-1101. | 1.1 | 43 |
| 42 | Reduced expression of <i>TRIM21/Ro52</i> predicts poor prognosis in diffuse large B-cell lymphoma patients with and without rheumatic disease. <i>Journal of Internal Medicine</i> , 2015, 278, 323-332. | 2.7 | 43 |
| 43 | <i>SORBS1</i> gene, a new candidate for diabetic nephropathy: results from a multi-stage genome-wide association study in patients with type 1 diabetes. <i>Diabetologia</i> , 2015, 58, 543-548. | 2.9 | 43 |
| 44 | Association of <i>TERC</i> and <i>OBFC1</i> Haplotypes with Mean Leukocyte Telomere Length and Risk for Coronary Heart Disease. <i>PLoS ONE</i> , 2013, 8, e83122. | 1.1 | 42 |
| 45 | Integration of Known DNA, RNA and Protein Biomarkers Provides Prediction of Anti-TNF Response in Rheumatoid Arthritis: Results from the COMBINE Study. <i>Molecular Medicine</i> , 2016, 22, 322-328. | 1.9 | 39 |
| 46 | The Chromosome 9p21.3 Coronary Heart Disease Risk Allele Is Associated with Altered Gene Expression in Normal Heart and Vascular Tissues. <i>PLoS ONE</i> , 2012, 7, e39574. | 1.1 | 37 |
| 47 | $\alpha 7$ Nicotinic Acetylcholine Receptor Is Expressed in Human Atherosclerosis and Inhibits Disease in Mice. <i>Brief Report. Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, 2632-2636. | 1.1 | 37 |
| 48 | Innate immune receptor NOD2 promotes vascular inflammation and formation of lipid-rich necrotic cores in hypercholesterolemic mice. <i>European Journal of Immunology</i> , 2014, 44, 3081-3092. | 1.6 | 36 |
| 49 | Common Genetic Determinants of Lung Function, Subclinical Atherosclerosis and Risk of Coronary Artery Disease. <i>PLoS ONE</i> , 2014, 9, e104082. | 1.1 | 36 |
| 50 | A Genome-Wide Association Study Identifies <i>KNG1</i> as a Genetic Determinant of Plasma Factor XI Level and Activated Partial Thromboplastin Time. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2012, 32, 2008-2016. | 1.1 | 33 |
| 51 | Impact of Common Variation in Bone-Related Genes on Type 2 Diabetes and Related Traits. <i>Diabetes</i> , 2012, 61, 2176-2186. | 0.3 | 31 |
| 52 | Low <i>TLR7</i> gene expression in atherosclerotic plaques is associated with major adverse cardiovascular and cerebrovascular events. <i>Cardiovascular Research</i> , 2017, 113, 30-39. | 1.8 | 31 |
| 53 | Network-based Analysis of Genome Wide Association Data Provides Novel Candidate Genes for Lipid and Lipoprotein Traits. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 3398-3408. | 2.5 | 28 |
| 54 | Genetic Variation in <i>SULF2</i> Is Associated with Postprandial Clearance of Triglyceride-Rich Remnant Particles and Triglyceride Levels in Healthy Subjects. <i>PLoS ONE</i> , 2013, 8, e79473. | 1.1 | 28 |

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|----|---|-----|-----------|
| 55 | MicroRNA 486-3P as a stability marker in acute coronary syndrome. Bioscience Reports, 2016, 36, . | 1.1 | 27 |
| 56 | <i>CARD8</i> gene encoding a protein of innate immunity is expressed in human atherosclerosis and associated with markers of inflammation. Clinical Science, 2013, 125, 401-407. | 1.8 | 26 |
| 57 | Neil3-dependent base excision repair regulates lipid metabolism and prevents atherosclerosis in ApoE-deficient mice. Scientific Reports, 2016, 6, 28337. | 1.6 | 26 |
| 58 | Discovery of new candidate genes for rheumatoid arthritis through integration of genetic association data with expression pathway analysis. Arthritis Research and Therapy, 2017, 19, 19. | 1.6 | 25 |
| 59 | Diverging Alternative Splicing Fingerprints in the Transforming Growth Factor- β Signaling Pathway Identified in Thoracic Aortic Aneurysms. Molecular Medicine, 2011, 17, 665-675. | 1.9 | 24 |
| 60 | A serum 25-hydroxyvitamin D concentration-associated genetic variant in DHCR7 interacts with type 2 diabetes status to influence subclinical atherosclerosis (measured by carotid intima-media thickness). Arteriosclerosis, Thrombosis, and Vascular Biology, 2017, 37, 1047-1053. | 0.0 | 24 |
| 61 | KLF12 Regulates Mouse NK Cell Proliferation. Journal of Immunology, 2019, 203, 981-989. | 0.4 | 24 |
| 62 | Why do people seek out polygenic risk scores for complex disorders, and how do they understand and react to results?. European Journal of Human Genetics, 2022, 30, 81-87. | 1.4 | 23 |
| 63 | Novel Genetic Approach to Investigate the Role of Plasma Secretory Phospholipase A2 (sPLA ₂) in Atherosclerosis. PLoS One, 2017, 12, e0174314. | 5.1 | 22 |
| 64 | PDE1A inhibition elicits cGMP-dependent relaxation of rat mesenteric arteries. British Journal of Pharmacology, 2017, 174, 4186-4198. | 2.7 | 22 |
| 65 | Early prediction of clinical response to anti-TNF treatment using multi-omics and machine learning in rheumatoid arthritis. Rheumatology, 2022, 61, 1680-1689. | 0.9 | 22 |
| 66 | Use of Allele-Specific FAIRE to Determine Functional Regulatory Polymorphism Using Large-Scale Genotyping Arrays. PLoS Genetics, 2012, 8, e1002908. | 1.5 | 21 |
| 67 | Identification of a novel flow-mediated gene expression signature in patients with bicuspid aortic valve. Journal of Molecular Medicine, 2013, 91, 129-139. | 1.7 | 20 |
| 68 | Genetic Landscape of the ACE2 Coronavirus Receptor. Circulation, 2022, 145, 1398-1411. | 1.6 | 20 |
| 69 | AllelicImbalance: an R/ bioconductor package for detecting, managing, and visualizing allele expression imbalance data from RNA sequencing. BMC Bioinformatics, 2015, 16, 194. | 1.2 | 19 |
| 70 | Systematic approach demonstrates enrichment of multiple interactions between non-HLA risk variants and HLA-DRB1 risk alleles in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2018, 77, 1454-1462. | 0.5 | 19 |
| 71 | Genetic stratification of depression in UK Biobank. Translational Psychiatry, 2020, 10, 163. | 2.4 | 19 |
| 72 | Identifying low density lipoprotein cholesterol associated variants in the Annexin A2 (ANXA2) gene. Atherosclerosis, 2017, 261, 60-68. | 0.4 | 18 |

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|----|---|-----|-----------|
| 73 | Mechanisms of Action of the KCa2-Negative Modulator AP30663, a Novel Compound in Development for Treatment of Atrial Fibrillation in Man. <i>Frontiers in Pharmacology</i> , 2020, 11, 610. | 1.6 | 18 |
| 74 | A tool for translating polygenic scores onto the absolute scale using summary statistics. <i>European Journal of Human Genetics</i> , 2022, 30, 339-348. | 1.4 | 18 |
| 75 | PLA2G10 Gene Variants, sPLA2 Activity, and Coronary Heart Disease Risk. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 356-362. | 5.1 | 17 |
| 76 | Functional Analysis of a Carotid Intima-Media Thickness Locus Implicates <i>BCAR1</i> and Suggests a Causal Variant. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 696-706. | 5.1 | 17 |
| 77 | Ubiquitin-specific peptidase 2 as a potential link between microRNA-125b and psoriasis. <i>British Journal of Dermatology</i> , 2017, 176, 723-731. | 1.4 | 17 |
| 78 | Functional Analysis of Two PLA2G2A Variants Associated with Secretory Phospholipase A2-IIA Levels. <i>PLoS ONE</i> , 2012, 7, e41139. | 1.1 | 16 |
| 79 | Dual roles of heparanase in human carotid plaque calcification. <i>Atherosclerosis</i> , 2019, 283, 127-136. | 0.4 | 16 |
| 80 | Endogenous control genes in complex vascular tissue samples. <i>BMC Genomics</i> , 2009, 10, 516. | 1.2 | 14 |
| 81 | High-Resolution Regulatory Maps Connect Vascular Risk Variants to Disease-Related Pathways. <i>Circulation Genomic and Precision Medicine</i> , 2019, 12, e002353. | 1.6 | 13 |
| 82 | Integration of Genetics into a Systems Model of Electrocardiographic Traits Using HumanCVD BeadChip. <i>Circulation: Cardiovascular Genetics</i> , 2012, 5, 630-638. | 5.1 | 12 |
| 83 | Translating polygenic risk scores for clinical use by estimating the confidence bounds of risk prediction. <i>Nature Communications</i> , 2021, 12, 5276. | 5.8 | 12 |
| 84 | ClusterSignificance: a bioconductor package facilitating statistical analysis of class cluster separations in dimensionality reduced data. <i>Bioinformatics</i> , 2017, 33, 3126-3128. | 1.8 | 11 |
| 85 | Imputed gene expression risk scores: a functionally informed component of polygenic risk. <i>Human Molecular Genetics</i> , 2021, 30, 727-738. | 1.4 | 11 |
| 86 | EBI3 regulates the NK cell response to mouse cytomegalovirus infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 1625-1630. | 3.3 | 10 |
| 87 | A gene-centric study of common carotid artery remodelling. <i>Atherosclerosis</i> , 2013, 226, 440-446. | 0.4 | 9 |
| 88 | Novel <i>TRAPPC11</i> Mutations in a Chinese Pedigree of Limb Girdle Muscular Dystrophy. <i>Case Reports in Genetics</i> , 2018, 2018, 1-6. | 0.1 | 9 |
| 89 | GeneRegionScan: a Bioconductor package for probe-level analysis of specific, small regions of the genome. <i>Bioinformatics</i> , 2009, 25, 1978-1979. | 1.8 | 8 |
| 90 | Human Genetic Evidence for Involvement of CD137 in Atherosclerosis. <i>Molecular Medicine</i> , 2014, 20, 456-465. | 1.9 | 8 |

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|-----|---|-----|-----------|
| 91 | Expression of CARD8 in human atherosclerosis and its regulation of inflammatory proteins in human endothelial cells. <i>Scientific Reports</i> , 2020, 10, 19108. | 1.6 | 8 |
| 92 | Aneurysm Development in Patients With Bicuspid Aortic Valve (BAV): Possible Connection to Repair Deficiency?. <i>Aorta</i> , 2013, 1, 13-22. | 0.1 | 7 |
| 93 | Applying genetics in inflammatory disease drug discovery. <i>Drug Discovery Today</i> , 2015, 20, 1176-1181. | 3.2 | 6 |
| 94 | Functional Analysis of the Coronary Heart Disease Risk Locus on Chromosome 21q22. <i>Disease Markers</i> , 2017, 2017, 1-10. | 0.6 | 6 |
| 95 | Enhanced base excision repair capacity in carotid atherosclerosis may protect nuclear DNA but not mitochondrial DNA. <i>Free Radical Biology and Medicine</i> , 2016, 97, 386-397. | 1.3 | 3 |
| 96 | A novel anti-inflammatory role links the CARS2 locus to protection from coronary artery disease. <i>Atherosclerosis</i> , 2022, 348, 8-15. | 0.4 | 3 |
| 97 | 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. | 0.5 | 3 |
| 98 | Vaccination of patients with primary Sjogren's syndrome reveals hyperreactive B cell compartment with a skewed maturation pattern. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, A67-A67. | 0.5 | 1 |
| 99 | The role of innate immune receptor nod2 in atherosclerosis. <i>Atherosclerosis</i> , 2014, 235, e20-e21. | 0.4 | 0 |
| 100 | Functional analysis of the chromosome 21q22 (gene desert) variant associated with CHD risk. <i>Atherosclerosis</i> , 2015, 241, e17. | 0.4 | 0 |
| 101 | Low TLR7 gene expression in atherosclerotic plaques is associated with major adverse cardio- and cerebrovascular events. <i>Atherosclerosis</i> , 2017, 263, e8. | 0.4 | 0 |
| 102 | Identifying LDL-C associated variants in the Annexin a2 (ANXA2) gene. <i>Atherosclerosis</i> , 2017, 263, e20. | 0.4 | 0 |
| 103 | M17 EVALUATING PREDICTIVE ABILITY OF FUNCTIONALLY INFORMED GENETIC RISK SCORES. <i>European Neuropsychopharmacology</i> , 2019, 29, S175. | 0.3 | 0 |
| 104 | M33 TRYGGVE2: PREDICTING POOR OUTCOMES IN MAJOR DEPRESSION USING REGISTER GENOMICS IN SWEDEN. <i>European Neuropsychopharmacology</i> , 2019, 29, S183-S184. | 0.3 | 0 |
| 105 | Comparison of quantitative trait loci methods: Total expression and allelic imbalance method in brain RNA-seq. <i>PLoS ONE</i> , 2019, 14, e0217765. | 1.1 | 0 |
| 106 | Photochemotherapy Induces Interferon Type III Expression via STING Pathway. <i>Cells</i> , 2020, 9, 2452. | 1.8 | 0 |
| 107 | Abstract 397: Analysis of Cell Phenotype in Relation to TGF β 2 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, . | 1.1 | 0 |
| 108 | Cholinergic signaling through the alpha 7 nicotinic receptor inhibits atherosclerosis in hypercholesterolemic mice (671.7). <i>FASEB Journal</i> , 2014, 28, 671.7. | 0.2 | 0 |

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|-----|--|-----|-----------|
| 109 | Abstract 52: The BiKE Project: Gene Expression Signatures, Pathways and Networks in Human Carotid Atherosclerosis. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, . | 1.1 | 0 |
| 110 | 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, . | 1.1 | 0 |
| 111 | The hunt for fatal myocardial infarction biomarkers: predictive circulating microRNAs. <i>Annals of Translational Medicine</i> , 2016, 4, S1-S1. | 0.7 | 0 |
| 112 | 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, . | 1.1 | 0 |
| 113 | Abstract 367: Pcsk6 Is a Key Protease Modulating Smooth Muscle Cell Activation in Vascular Remodeling and Plaque Vulnerability. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, . | 1.1 | 0 |