Vasiliki Lagou

List of Publications by Year in descending order

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218677 302126 9,556 37 26 39 citations h-index g-index papers 43 43 43 18501 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Large-scale association analysis provides insights into the genetic architecture and pathophysiology of type 2 diabetes. Nature Genetics, 2012, 44, 981-990.	21.4	1,748
2	Genome-wide trans-ancestry meta-analysis provides insight into the genetic architecture of type 2 diabetes susceptibility. Nature Genetics, 2014, 46, 234-244.	21.4	959
3	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. Nature Genetics, 2012, 44, 659-669.	21.4	762
4	Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. Nature Genetics, 2012, 44, 991-1005.	21.4	746
5	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. Nature Genetics, 2011, 43, 1131-1138.	21.4	501
6	Genome-wide associations for birth weight and correlations with adult disease. Nature, 2016, 538, 248-252.	27.8	406
7	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. Nature Genetics, 2019, 51, 804-814.	21.4	402
8	New gene functions in megakaryopoiesis and platelet formation. Nature, 2011, 480, 201-208.	27.8	401
9	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. PLoS Genetics, 2015, 11, e1005378.	3.5	331
10	Seventy-five genetic loci influencing the human red blood cell. Nature, 2012, 492, 369-375.	27.8	320
11	The impact of low-frequency and rare variants on lipid levels. Nature Genetics, 2015, 47, 589-597.	21.4	310
12	Impact of Type 2 Diabetes Susceptibility Variants on Quantitative Glycemic Traits Reveals Mechanistic Heterogeneity. Diabetes, 2014, 63, 2158-2171.	0.6	297
13	New loci associated with birth weight identify genetic links between intrauterine growth and adult height and metabolism. Nature Genetics, 2013, 45, 76-82.	21.4	293
14	The cellular composition of the human immune system is shaped by age and cohabitation. Nature Immunology, 2016, 17, 461-468.	14.5	258
15	Genetic loci influencing kidney function and chronic kidney disease. Nature Genetics, 2010, 42, 373-375.	21.4	246
16	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	12.8	245
17	Familial autoinflammation with neutrophilic dermatosis reveals a regulatory mechanism of pyrin activation. Science Translational Medicine, 2016, 8, 332ra45.	12.4	241
18	Microglia Require CD4ÂT Cells to Complete the Fetal-to-Adult Transition. Cell, 2020, 182, 625-640.e24.	28.9	191

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19	A Central Role for GRB10 in Regulation of Islet Function in Man. PLoS Genetics, 2014, 10, e1004235.	3.5	164
20	A Genome-Wide Association Meta-Analysis of Circulating Sex Hormone–Binding Globulin Reveals Multiple Loci Implicated in Sex Steroid Hormone Regulation. PLoS Genetics, 2012, 8, e1002805.	3.5	151
21	Genetic predisposition for beta cell fragility underlies type 1 and type 2 diabetes. Nature Genetics, 2016, 48, 519-527.	21.4	117
22	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. Nature Communications, 2021, 12, 24.	12.8	87
23	Six Novel Loci Associated with Circulating VEGF Levels Identified by a Meta-analysis of Genome-Wide Association Studies. PLoS Genetics, 2016, 12, e1005874.	3.5	56
24	Beta-Cell Fragility As a Common Underlying Risk Factor in Type 1 and Type 2 Diabetes. Trends in Molecular Medicine, 2017, 23, 181-194.	6.7	53
25	Gene–Lifestyle Interactions in Obesity. Current Nutrition Reports, 2012, 1, 184-196.	4.3	46
26	Machine learning identifies an immunological pattern associated with multiple juvenile idiopathic arthritis subtypes. Annals of the Rheumatic Diseases, 2019, 78, 617-628.	0.9	38
27	Genetic Architecture of Adaptive Immune System Identifies Key Immune Regulators. Cell Reports, 2018, 25, 798-810.e6.	6.4	36
28	Defective Sec $61\hat{l}\pm1$ underlies a novel cause of autosomal dominant severe congenital neutropenia. Journal of Allergy and Clinical Immunology, 2020, 146, 1180-1193.	2.9	32
29	Common Variants Near Melanocortin 4 Receptor Are Associated with General and Visceral Adiposity in European- and African-American Youth. Journal of Pediatrics, 2010, 156, 598-605.e1.	1.8	26
30	NFIL3 mutations alter immune homeostasis and sensitise for arthritis pathology. Annals of the Rheumatic Diseases, 2019, 78, 342-349.	0.9	21
31	Lifestyle and Socioeconomicâ€Status Modify the Effects of <i>ADRB2</i> and <i>NOS3</i> on Adiposity in Europeanâ€American and Africanâ€American Adolescents. Obesity, 2011, 19, 595-603.	3.0	14
32	No Effect of Dietary Aspartame or Stevia on Pancreatic Acinar Carcinoma Development, Growth, or Induced Mortality in a Murine Model. Frontiers in Oncology, 2017, 7, 18.	2.8	7
33	No Functional Role for microRNA-342 in a Mouse Model of Pancreatic Acinar Carcinoma. Frontiers in Oncology, 2017, 7, 101.	2.8	7
34	Heterogeneous Effects of Calorie Content and Nutritional Components Underlie Dietary Influence on Pancreatic Cancer Susceptibility. Cell Reports, 2020, 32, 107880.	6.4	6
35	Role of Gene-Stress Interactions in Gene-Finding Studies. Novartis Foundation Symposium, 0, , 71-86.	1.1	5
36	NOD mice, susceptible to pancreatic autoimmunity, demonstrate delayed growth of pancreatic cancer. Oncotarget, 2017, 8, 80167-80174.	1.8	2

#	Article	IF	CITATIONS
37	Murine Pancreatic Acinar Cell Carcinoma Growth Kinetics Are Independent of Dietary Vitamin D Deficiency or Supplementation. Frontiers in Oncology, 2017, 7, 133.	2.8	1