Yann C Klimentidis

List of Publications by Year in descending order

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Version: 2024-02-01

81 papers 2,806 citations

279798 23 h-index 214800 47 g-index

90 all docs 90 docs citations

90 times ranked 5296 citing authors

#	Article	IF	CITATIONS
1	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults. JAMA Psychiatry, 2019, 76, 399.	11.0	399
2	Prediction of Complex Human Traits Using the Genomic Best Linear Unbiased Predictor. PLoS Genetics, 2013, 9, e1003608.	3.5	318
3	Genome-wide association study of habitual physical activity in over 377,000 UK Biobank participants identifies multiple variants including CADM2 and APOE. International Journal of Obesity, 2018, 42, 1161-1176.	3.4	249
4	Beyond Missing Heritability: Prediction of Complex Traits. PLoS Genetics, 2011, 7, e1002051.	3.5	224
5	Canaries in the coal mine: a cross-species analysis of the plurality of obesity epidemics. Proceedings of the Royal Society B: Biological Sciences, 2011, 278, 1626-1632.	2.6	123
6	Post-acute sequelae of COVID-19 in a non-hospitalized cohort: Results from the Arizona CoVHORT. PLoS ONE, 2021, 16, e0254347.	2.5	110
7	Natural selection at genomic regions associated with obesity and type-2 diabetes: East Asians and sub-Saharan Africans exhibit high levels of differentiation at type-2 diabetes regions. Human Genetics, 2011, 129, 407-418.	3.8	71
8	Physical exercise is a risk factor for amyotrophic lateral sclerosis: Convergent evidence from Mendelian randomisation, transcriptomics and risk genotypes. EBioMedicine, 2021, 68, 103397.	6.1	65
9	Genetic admixture, selfâ€reported ethnicity, selfâ€estimated admixture, and skin pigmentation among Hispanics and Native Americans. American Journal of Physical Anthropology, 2009, 138, 375-383.	2.1	61
10	Infection with "Escaped―Virus Variants Impairs Control of Simian Immunodeficiency Virus SIVmac239 Replication in <i>Mamu-B*08</i> -Positive Macaques. Journal of Virology, 2009, 83, 11514-11527.	3.4	53
11	Phenotypic and Genetic Characterization of Lower LDL Cholesterol and Increased Type 2 Diabetes Risk in the UK Biobank. Diabetes, 2020, 69, 2194-2205.	0.6	52
12	Associations of air pollution with obesity and body fat percentage, and modification by polygenic risk score for BMI in the UK Biobank. Environmental Research, 2020, 185, 109364.	7. 5	52
13	A Comprehensive Genetic Approach for Improving Prediction of Skin Cancer Risk in Humans. Genetics, 2012, 192, 1493-1502.	2.9	50
14	OPRM1 and EGFR contribute to skin pigmentation differences between Indigenous Americans and Europeans. Human Genetics, 2012, 131, 1073-1080.	3.8	48
15	Heritability of pulmonary function estimated from pedigree and whole-genome markers. Frontiers in Genetics, 2013, 4, 174.	2.3	44
16	A Conserved Role for Syndecan Family Members in the Regulation of Whole-Body Energy Metabolism. PLoS ONE, 2010, 5, e11286.	2.5	41
17	T-Cell Correlates of Vaccine Efficacy after a Heterologous Simian Immunodeficiency Virus Challenge. Journal of Virology, 2010, 84, 4352-4365.	3.4	40
18	Ordered multinomial regression for genetic association analysis of ordinal phenotypes at Biobank scale. Genetic Epidemiology, 2020, 44, 248-260.	1.3	37

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19	Genetic influences in childhood obesity: recent progress and recommendations for experimental designs. International Journal of Obesity, 2012, 36, 479-484.	3.4	35
20	Differential associations of engagement in physical activity and estimated cardiorespiratory fitness with brain volume in middle-aged to older adults. Brain Imaging and Behavior, 2020, 14, 1994-2003.	2.1	33
21	Association of physical activity with lower type 2 diabetes incidence is weaker among individuals at high genetic risk. Diabetologia, 2014, 57, 2530-2534.	6.3	26
22	CYP24A1 and CYP27B1 Polymorphisms, Concentrations of Vitamin D Metabolites, and Odds of Colorectal Adenoma Recurrence. Nutrition and Cancer, 2015, 67, 1131-1141.	2.0	26
23	Body Mass Index, Waist Circumference, and Mortality in a Large Multiethnic Postmenopausal Cohort—Results from the Women's Health Initiative. Journal of the American Geriatrics Society, 2017, 65, 1907-1915.	2.6	26
24	Genetic admixture, social–behavioural factors and body composition are associated with blood pressure differently by racial–ethnic group among children. Journal of Human Hypertension, 2012, 26, 98-107.	2.2	25
25	A unified GMDR method for detecting gene–gene interactions in family and unrelated samples with application to nicotine dependence. Human Genetics, 2014, 133, 139-150.	3.8	23
26	Diminished Milk Synthesis in Upstream Stimulatory Factor 2 Null Mice Is Associated With Decreased Circulating Oxytocin and Decreased Mammary Gland Expression of Eukaryotic Initiation Factors 4E and 4G. Molecular Endocrinology, 2003, 17, 2251-2267.	3.7	21
27	Triglyceride-Increasing Alleles Associated with Protection against Type-2 Diabetes. PLoS Genetics, 2015, 11, e1005204.	3. 5	21
28	Indigenous American Ancestry is Associated with Arsenic Methylation Efficiency in an Admixed Population of Northwest Mexico. Journal of Toxicology and Environmental Health - Part A: Current Issues, 2012, 75, 36-49.	2.3	20
29	Multiple Metabolic Genetic Risk Scores and Type 2 Diabetes Risk in Three Racial/Ethnic Groups. Journal of Clinical Endocrinology and Metabolism, 2014, 99, E1814-E1818.	3.6	20
30	Ancestry-informative markers on chromosomes 2, 8 and 15 are associated with insulin-related traits in a racially diverse sample of children. Human Genomics, 2011, 5, 79.	2.9	19
31	Obesity polymorphisms identified in genome-wide association studies interact with n-3 polyunsaturated fatty acid intake and modify the genetic association with adiposity phenotypes in Yup'ik people. Genes and Nutrition, 2013, 8, 495-505.	2.5	19
32	The relationship between European genetic admixture and body composition among Hispanics and Native Americans. American Journal of Human Biology, 2009, 21, 377-382.	1.6	18
33	High genetic risk individuals benefit less from resistance exercise intervention. International Journal of Obesity, 2015, 39, 1371-1375.	3.4	18
34	Fractal Complexity of Daily Physical Activity Patterns Differs With Age Over the Life Span and Is Associated With Mortality in Older Adults. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2019, 74, 1461-1467.	3.6	18
35	Prediction of Expected Years of Life Using Whole-Genome Markers. PLoS ONE, 2012, 7, e40964.	2.5	17
36	FTO association and interaction with time spent sitting. International Journal of Obesity, 2016, 40, 411-416.	3 . 4	17

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37	Genetically predicted physical activity levels are associated with lower colorectal cancer risk: a Mendelian randomisation study. British Journal of Cancer, 2021, 124, 1330-1338.	6.4	17
38	<i><scp>CDKAL1</scp></i> and <i><scp>HHEX</scp></i> are associated with type 2 diabetesâ€related traits among <scp>Y</scp> up'ik people (在å°Çš®å⟨人羼¸ <i>CDKAL1</i> å'Œ <i>HHEX</i> 与²åž⟨ç³−å°¿ç−251-259.	ç> jå8. ³ç%	‰¹å¥#有å.
39	Racial, ethnic, and gender differences in obesity and body fat distribution: An All of Us Research Program demonstration project. PLoS ONE, 2021, 16, e0255583.	2.5	16
40	Associations of Obesity Genes with Obesity-related Outcomes in Multiethnic Children. Archives of Medical Research, 2011, 42, 509-514.	3.3	15
41	Interaction of Age and Self-reported Physical Sports Activity on White Matter Hyperintensity Volume in Healthy Older Adults. Frontiers in Aging Neuroscience, 2020, 12, 576025.	3.4	15
42	Design of the Arizona CoVHORT: A Population-Based COVID-19 Cohort. Frontiers in Public Health, 2021, 9, 620060.	2.7	15
43	The E3 ubiquitin ligase MARCH1 regulates glucose-tolerance and lipid storage in a sex-specific manner. PLoS ONE, 2018, 13, e0204898.	2.5	14
44	Bed net care practices and associated factors in western Kenya. Malaria Journal, 2019, 18, 274.	2.3	14
45	Estimating Genetic Ancestry Proportions from Faces. PLoS ONE, 2009, 4, e4460.	2.5	12
46	High Viremia Is Associated with High Levels of <i>In Vivo</i> Major Histocompatibility Complex Class I Downregulation in Rhesus Macaques Infected with Simian Immunodeficiency Virus SIVmac239. Journal of Virology, 2010, 84, 5443-5447.	3.4	12
47	The Genetic Contribution of West-African Ancestry to Protection against Central Obesity in African-American Men but Not Women: Results from the ARIC and MESA Studies. Frontiers in Genetics, 2016, 7, 89.	2.3	11
48	Nonalcoholic Fatty Liver Disease and Associated Risk Factors in a Communityâ€Based Sample of Mexicanâ€Origin Adults. Hepatology Communications, 2022, 6, 1322-1335.	4.3	11
49	Associations of the lactase persistence allele and lactose intake with body composition among multiethnic children. Genes and Nutrition, 2013, 8, 487-494.	2.5	10
50	Evidence for novel genetic loci associated with metabolic traits in Yup'ik people. American Journal of Human Biology, 2013, 25, 673-680.	1.6	10
51	<i>Taq1a</i> polymorphism (rs1800497) is associated with obesityâ€related outcomes and dietary intake in a multiâ€ethnic sample of children. Pediatric Obesity, 2019, 14, e12470.	2.8	10
52	Association of Air Pollution and Physical Activity With Brain Volumes. Neurology, 2022, 98, e416-e426.	1.1	10
53	Genetic Variant in ACVR2B Is Associated with Lean Mass. Medicine and Science in Sports and Exercise, 2016, 48, 1270-1275.	0.4	9
54	Psychosocial stress and bone loss among postmenopausal women: results from the Women's Health Initiative. Journal of Epidemiology and Community Health, 2019, 73, 888-892.	3.7	9

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55	Genome-wide association study of body fat distribution traits in Hispanics/Latinos from the HCHS/SOL. Human Molecular Genetics, 2021, 30, 2190-2204.	2.9	8
56	Inability of Overexpressed des(1-3)Human Insulin-Like Growth Factor I (IGF-I) to Inhibit Forced Mammary Gland Involution Is Associated with Decreased Expression of IGF Signaling Molecules. Endocrinology, 2001, 142, 1479-1488.	2.8	8
57	Interaction of Insulin Resistance and Related Genetic Variants With Triglyceride-Associated Genetic Variants. Circulation: Cardiovascular Genetics, 2016, 9, 154-161.	5.1	7
58	Associations between ACE-Inhibitors, Angiotensin Receptor Blockers, and Lean Body Mass in Community Dwelling Older Women. Journal of Aging Research, 2018, 2018, 1-8.	0.9	7
59	Identification of Allelic Heterogeneity at Type-2 Diabetes Loci and Impact on Prediction. PLoS ONE, 2014, 9, e113072.	2.5	6
60	"After those nets are torn, most people use them for other purposes†an examination of alternative bed net use in western Kenya. Malaria Journal, 2020, 19, 272.	2.3	6
61	Circulating Fibroblast Growth Factor-21 and Risk of Metachronous Colorectal Adenoma. Journal of Gastrointestinal Cancer, 2021, 52, 940-946.	1.3	6
62	Malaria education interventions addressing bed net care and repair practices: a systematic review. Pathogens and Global Health, 2020, 114, 2-15.	2.3	6
63	Weight Loss Interventions for Hispanic Women in the United States: A Systematic Review. Journal of Environmental and Public Health, 2021, 2021, 1-14.	0.9	6
64	Geographic Variation in Obesity at the State Level in the All of Us Research Program. Preventing Chronic Disease, 2021, 18, E104.	3.4	6
65	Natural selection among Eurasians at genomic regions associated with HIV-1 control. BMC Evolutionary Biology, 2011, 11, 173.	3.2	5
66	A hybrid Bayesian Network/Structural Equation Modeling (BN/SEM) approach for detecting physiological networks for obesity-related genetic variants., 2011,, 696-702.		5
67	Assessment of Whole-Genome Regression for Type II Diabetes. PLoS ONE, 2015, 10, e0123818.	2.5	5
68	Discovery of phenotypic networks from genotypic association studies with application to obesity. International Journal of Data Mining and Bioinformatics, 2015, 12, 129.	0.1	4
69	Serum urate gene associations with incident gout, measured in the Framingham Heart Study, are modified by renal disease and not by body mass index. Rheumatology International, 2016, 36, 263-270.	3.0	4
70	Weight loss interventions for Hispanic women in the USA: a protocol for a systematic review. Systematic Reviews, 2019, 8, 301.	5.3	4
71	Assessing Interactions between PNPLA3 and Dietary Intake on Liver Steatosis in Mexican-Origin Adults. International Journal of Environmental Research and Public Health, 2021, 18, 7055.	2.6	4
72	Association of <i>PNPLA3</i> I148M with Liver Disease Biomarkers in Latinos. Human Heredity, 2021, 86, 21-27.	0.8	4

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73	Identification of genetic loci simultaneously associated with multiple cardiometabolic traits. Nutrition, Metabolism and Cardiovascular Diseases, 2022, 32, 1027-1034.	2.6	4
74	Polymorphisms in stearoyl coa desaturase and sterol regulatory element binding protein interact with N-3 polyunsaturated fatty acid intake to modify associations with anthropometric variables and metabolic phenotypes in Yup'ik people. Molecular Nutrition and Food Research, 2016, 60, 2642-2653.	3.3	3
75	Chronic kidney disease unawareness and determinants using 1999–2014 National Health and Nutrition Examination Survey Data. Journal of Public Health, 2021, , .	1.8	3
76	Elucidating symptoms of COVID-19 illness in the Arizona CoVHORT: a longitudinal cohort study. BMJ Open, 2022, 12, e053403.	1.9	3
77	Genome-wide Association Study of Liking for Several Types of Physical Activity in the UK Biobank and Two Replication Cohorts. Medicine and Science in Sports and Exercise, 2022, 54, 1252-1260.	0.4	3
78	The intersectional role of social stress in fracture risk: results from the Women's Health Initiative. Journal of Epidemiology and Community Health, 2021, 75, 1208-1214.	3.7	2
79	Integrated genomic and BMI analysis for type 2 diabetes risk assessment. Frontiers in Genetics, 2015, 6, 75.	2.3	1
80	In response to: †Information bias in measures of self-reported physical activity'. International Journal of Obesity, 2018, 42, 2064-2065.	3.4	0
81	Obesity polymorphisms identified in genomeâ€wide association studies interact with nâ€3 polyunsaturated fatty acid intake and modify genetic associations with adiposity phenotypes in Yup'ik people. FASEB Journal, 2013, 27, 608.2.	0.5	0