

Manuela Uda

List of Publications by Year in descending order

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Version: 2024-02-01

109
papers

44,961
citations

8732

75
h-index

24915

109
g-index

113
all docs

113
docs citations

113
times ranked

45852
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	Biological, clinical and population relevance of 95 loci for blood lipids. <i>Nature</i> , 2010, 466, 707-713.	13.7	3,249
3	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010, 42, 937-948.	9.4	2,634
4	New genetic loci implicated in fasting glucose homeostasis and their impact on type 2 diabetes risk. <i>Nature Genetics</i> , 2010, 42, 105-116.	9.4	1,982
5	Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. <i>Nature</i> , 2011, 478, 103-109.	13.7	1,855
6	Hundreds of variants clustered in genomic loci and biological pathways affect human height. <i>Nature</i> , 2010, 467, 832-838.	13.7	1,789
7	Six new loci associated with body mass index highlight a neuronal influence on body weight regulation. <i>Nature Genetics</i> , 2009, 41, 25-34.	9.4	1,572
8	Newly identified loci that influence lipid concentrations and risk of coronary artery disease. <i>Nature Genetics</i> , 2008, 40, 161-169.	9.4	1,488
9	Genome-Wide Association Scan Shows Genetic Variants in the FTO Gene Are Associated with Obesity-Related Traits. <i>PLoS Genetics</i> , 2007, 3, e115.	1.5	1,446
10	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196.	13.7	1,328
11	Common variants at 30 loci contribute to polygenic dyslipidemia. <i>Nature Genetics</i> , 2009, 41, 56-65.	9.4	1,234
12	Common variants near MC4R are associated with fat mass, weight and risk of obesity. <i>Nature Genetics</i> , 2008, 40, 768-775.	9.4	1,179
13	Genome-wide association study identifies eight loci associated with blood pressure. <i>Nature Genetics</i> , 2009, 41, 666-676.	9.4	1,104
14	The putative forkhead transcription factor FOXL2 is mutated in blepharophimosis/ptosis/epicanthus inversus syndrome. <i>Nature Genetics</i> , 2001, 27, 159-166.	9.4	886
15	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. <i>Nature Genetics</i> , 2010, 42, 949-960.	9.4	836
16	Genome-Wide Association Analysis Identifies Variants Associated with Nonalcoholic Fatty Liver Disease That Have Distinct Effects on Metabolic Traits. <i>PLoS Genetics</i> , 2011, 7, e1001324.	1.5	796
17	Interleukin-6 receptor pathways in coronary heart disease: a collaborative meta-analysis of 82 studies. <i>Lancet</i> , The, 2012, 379, 1205-1213.	6.3	668
18	Variants in MTNR1B influence fasting glucose levels. <i>Nature Genetics</i> , 2009, 41, 77-81.	9.4	662

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19	Meta-analysis and imputation refines the association of 15q25 with smoking quantity. <i>Nature Genetics</i> , 2010, 42, 436-440.	9.4	581
20	Meta-Analysis of 28,141 Individuals Identifies Common Variants within Five New Loci That Influence Uric Acid Concentrations. <i>PLoS Genetics</i> , 2009, 5, e1000504.	1.5	572
21	Genome-wide association study shows <i>BCL11A</i> associated with persistent fetal hemoglobin and amelioration of the phenotype of β^2 -thalassemia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 1620-1625.	3.3	561
22	Identification of ten loci associated with height highlights new biological pathways in human growth. <i>Nature Genetics</i> , 2008, 40, 584-591.	9.4	537
23	DNA polymorphisms at the <i>BCL11A</i> , <i>HBS1L-MYB</i> , and β^2 -globin loci associate with fetal hemoglobin levels and pain crises in sickle cell disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 11869-11874.	3.3	510
24	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. <i>Nature Genetics</i> , 2011, 43, 1131-1138.	9.4	501
25	<i>Foxl2</i> disruption causes mouse ovarian failure by pervasive blockage of follicle development. <i>Human Molecular Genetics</i> , 2004, 13, 1171-1181.	1.4	468
26	Meta-Analysis of Genome-Wide Association Studies in >80 000 Subjects Identifies Multiple Loci for C-Reactive Protein Levels. <i>Circulation</i> , 2011, 123, 731-738.	1.6	461
27	Genome-Wide Association Scan Meta-Analysis Identifies Three Loci Influencing Adiposity and Fat Distribution. <i>PLoS Genetics</i> , 2009, 5, e1000508.	1.5	453
28	Thirty new loci for age at menarche identified by a meta-analysis of genome-wide association studies. <i>Nature Genetics</i> , 2010, 42, 1077-1085.	9.4	445
29	Novel Loci for Adiponectin Levels and Their Influence on Type 2 Diabetes and Metabolic Traits: A Multi-Ethnic Meta-Analysis of 45,891 Individuals. <i>PLoS Genetics</i> , 2012, 8, e1002607.	1.5	419
30	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. <i>Nature Communications</i> , 2016, 7, 10023.	5.8	412
31	Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. <i>Nature Genetics</i> , 2011, 43, 1005-1011.	9.4	403
32	New gene functions in megakaryopoiesis and platelet formation. <i>Nature</i> , 2011, 480, 201-208.	13.7	401
33	Genome-wide association study of PR interval. <i>Nature Genetics</i> , 2010, 42, 153-159.	9.4	400
34	Common Variants at 10 Genomic Loci Influence Hemoglobin A1C Levels via Glycemic and Nonglycemic Pathways. <i>Diabetes</i> , 2010, 59, 3229-3239.	0.3	387
35	Common variants in the GDF5-UQCC region are associated with variation in human height. <i>Nature Genetics</i> , 2008, 40, 198-203.	9.4	369
36	Common variants at ten loci modulate the QT interval duration in the QTSCD Study. <i>Nature Genetics</i> , 2009, 41, 407-414.	9.4	356

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37	Novel insights into the genetics of smoking behaviour, lung function, and chronic obstructive pulmonary disease (UK BiLEVE): a genetic association study in UK Biobank. <i>Lancet Respiratory Medicine</i> , 2015, 3, 769-781.	5.2	346
38	Meta-analysis of genome-wide association studies for personality. <i>Molecular Psychiatry</i> , 2012, 17, 337-349.	4.1	340
39	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
40	Seventy-five genetic loci influencing the human red blood cell. <i>Nature</i> , 2012, 492, 369-375.	13.7	320
41	Meta-analyses identify 13 loci associated with age at menopause and highlight DNA repair and immune pathways. <i>Nature Genetics</i> , 2012, 44, 260-268.	9.4	303
42	Foxl2 is required for commitment to ovary differentiation. <i>Human Molecular Genetics</i> , 2005, 14, 2053-2062.	1.4	298
43	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. <i>Nature Genetics</i> , 2014, 46, 826-836.	9.4	281
44	Genome-wide association scan for five major dimensions of personality. <i>Molecular Psychiatry</i> , 2010, 15, 647-656.	4.1	250
45	The GLUT9 Gene Is Associated with Serum Uric Acid Levels in Sardinia and Chianti Cohorts. <i>PLoS Genetics</i> , 2007, 3, e194.	1.5	249
46	Genetic loci influencing kidney function and chronic kidney disease. <i>Nature Genetics</i> , 2010, 42, 373-375.	9.4	246
47	Genome-wide Association Study of Vitamin B6, Vitamin B12, Folate, and Homocysteine Blood Concentrations. <i>American Journal of Human Genetics</i> , 2009, 84, 477-482.	2.6	225
48	High Neuroticism and low Conscientiousness are associated with interleukin-6. <i>Psychological Medicine</i> , 2010, 40, 1485-1493.	2.7	202
49	A Genome-Wide Association Search for Type 2 Diabetes Genes in African Americans. <i>PLoS ONE</i> , 2012, 7, e29202.	1.1	197
50	Novel loci affecting iron homeostasis and their effects in individuals at risk for hemochromatosis. <i>Nature Communications</i> , 2014, 5, 4926.	5.8	192
51	Meta-analysis of genome-wide association studies from the CHARGE consortium identifies common variants associated with carotid intima media thickness and plaque. <i>Nature Genetics</i> , 2011, 43, 940-947.	9.4	191
52	Variants within the immunoregulatory CBLB gene are associated with multiple sclerosis. <i>Nature Genetics</i> , 2010, 42, 495-497.	9.4	164
53	Facets of Personality Linked to Underweight and Overweight. <i>Psychosomatic Medicine</i> , 2009, 71, 682-689.	1.3	157
54	Variations in the G6PC2/ABCB11 genomic region are associated with fasting glucose levels. <i>Journal of Clinical Investigation</i> , 2008, 118, 2620-8.	3.9	146

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55	IRAK-M Is Involved in the Pathogenesis of Early-Onset Persistent Asthma. <i>American Journal of Human Genetics</i> , 2007, 80, 1103-1114.	2.6	144
56	Genome-Wide Association Scan of Trait Depression. <i>Biological Psychiatry</i> , 2010, 68, 811-817.	0.7	143
57	A Genome-Wide Association Scan on the Levels of Markers of Inflammation in Sardinians Reveals Associations That Underpin Its Complex Regulation. <i>PLoS Genetics</i> , 2012, 8, e1002480.	1.5	141
58	Amelioration of Sardinian α^0 thalassemia by genetic modifiers. <i>Blood</i> , 2009, 114, 3935-3937.	0.6	137
59	Fine Mapping of Five Loci Associated with Low-Density Lipoprotein Cholesterol Detects Variants That Double the Explained Heritability. <i>PLoS Genetics</i> , 2011, 7, e1002198.	1.5	134
60	Genome-wide association analysis identifies multiple loci related to resting heart rate. <i>Human Molecular Genetics</i> , 2010, 19, 3885-3894.	1.4	133
61	Common variants in the <i>SLCO1B3</i> locus are associated with bilirubin levels and unconjugated hyperbilirubinemia. <i>Human Molecular Genetics</i> , 2009, 18, 2711-2718.	1.4	126
62	Phosphodiesterase 8B Gene Variants Are Associated with Serum TSH Levels and Thyroid Function. <i>American Journal of Human Genetics</i> , 2008, 82, 1270-1280.	2.6	124
63	<i>Foxl2</i> functions in sex determination and histogenesis throughout mouse ovary development. <i>BMC Developmental Biology</i> , 2009, 9, 36.	2.1	113
64	<i>Gpc3</i> expression correlates with the phenotype of the Simpson-Golabi-Behmel syndrome. <i>Developmental Dynamics</i> , 1998, 213, 431-439.	0.8	104
65	Meta-analysis of genome-wide association studies identifies common variants in <i>CTNNA2</i> associated with excitement-seeking. <i>Translational Psychiatry</i> , 2011, 1, e49-e49.	2.4	97
66	<i>COL4A1</i> Is Associated With Arterial Stiffness by Genome-Wide Association Scan. <i>Circulation: Cardiovascular Genetics</i> , 2009, 2, 151-158.	5.1	91
67	Personality and metabolic syndrome. <i>Age</i> , 2010, 32, 513-519.	3.0	90
68	The central arterial burden of the metabolic syndrome is similar in men and women: the SardiNIA Study. <i>European Heart Journal</i> , 2010, 31, 602-613.	1.0	90
69	The genetic association between personality and major depression or bipolar disorder. A polygenic score analysis using genome-wide association data. <i>Translational Psychiatry</i> , 2011, 1, e50-e50.	2.4	90
70	Common Genetic Variation in the β^2 - <i>BCL11B</i> Gene Desert Is Associated With Carotid-Femoral Pulse Wave Velocity and Excess Cardiovascular Disease Risk. <i>Circulation: Cardiovascular Genetics</i> , 2012, 5, 81-90.	5.1	90
71	<i>BDNF</i> Val66Met is Associated with Introversion and Interacts with 5-HTTLPR to Influence Neuroticism. <i>Neuropsychopharmacology</i> , 2010, 35, 1083-1089.	2.8	89
72	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

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73	Cholesterol, triglycerides, and the Five-Factor Model of personality. <i>Biological Psychology</i> , 2010, 84, 186-191.	1.1	85
74	Associations of large artery structure and function with adiposity: Effects of age, gender, and hypertension. <i>The SardiNIA Study. Atherosclerosis</i> , 2012, 221, 189-197.	0.4	85
75	Genome-wide association study of sexual maturation in males and females highlights a role for body mass and menarche loci in male puberty. <i>Human Molecular Genetics</i> , 2014, 23, 4452-4464.	1.4	82
76	Independent and additive effects of cytokine patterns and the metabolic syndrome on arterial aging in the SardiNIA Study. <i>Atherosclerosis</i> , 2011, 215, 459-464.	0.4	80
77	Ageing of Oocyte, Ovary, and Human Reproduction. <i>Annals of the New York Academy of Sciences</i> , 2004, 1034, 117-131.	1.8	77
78	Crisponi Syndrome Is Caused by Mutations in the CRLF1 Gene and Is Allelic to Cold-Induced Sweating Syndrome Type 1. <i>American Journal of Human Genetics</i> , 2007, 80, 971-981.	2.6	76
79	Effects of Long-Term Averaging of Quantitative Blood Pressure Traits on the Detection of Genetic Associations. <i>American Journal of Human Genetics</i> , 2014, 95, 49-65.	2.6	73
80	Increased Genetic Vulnerability to Smoking at CHRNA5 in Early-Onset Smokers. <i>Archives of General Psychiatry</i> , 2012, 69, 854.	13.8	71
81	Impulsivity is Associated with Uric Acid: Evidence from Humans and Mice. <i>Biological Psychiatry</i> , 2014, 75, 31-37.	0.7	68
82	Sex-Specific Correlates of Walking Speed in a Wide Age-Ranged Population. <i>Journals of Gerontology - Series B Psychological Sciences and Social Sciences</i> , 2010, 65B, 174-184.	2.4	67
83	Neuroticism, Depressive Symptoms, and Serum BDNF. <i>Psychosomatic Medicine</i> , 2011, 73, 638-642.	1.3	67
84	Identification of a common variant in the TFR2 gene implicated in the physiological regulation of serum iron levels. <i>Human Molecular Genetics</i> , 2011, 20, 1232-1240.	1.4	67
85	Personality traits prospectively predict verbal fluency in a lifespan sample.. <i>Psychology and Aging</i> , 2011, 26, 994-999.	1.4	67
86	Genome-wide association analyses based on whole-genome sequencing in Sardinia provide insights into regulation of hemoglobin levels. <i>Nature Genetics</i> , 2015, 47, 1264-1271.	9.4	66
87	Evidence for three genetic loci involved in both anorexia nervosa risk and variation of body mass index. <i>Molecular Psychiatry</i> , 2017, 22, 192-201.	4.1	63
88	Genetics of serum BDNF: Meta-analysis of the Val66Met and genome-wide association study. <i>World Journal of Biological Psychiatry</i> , 2013, 14, 583-589.	1.3	57
89	A common biological basis of obesity and nicotine addiction. <i>Translational Psychiatry</i> , 2013, 3, e308-e308.	2.4	51
90	FOXL2 inactivation by a translocation 171 kb away: analysis of 500 kb of chromosome 3 for candidate long-range regulatory sequences. <i>Genomics</i> , 2004, 83, 757-764.	1.3	50

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91	Determination and stability of sex. <i>BioEssays</i> , 2007, 29, 15-25.	1.2	49
92	Determination and Stability of Gonadal Sex. <i>Journal of Andrology</i> , 2010, 31, 16-25.	2.0	46
93	Jagged-1 mutation analysis in Italian Alagille syndrome patients. , 1999, 14, 394-400.		44
94	Age- and gender-specific awareness, treatment, and control of cardiovascular risk factors and subclinical vascular lesions in a founder population: The SardiNIA Study. <i>Nutrition, Metabolism and Cardiovascular Diseases</i> , 2009, 19, 532-541.	1.1	44
95	Impulsivity-related traits are associated with higher white blood cell counts. <i>Journal of Behavioral Medicine</i> , 2012, 35, 616-623.	1.1	41
96	Variants of the serotonin transporter gene and NEOâ€”Neuroticism: No association in the BLSA and SardiNIA samples. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2009, 150B, 1070-1077.	1.1	36
97	Personality Traits and Leptin. <i>Psychosomatic Medicine</i> , 2013, 75, 505-509.	1.3	31
98	Personality Traits in Sardinia: Testing Founder Population Effects on Trait Means and Variances. <i>Behavior Genetics</i> , 2007, 37, 376-387.	1.4	27
99	FOXL2 modulates cartilage, skeletal development and IGF1-dependent growth in mice. <i>BMC Developmental Biology</i> , 2015, 15, 27.	2.1	27
100	Refining genome-wide linkage intervals using a meta-analysis of genome-wide association studies identifies loci influencing personality dimensions. <i>European Journal of Human Genetics</i> , 2013, 21, 876-882.	1.4	24
101	Trait Antagonism and the Progression of Arterial Thickening. <i>Hypertension</i> , 2010, 56, 617-622.	1.3	20
102	Reversal of Agingâ€”Induced Increases in Aortic Stiffness by Targeting Cytoskeletal Proteinâ€”Protein Interfaces. <i>Journal of the American Heart Association</i> , 2018, 7, .	1.6	17
103	Genome-wide association study of susceptibility loci for breast cancer in Sardinian population. <i>BMC Cancer</i> , 2015, 15, 383.	1.1	12
104	Novel and recurrent JAG1 mutations in patients with tetralogy of Fallot. <i>Clinical Genetics</i> , 2011, 80, 591-594.	1.0	11
105	Transcriptional Control of Ovarian Development in Somatic Cells. <i>Seminars in Reproductive Medicine</i> , 2007, 25, 252-263.	0.5	5
106	Using New Tools to Define the Genetic Underpinnings of Risky Traits Associated With Coronary Artery Disease: The SardiNIA Study. <i>Trends in Cardiovascular Medicine</i> , 2009, 19, 69-75.	2.3	1
107	Genome-wide Association Study of Vitamin B6, Vitamin B12, Folate, and Homocysteine Blood Concentrations. <i>American Journal of Human Genetics</i> , 2009, 84, 712.	2.6	1
108	Jagged-1 MUTATIONS IN ITALIAN PATIENTS WITH ALAGILLE SYNDROME. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 1999, 28, 583.	0.9	0

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109	Editorial: Female Infertility: Genetics of Reproductive Ageing, Menopause and Primary Ovarian Insufficiency. <i>Frontiers in Genetics</i> , 2022, 13, 839758.	1.1	0