Kathleen A Ryan

List of Publications by Year in descending order

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279798 182427 3,234 55 23 51 citations h-index g-index papers 61 61 61 6745 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A catalog of genetic loci associated with kidney function from analyses of a million individuals. Nature Genetics, 2019, 51, 957-972.	21.4	549
2	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	21.4	341
3	Genome Analyses of >200,000 Individuals Identify 58 Loci for Chronic Inflammation and Highlight Pathways that Link Inflammation and Complex Disorders. American Journal of Human Genetics, 2018, 103, 691-706.	6.2	326
4	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. Nature Genetics, 2016, 48, 1462-1472.	21.4	284
5	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. Nature Genetics, 2019, 51, 1459-1474.	21.4	251
6	Obesity Increases Risk of Ischemic Stroke in Young Adults. Stroke, 2015, 46, 1690-1692.	2.0	159
7	Genome-wide association meta-analyses and fine-mapping elucidate pathways influencing albuminuria. Nature Communications, 2019, 10, 4130.	12.8	133
8	The genetic response to short-term interventions affecting cardiovascular function: Rationale and design of the Heredity and Phenotype Intervention (HAPI) Heart Study. American Heart Journal, 2008, 155, 823-828.	2.7	109
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	Familial Defective Apolipoprotein B-100 and Increased Low-Density Lipoprotein Cholesterol and Coronary Artery Calcification in the Old Order Amish. Archives of Internal Medicine, 2010, 170, 1850-5.	3.8	79
11	Deep coverage whole genome sequences and plasma lipoprotein(a) in individuals of European and African ancestries. Nature Communications, 2018, 9, 2606.	12.8	79
12	A multiancestry genome-wide association study of unexplained chronic ALT elevation as a proxy for nonalcoholic fatty liver disease with histological and radiological validation. Nature Genetics, 2022, 54, 761-771.	21.4	68
13	Cocaine Use and Risk of Ischemic Stroke in Young Adults. Stroke, 2016, 47, 918-922.	2.0	64
14	Multi-ancestry GWAS of the electrocardiographic PR interval identifies 202 loci underlying cardiac conduction. Nature Communications, 2020, 11, 2542.	12.8	59
15	Genetic Variants Associated with Circulating Parathyroid Hormone. Journal of the American Society of Nephrology: JASN, 2017, 28, 1553-1565.	6.1	52
16	Genome-wide association study of knee pain identifies associations with GDF5 and COL27A1 in UK Biobank. Communications Biology, 2019, 2, 321.	4.4	48
17	A Common Variant in the SETD7 Gene Predicts Serum Lycopene Concentrations. Nutrients, 2016, 8, 82.	4.1	45
18	Insulin Resistance Exacerbates Genetic Predisposition to Nonalcoholic Fatty Liver Disease in Individuals Without Diabetes. Hepatology Communications, 2019, 3, 894-907.	4.3	41

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19	The <i>ABCG8</i> G574R Variant, Serum Plant Sterol Levels, and Cardiovascular Disease Risk in the Old Order Amish. Arteriosclerosis, Thrombosis, and Vascular Biology, 2013, 33, 413-419.	2.4	33
20	Genome sequencing unveils a regulatory landscape of platelet reactivity. Nature Communications, 2021, 12, 3626.	12.8	29
21	Chronotype and seasonality: Morningness is associated with lower seasonal mood and behavior changes in the Old Order Amish. Journal of Affective Disorders, 2015, 174, 209-214.	4.1	28
22	Familial Hypercholesterolemia and Type 2 Diabetes in the Old Order Amish. Diabetes, 2017, 66, 2054-2058.	0.6	28
23	Whole-genome sequencing association analysis of quantitative red blood cell phenotypes: The NHLBI TOPMed program. American Journal of Human Genetics, 2021, 108, 874-893.	6.2	28
24	Genetic Variation in the Platelet Endothelial Aggregation Receptor 1 Gene Results in Endothelial Dysfunction. PLoS ONE, 2015, 10, e0138795.	2.5	24
25	Insulin Pump and Continuous Glucose Monitor Initiation in Hospitalized Patients with Type 2 Diabetes Mellitus. Diabetes Technology and Therapeutics, 2018, 20, 32-38.	4.4	22
26	Differential and shared genetic effects on kidney function between diabetic and non-diabetic individuals. Communications Biology, 2022, 5, .	4.4	17
27	Whole-genome sequencing in diverse subjects identifies genetic correlates of leukocyte traits: The NHLBI TOPMed program. American Journal of Human Genetics, 2021, 108, 1836-1851.	6.2	14
28	Genetic and functional evidence links a missense variant in <i>B4GALT1</i> to lower LDL and fibrinogen. Science, 2021, 374, 1221-1227.	12.6	14
29	Genetic variants affecting bone mineral density and bone mineral content at multiple skeletal sites in Hispanic children. Bone, 2020, 132, 115175.	2.9	13
30	Marijuana Use and the Risk of Early Ischemic Stroke. Stroke, 2021, 52, 3184-3190.	2.0	13
31	Prevalence, control, and treatment of diabetes, hypertension, and high cholesterol in the Amish. BMJ Open Diabetes Research and Care, 2020, 8, e000912.	2.8	12
32	Prevention Opportunities for Oral Contraceptive–Associated Ischemic Stroke. Stroke, 2014, 45, 893-895.	2.0	11
33	Ethnic differences in ischemic stroke subtypes in young-onset stroke: the Stroke Prevention in Young Adults Study. BMC Neurology, 2015, 15, 221.	1.8	11
34	Heterozygosity for a Pathogenic Variant in SLC12A3 That Causes Autosomal Recessive Gitelman Syndrome Is Associated with Lower Serum Potassium. Journal of the American Society of Nephrology: JASN, 2021, 32, 756-765.	6.1	11
35	An exome-wide sequencing study of lipid response to high-fat meal and fenofibrate in Caucasians from the GOLDN cohort. Journal of Lipid Research, 2018, 59, 722-729.	4.2	10
36	An <i>APOO</i> Pseudogene on Chromosome 5q Is Associated With Low-Density Lipoprotein Cholesterol Levels. Circulation, 2018, 138, 1343-1355.	1.6	10

#	Article	IF	Citations
37	Increased usual physical activity is associated with a blunting of the triglyceride response to a high-fat meal. Journal of Clinical Lipidology, 2019, 13, 109-114.	1.5	9
38	Genomeâ€wide metaâ€analysis identified novel variant associated with hallux valgus in Caucasians. Journal of Foot and Ankle Research, 2020, 13, 11.	1.9	9
39	Allelic Heterogeneity at the CRP Locus Identified by Whole-Genome Sequencing in Multi-ancestry Cohorts. American Journal of Human Genetics, 2020, 106, 112-120.	6.2	9
40	Genetics of the thrombomodulin-endothelial cell protein C receptor system and the risk of early-onset ischemic stroke. PLoS ONE, 2018, 13, e0206554.	2.5	8
41	Toxoplasma gondii Serointensity and Seropositivity: Heritability and Household-Related Associations in the Old Order Amish. International Journal of Environmental Research and Public Health, 2019, 16, 3732.	2.6	8
42	<i>KCNQ1</i> and Long QT Syndrome in 1/45 Amish. Circulation Genomic and Precision Medicine, 2020, 13, e003133.	3.6	7
43	An Amish founder population reveals rare-population genetic determinants of the human lipidome. Communications Biology, 2022, 5, 334.	4.4	7
44	Clopidogrel Improves Skin Microcirculatory Endothelial Function in Persons With Heightened Platelet Aggregation. Journal of the American Heart Association, 2016, 5, .	3.7	6
45	Toxoplasma gondii IgG associations with sleepwake problems, sleep duration and timing. Pteridines, 2019, 30, 1-9.	0.5	6
46	Self-Reported Sleep Duration and Pattern in Old Order Amish and Non-Amish Adults. Journal of Clinical Sleep Medicine, 2019, 15, 1321-1328.	2.6	6
47	Heritability of plasma neopterin levels in the Old Order Amish. Journal of Neuroimmunology, 2017, 307, 37-41.	2.3	5
48	Exome Array Analysis of Early-Onset Ischemic Stroke. Stroke, 2020, 51, 3356-3360.	2.0	5
49	Black-White Differences in Ischemic Stroke Risk Factor Burden in Young Adults. Stroke, 2022, 53, STROKEAHA121034314.	2.0	5
50	Identification of novel and rare variants associated with handgrip strength using whole genome sequence data from the NHLBI Trans-Omics in Precision Medicine (TOPMed) Program. PLoS ONE, 2021, 16, e0253611.	2.5	4
51	Evaluation of WISP1 as a candidate gene for bone mineral density in the Old Order Amish. Scientific Reports, 2018, 8, 7141.	3.3	3
52	An Exome-Wide Sequencing Study of the GOLDN Cohort Reveals Novel Associations of Coding Variants and Fasting Plasma Lipids. Frontiers in Genetics, 2019, 10, 158.	2.3	2
53	Genomics of Postprandial Lipidomics in the Genetics of Lipid-Lowering Drugs and Diet Network Study. Nutrients, 2021, 13, 4000.	4.1	2
54	Sickle Cell Trait and Risk of Ischemic Stroke in Young Adults. Stroke, 2020, 51, e238-e241.	2.0	1

#	Article	IF	CITATIONS
55	Eating behavior disinhibition predicts insulin resistance in the Old Order Amish. Journal of Human Nutrition and Food Science, 2017, 5, .	0.0	0