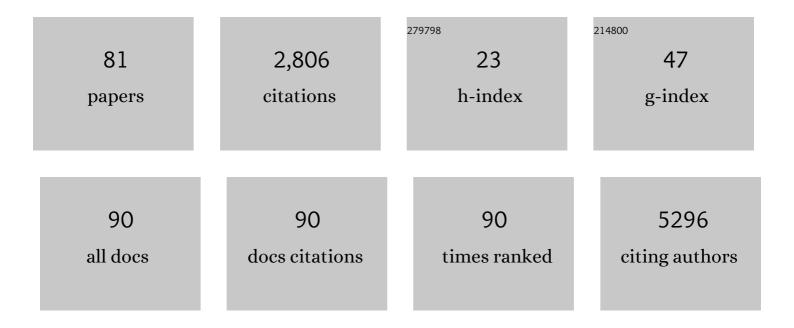
## Yann C Klimentidis

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	ldentification of genetic loci simultaneously associated with multiple cardiometabolic traits. Nutrition, Metabolism and Cardiovascular Diseases, 2022, 32, 1027-1034.	2.6	4
2	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
3	Elucidating symptoms of COVID-19 illness in the Arizona CoVHORT: a longitudinal cohort study. BMJ Open, 2022, 12, e053403.	1.9	3
4	Association of Air Pollution and Physical Activity With Brain Volumes. Neurology, 2022, 98, e416-e426.	1.1	10
5	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
6	Circulating Fibroblast Growth Factor-21 and Risk of Metachronous Colorectal Adenoma. Journal of Gastrointestinal Cancer, 2021, 52, 940-946.	1.3	6
7	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
8	Design of the Arizona CoVHORT: A Population-Based COVID-19 Cohort. Frontiers in Public Health, 2021, 9, 620060.	2.7	15
9	Chronic kidney disease unawareness and determinants using 1999–2014 National Health and Nutrition Examination Survey Data. Journal of Public Health, 2021, , .	1.8	3
10	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
11	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
12	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
13	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
14	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
15	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
16	Post-acute sequelae of COVID-19 in a non-hospitalized cohort: Results from the Arizona CoVHORT. PLoS ONE, 2021, 16, e0254347.	2.5	110
17	Association of <b><i>PNPLA3</i></b> I148M with Liver Disease Biomarkers in Latinos. Human Heredity, 2021, 86, 21-27.	0.8	4
18	Geographic Variation in Obesity at the State Level in the All of Us Research Program. Preventing Chronic Disease, 2021, 18, E104.	3.4	6

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19	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
20	Ordered multinomial regression for genetic association analysis of ordinal phenotypes at Biobank scale. Genetic Epidemiology, 2020, 44, 248-260.	1.3	37
21	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
22	"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
23	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
24	Malaria education interventions addressing bed net care and repair practices: a systematic review. Pathogens and Global Health, 2020, 114, 2-15.	2.3	6
25	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
26	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
27	Bed net care practices and associated factors in western Kenya. Malaria Journal, 2019, 18, 274.	2.3	14
28	Assessment of Bidirectional Relationships Between Physical Activity and Depression Among Adults. JAMA Psychiatry, 2019, 76, 399.	11.0	399
29	Weight loss interventions for Hispanic women in the USA: a protocol for a systematic review. Systematic Reviews, 2019, 8, 301.	5.3	4
30	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
31	<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
32	The E3 ubiquitin ligase MARCH1 regulates glucose-tolerance and lipid storage in a sex-specific manner. PLoS ONE, 2018, 13, e0204898.	2.5	14
33	In response to: †Information bias in measures of self-reported physical activity'. International Journal of Obesity, 2018, 42, 2064-2065.	3.4	Ο
34	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
35	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
36	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

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37	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
38	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
39	Genetic Variant in ACVR2B Is Associated with Lean Mass. Medicine and Science in Sports and Exercise, 2016, 48, 1270-1275.	0.4	9
40	Interaction of Insulin Resistance and Related Genetic Variants With Triglyceride-Associated Genetic Variants. Circulation: Cardiovascular Genetics, 2016, 9, 154-161.	5.1	7
41	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
42	FTO association and interaction with time spent sitting. International Journal of Obesity, 2016, 40, 411-416.	3.4	17
43	Assessment of Whole-Genome Regression for Type II Diabetes. PLoS ONE, 2015, 10, e0123818.	2.5	5
44	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
45	Triglyceride-Increasing Alleles Associated with Protection against Type-2 Diabetes. PLoS Genetics, 2015, 11, e1005204.	3.5	21
46	Integrated genomic and BMI analysis for type 2 diabetes risk assessment. Frontiers in Genetics, 2015, 6, 75.	2.3	1
47	High genetic risk individuals benefit less from resistance exercise intervention. International Journal of Obesity, 2015, 39, 1371-1375.	3.4	18
48	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
49	Identification of Allelic Heterogeneity at Type-2 Diabetes Loci and Impact on Prediction. PLoS ONE, 2014, 9, e113072.	2.5	6
50	<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°ਝूš®å‹ä≌ç¾ष्ठ, <i>CDKAL1</i> å'Œ <i>HHEX</i> ä,Ž2型糖尿ç- 251-259.	ç> <b>jå8.</b> ³ç%	‰¹å¥4œœ‰å
51	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
52	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
53	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
54	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

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55	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
56	Prediction of Complex Human Traits Using the Genomic Best Linear Unbiased Predictor. PLoS Genetics, 2013, 9, e1003608.	3.5	318
57	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
58	Heritability of pulmonary function estimated from pedigree and whole-genome markers. Frontiers in Genetics, 2013, 4, 174.	2.3	44
59	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
60	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
61	A Comprehensive Genetic Approach for Improving Prediction of Skin Cancer Risk in Humans. Genetics, 2012, 192, 1493-1502.	2.9	50
62	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
63	Genetic influences in childhood obesity: recent progress and recommendations for experimental designs. International Journal of Obesity, 2012, 36, 479-484.	3.4	35
64	Prediction of Expected Years of Life Using Whole-Genome Markers. PLoS ONE, 2012, 7, e40964.	2.5	17
65	OPRM1 and EGFR contribute to skin pigmentation differences between Indigenous Americans and Europeans. Human Genetics, 2012, 131, 1073-1080.	3.8	48
66	Natural selection among Eurasians at genomic regions associated with HIV-1 control. BMC Evolutionary Biology, 2011, 11, 173.	3.2	5
67	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
68	Associations of Obesity Genes with Obesity-related Outcomes in Multiethnic Children. Archives of Medical Research, 2011, 42, 509-514.	3.3	15
69	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
70	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
71	A hybrid Bayesian Network/Structural Equation Modeling (BN/SEM) approach for detecting physiological networks for obesity-related genetic variants. , 2011, , 696-702.		5
72	Beyond Missing Heritability: Prediction of Complex Traits. PLoS Genetics, 2011, 7, e1002051.	3.5	224

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73	A Conserved Role for Syndecan Family Members in the Regulation of Whole-Body Energy Metabolism. PLoS ONE, 2010, 5, e11286.	2.5	41
74	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
75	T-Cell Correlates of Vaccine Efficacy after a Heterologous Simian Immunodeficiency Virus Challenge. Journal of Virology, 2010, 84, 4352-4365.	3.4	40
76	Estimating Genetic Ancestry Proportions from Faces. PLoS ONE, 2009, 4, e4460.	2.5	12
77	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
78	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
79	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
80	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
81	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