## Lisa de las Fuentes

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4640579/publications.pdf

Version: 2024-02-01

98 papers 4,319 citations

32 h-index 62 g-index

101 all docs

101 docs citations

times ranked

101

7783 citing authors

#	Article	IF	CITATIONS
1	Effects of Moderate and Subsequent Progressive Weight Loss on Metabolic Function and Adipose Tissue Biology in Humans with Obesity. Cell Metabolism, 2016, 23, 591-601.	16.2	592
2	Altered myocardial fatty acid and glucose metabolism in idiopathic dilated cardiomyopathy. Journal of the American College of Cardiology, 2002, 40, 271-277.	2.8	432
3	Cardiovascular Phenotype in HFpEF Patients With or Without Diabetes. Journal of the American College of Cardiology, 2014, 64, 541-549.	2.8	157
4	Myocardial Fatty Acid Metabolism. Hypertension, 2003, 41, 83-87.	2.7	141
5	Metabolic syndrome is associated with abnormal left ventricular diastolic function independent of left ventricular mass. European Heart Journal, 2006, 28, 553-559.	2.2	140
6	Effect of Moderate Diet-Induced Weight Loss and Weight Regain on Cardiovascular Structure and Function. Journal of the American College of Cardiology, 2009, 54, 2376-2381.	2.8	130
7	A Large-Scale Multi-ancestry Genome-wide Study Accounting for Smoking Behavior Identifies Multiple Significant Loci for Blood Pressure. American Journal of Human Genetics, 2018, 102, 375-400.	6.2	123
8	Multi-ancestry genome-wide gene–smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. Nature Genetics, 2019, 51, 636-648.	21.4	112
9	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. American Journal of Human Genetics, 2019, 104, 112-138.	6.2	106
10	Vitamin D Suppression of Endoplasmic Reticulum Stress Promotes an Antiatherogenic Monocyte/Macrophage Phenotype in Type 2 Diabetic Patients. Journal of Biological Chemistry, 2012, 287, 38482-38494.	3.4	96
11	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. PLoS ONE, 2018, 13, e0198166.	2.5	94
12	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. American Journal of Epidemiology, 2019, 188, 1033-1054.	3.4	85
13	Insulin-like growth factor regulation of human endometrial stromal cell function: coordinate effects on insulin-like growth factor binding protein-1, cell proliferation and prolactin secretion. Regulatory Peptides, 1993, 48, 165-177.	1.9	84
14	Timing and Causes of Readmission After Acute Heart Failure Hospitalizationâ€"Insights From the Heart Failure Network Trials. Journal of Cardiac Failure, 2016, 22, 875-883.	1.7	78
15	Timing of cardiac transplantation in patients with heart failure receiving $\hat{l}^2$ -adrenergic blockers. Journal of Heart and Lung Transplantation, 2003, 22, 1141-1148.	0.6	74
16	Williams Syndrome Predisposes to Vascular Stiffness Modified by Antihypertensive Use and Copy Number Changes in <i>NCF1</i> . Hypertension, 2014, 63, 74-79.	2.7	69
17	Effects of Phosphate Binder Therapy on Vascular Stiffness in Early-Stage Chronic Kidney Disease. American Journal of Nephrology, 2013, 38, 158-167.	3.1	65
18	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. Nature Communications, 2019, 10, 376.	12.8	64

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19	Improvements in Left Ventricular Diastolic Function After Cardiac Resynchronization Therapy Are Coupled to Response in Systolic Performance. Journal of the American College of Cardiology, 2005, 46, 2244-2249.	2.8	62
20	Multi-ancestry sleep-by-SNP interaction analysis in 126,926 individuals reveals lipid loci stratified by sleep duration. Nature Communications, 2019, 10, 5121.	12.8	62
21	Growth Factors and Decidualization in Vitro. Annals of the New York Academy of Sciences, 1994, 734, 7-18.	3.8	56
22	Gene–Smoking Interactions Identify Several Novel Blood Pressure Loci in the Framingham Heart Study. American Journal of Hypertension, 2015, 28, 343-354.	2.0	52
23	Hypertensive left ventricular hypertrophy is associated with abnormal myocardial fatty acid metabolism and myocardial efficiency. Journal of Nuclear Cardiology, 2006, 13, 369-377.	2.1	50
24	Intensification of Medication Therapy for Cardiorenal Syndrome in Acute Decompensated Heart Failure. Journal of Cardiac Failure, 2016, 22, 26-32.	1.7	48
25	Effects of Sodium Thiosulfate on Vascular Calcification in End-Stage Renal Disease: A Pilot Study of Feasibility, Safety and Efficacy. American Journal of Nephrology, 2011, 33, 131-138.	3.1	45
26	Characterization of Left Ventricular Diastolic Function in Hypertension by Use of Doppler Tissue Imaging and Color M-Mode Techniques. Journal of the American Society of Echocardiography, 2006, 19, 872-879.	2.8	44
27	Cardiac resynchronization therapy acutely improves diastolic function. Journal of the American Society of Echocardiography, 2005, 18, 216-220.	2.8	43
28	Population sequencing data reveal a compendium of mutational processes in the human germ line. Science, 2021, 373, 1030-1035.	12.6	43
29	Plasma Triglyceride Level is an Independent Predictor of Altered Left Ventricular Relaxation. Journal of the American Society of Echocardiography, 2005, 18, 1285-1291.	2.8	41
30	Recent Innovations, Modifications, and Evolution of ACC/AHA Clinical Practice Guidelines: An Update for Our Constituencies: A Report of the American College of Cardiology/American Heart Association Task Force on Clinical Practice Guidelines. Circulation, 2019, 139, e879-e886.	1.6	41
31	Genes for left ventricular hypertrophy. Current Hypertension Reports, 2004, 6, 36-41.	3.5	40
32	Clinical Outcomes After Cardiac Resynchronization Therapy: Importance of Left Ventricular Diastolic Function and Origin of Heart Failure. Journal of the American Society of Echocardiography, 2006, 19, 307-313.	2.8	39
33	A Review of the Genetics of Hypertension with a Focus on Gene-Environment Interactions. Current Hypertension Reports, 2017, 19, 23.	3.5	39
34	A Custom Correlation Coefficient (CCC) Approach for Fast Identification of Multiâ€SNP Association Patterns in Genomeâ€Wide SNPs Data. Genetic Epidemiology, 2014, 38, 610-621.	1.3	38
35	Mendelian randomization supports bidirectional causality between telomere length and clonal hematopoiesis of indeterminate potential. Science Advances, 2022, 8, eabl6579.	10.3	36
36	An Overview of the Genomics of Metabolic Syndrome. Journal of Nursing Scholarship, 2013, 45, 52-59.	2.4	33

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37	Impaired cardiac and skeletal muscle bioenergetics in children, adolescents, and young adults with Barth syndrome. Physiological Reports, 2017, 5, e13130.	1.7	33
38	2020 AHA/ACC guideline for the diagnosis and treatment of patients with hypertrophic cardiomyopathy. Journal of Thoracic and Cardiovascular Surgery, 2021, 162, e23-e106.	0.8	33
39	Role of Serotoninergic Pathways in Drug-Induced Valvular Heart Disease and Diagnostic Features by Echocardiography. Journal of the American Society of Echocardiography, 2009, 22, 883-889.	2.8	32
40	Nitrate's Effect on Activity Tolerance in Heart Failure With Preserved Ejection Fraction Trial. Circulation: Heart Failure, 2015, 8, 221-228.	3.9	31
41	Determinants of Diuretic Responsiveness and Associated Outcomes During Acute Heart Failure Hospitalization: An Analysis From the NHLBI Heart Failure Network Clinical Trials. Journal of Cardiac Failure, 2018, 24, 428-438.	1.7	31
42	A multi-ancestry genome-wide study incorporating gene–smoking interactions identifies multiple new loci for pulse pressure and mean arterial pressure. Human Molecular Genetics, 2019, 28, 2615-2633.	2.9	31
43	Left Ventricular Mass Progression despite Stable Blood Pressure and Kidney Function in Stage 3 Chronic Kidney Disease. American Journal of Nephrology, 2014, 39, 392-399.	3.1	30
44	Recent Innovations, Modifications, Âand ÂEvolution of ACC/AHA Clinical Practice Guidelines: An Update for Our ÂConstituencies. Journal of the American College of Cardiology, 2019, 73, 1990-1998.	2.8	30
45	Insulin resistance predicts endothelial dysfunction and cardiovascular risk in HIV-infected persons on long-term highly active antiretroviral therapy. Aids, 2008, 22, 849-856.	2.2	29
46	Genetic determinants of telomere length from 109,122 ancestrally diverse whole-genome sequences in TOPMed. Cell Genomics, 2022, 2, 100084.	6.5	29
47	N-terminal Pro B-type Natriuretic Peptide Levels: Correlation with Echocardiographically Determined Left Ventricular Diastolic Function in an Ambulatory Cohort. Journal of the American Society of Echocardiography, 2006, 19, 1017-1025.	2.8	27
48	Pathway-based genome-wide association analysis of coronary heart disease identifies biologically important gene sets. European Journal of Human Genetics, 2012, 20, 1168-1173.	2.8	26
49	Osteopontin Promoter Polymorphism Is Associated With Increased Carotid Intima-Media Thickness. Journal of the American Society of Echocardiography, 2008, 21, 954-960.	2.8	25
50	Blunted fat oxidation upon submaximal exercise is partially compensated by enhanced glucose metabolism in children, adolescents, and young adults with Barth syndrome. Journal of Inherited Metabolic Disease, 2019, 42, 480-493.	3.6	24
51	Alterations in Left Ventricular Structure and Function in Type-1 Diabetics: A Focus on Left Atrial Contribution to Function. Journal of the American Society of Echocardiography, 2006, 19, 749-755.	2.8	23
52	Doppler Echocardiographic Methods for Optimization of the Atrioventricular Delay during Cardiac Resynchronization Therapy. Echocardiography, 2008, 25, 1047-1055.	0.9	23
53	Relation of Serum Fetuin-A Levels to Coronary Artery Calcium in African-American Patients on Chronic Hemodialysis. American Journal of Cardiology, 2009, 103, 46-49.	1.6	23
54	Mentored Training to Increase Diversity among Faculty in the Biomedical Sciences: The NHLBI Summer Institute Programs to Increase Diversity (SIPID) and the Programs to Increase Diversity among Individuals Engaged in Health-related Research (PRIDE). Ethnicity and Disease, 2017, 27, 249.	2.3	23

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55	Interatrial Conduction Time and Left Atrial Function in Patients With Left Ventricular Systolic Dysfunction: Effects of Cardiac Resynchronization Therapy. Journal of the American Society of Echocardiography, 2009, 22, 472-477.	2.8	21
56	Myocardial glucose and fatty acid metabolism is altered and associated with lower cardiac function in young adults with Barth syndrome. Journal of Nuclear Cardiology, 2021, 28, 1649-1659.	2.1	21
57	Gene-Education Interactions Identify Novel Blood Pressure Loci in the Framingham Heart Study. American Journal of Hypertension, 2014, 27, 431-444.	2.0	19
58	Relation of Left Ventricular Lead Placement in Cardiac Resynchronization Therapy to Left Ventricular Reverse Remodeling and to Diastolic Dyssynchrony. American Journal of Cardiology, 2007, 99, 239-241.	1.6	17
59	Influence of Smoking Status and Intensity on Discovery of Blood Pressure Loci Through Geneâ€Smoking Interactions. Genetic Epidemiology, 2015, 39, 480-488.	1.3	17
60	Gene-educational attainment interactions in a multi-ancestry genome-wide meta-analysis identify novel blood pressure loci. Molecular Psychiatry, 2020, 26, 2111-2125.	7.9	17
61	Abnormalities in Cardiac Structure and Function in Adults with Sickle Cell Disease are not Associated with Pulmonary Hypertension. Journal of the American Society of Echocardiography, 2011, 24, 1285-1290.	2.8	16
62	Central aortic pressure is independently associated with diastolic function. American Heart Journal, 2010, 159, 1081-1088.	2.7	15
63	Low dose chloroquine decreases insulin resistance in human metabolic syndrome but does not reduce carotid intima-media thickness. Diabetology and Metabolic Syndrome, 2019, 11, 61.	2.7	15
64	Variable set enrichment analysis in genome-wide association studies. European Journal of Human Genetics, 2011, 19, 893-900.	2.8	14
65	Whole genome sequence analyses of eGFR in 23,732 people representing multiple ancestries in the NHLBI trans-omics for precision medicine (TOPMed) consortium. EBioMedicine, 2021, 63, 103157.	6.1	14
66	Reduced Muscle Strength in Barth Syndrome May Be Improved by Resistance Exercise Training: A Pilot Study. JIMD Reports, 2018, 41, 63-72.	1.5	13
67	The TDR MOOC training in implementation research: evaluation of feasibility and lessons learned in Rwanda. Pilot and Feasibility Studies, 2020, 6, 66.	1.2	12
68	Left Ventricular Diastolic Filling Prior to Cardiac Resynchronization Therapy: Implications for Atrioventricular Delay Programming. PACE - Pacing and Clinical Electrophysiology, 2008, 31, 838-844.	1.2	11
69	The role of SNP-loop diuretic interactions in hypertension across ethnic groups in HyperGEN. Frontiers in Genetics, 2013, 4, 304.	2.3	11
70	Development of a Health Information Technology Tool for Behavior Change to Address Obesity and Prevent Chronic Disease Among Adolescents: Designing for Dissemination and Sustainment Using the ORBIT Model. Frontiers in Digital Health, 2021, 3, 648777.	2.8	11
71	Obesity–insulin targeted genes in the 3p26-25 region in human studies and LG/J and SM/J mice. Metabolism: Clinical and Experimental, 2012, 61, 1129-1141.	3.4	9
72	Research Education and Mentoring Program in Cardiovascular Diseases for Under-Represented Junior Faculty From NHLBI SIPID/PRIDE. Journal of the American College of Cardiology, 2019, 73, 1861-1865.	2.8	9

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73	Dissemination and Implementation Program in Hypertension in Rwanda: Report on Initial Training and Evaluation. Global Heart, 2019, 14, 135.	2.3	9
74	A risk assessment tool for resumption of research activities during the COVID-19 pandemic for field trials in low resource settings. BMC Medical Research Methodology, 2021, 21, 68.	3.1	8
75	Enhanced detection of genetic association of hypertensive heart disease by analysis of latent phenotypes. Genetic Epidemiology, 2008, 32, 528-538.	1.3	7
76	Molecular Determinants of the Cardiometabolic Phenotype. Endocrine, Metabolic and Immune Disorders - Drug Targets, 2010, 10, 109-123.	1.2	7
77	Are Normative Values for LV Geometry and Mass Based on Fundamental Imaging Valid with Use ofÂHarmonic Imaging?. Journal of the American Society of Echocardiography, 2010, 23, 1317-1322.	2.8	6
78	Genetic association of left ventricular mass assessed by M-mode and two-dimensional echocardiography. Journal of Hypertension, 2016, 34, 88-96.	0.5	6
79	Identifying blood pressure loci whose effects are modulated by multiple lifestyle exposures. Genetic Epidemiology, 2020, 44, 629-641.	1.3	6
80	Association and interaction of PPAR-complex gene variants with latent traits of left ventricular diastolic function. BMC Medical Genetics, 2010, 11, 65.	2.1	5
81	Single Nucleotide Polymorphism–Single Nucleotide Polymorphism Interactions Among Inflammation Genes in the Genetic Architecture of Blood Pressure in the Framingham Heart Study. American Journal of Hypertension, 2015, 28, 248-255.	2.0	5
82	A Novel Method Combining Linkage Disequilibrium Information and Imputed Functional Knowledge for <i>tag</i> SNP Selection. Human Heredity, 2007, 64, 243-249.	0.8	4
83	The St. Louis African American health-heart study: methodology for the study of cardiovascular disease and depression in young-old African Americans. BMC Cardiovascular Disorders, 2013, 13, 66.	1.7	4
84	Genome-wide meta-analysis of variant-by-diuretic interactions as modulators of lipid traits in persons of European and African ancestry. Pharmacogenomics Journal, 2020, 20, 482-493.	2.0	4
85	Ultrasound Core Laboratory for the Household Air Pollution Intervention Network Trial: Standardized Training and Image Management for Field Studies Using Portable Ultrasound in Fetal, Lung, and Vascular Evaluations. Ultrasound in Medicine and Biology, 2021, 47, 1506-1513.	1.5	4
86	Relationships Among HIV Infection, Metabolic Risk Factors, and Left Ventricular Structure and Function. AIDS Research and Human Retroviruses, 2013, 29, 1151-1160.	1.1	3
87	Whole-Exome Sequencing and hiPSC Cardiomyocyte Models Identify MYRIP, TRAPPC11, and SLC27A6 of Potential Importance to Left Ventricular Hypertrophy in an African Ancestry Population. Frontiers in Genetics, 2021, 12, 588452.	2.3	3
88	High rates of undiagnosed and uncontrolled hypertension upon a screening campaign in rural Rwanda: a cross-sectional study. BMC Cardiovascular Disorders, 2022, 22, 197.	1.7	3
89	Adiposity and Cardiometabolic Risk in Children With and Without Antipsychotic Drug Treatment. Journal of Clinical Endocrinology and Metabolism, 2015, 100, 3418-3426.	3.6	2
90	Three Approaches to Modeling Geneâ€Environment Interactions in Longitudinal Family Data: Geneâ€6moking Interactions in Blood Pressure. Genetic Epidemiology, 2016, 40, 73-80.	1.3	2

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91	Diagnostic accuracy of damage-associated molecular patterns (DAMPs) in patients with heart failure with a reduced ejection fraction. Journal of Clinical and Translational Science, 2017, 1, 208-209.	0.6	2
92	The Promise of Selecting Individuals from the Extremes of Exposure in the Analysis of Gene-Physical Activity Interactions. Human Heredity, 2018, 83, 315-332.	0.8	2
93	Multi-ancestry genome-wide association study accounting for gene-psychosocial factor interactions identifies novel loci for blood pressure traits. Human Genetics and Genomics Advances, 2021, 2, 100013.	1.7	2
94	Role of Tissue Doppler and Color M-Mode Imaging for Evaluation of Diastolic Function in Ambulatory Patients with LV Systolic Dysfunction. Echocardiography, 2007, 24, 478-484.	0.9	1
95	Resistance exercise training with protein supplementation improves skeletal muscle strength and improves quality of life in late adolescents and young adults with Barth syndrome: A pilot study. JIMD Reports, 2021, 62, 74-84.	1.5	1
96	Exploring contextual factors influencing the implementation of evidence-based care for hypertension in Rwanda: a cross-sectional study using the COACH questionnaire. BMJ Open, 2021, 11, e048425.	1.9	1
97	Aggregate blood pressure responses to serial dietary sodium and potassium intervention: defining responses using independent component analysis. BMC Genetics, 2015, 16, 64.	2.7	O
98	Lifestyle Risk Score: handling missingness of individual lifestyle components in meta-analysis of gene-by-lifestyle interactions. European Journal of Human Genetics, 2021, 29, 839-850.	2.8	0