

# Kevin B Jacobs

## List of Publications by Year in descending order

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Version: 2024-02-01

77  
papers

28,059  
citations

28274

55  
h-index

66911

78  
g-index

78  
all docs

78  
docs citations

78  
times ranked

34106  
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.	27.8	3,823
2	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010, 42, 937-948.	21.4	2,634
3	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	21.4	1,818
4	Hundreds of variants clustered in genomic loci and biological pathways affect human height. <i>Nature</i> , 2010, 467, 832-838.	27.8	1,789
5	Six new loci associated with body mass index highlight a neuronal influence on body weight regulation. <i>Nature Genetics</i> , 2009, 41, 25-34.	21.4	1,572
6	A genome-wide association study identifies alleles in <i>FGFR2</i> associated with risk of sporadic postmenopausal breast cancer. <i>Nature Genetics</i> , 2007, 39, 870-874.	21.4	1,370
7	Common variants near <i>MC4R</i> are associated with fat mass, weight and risk of obesity. <i>Nature Genetics</i> , 2008, 40, 768-775.	21.4	1,179
8	Genome-wide association study of prostate cancer identifies a second risk locus at 8q24. <i>Nature Genetics</i> , 2007, 39, 645-649.	21.4	1,059
9	Multiple loci identified in a genome-wide association study of prostate cancer. <i>Nature Genetics</i> , 2008, 40, 310-315.	21.4	871
10	Genome-wide association study of circulating vitamin D levels. <i>Human Molecular Genetics</i> , 2010, 19, 2739-2745.	2.9	700
11	Estimation of effect size distribution from genome-wide association studies and implications for future discoveries. <i>Nature Genetics</i> , 2010, 42, 570-575.	21.4	609
12	Genome-wide association study identifies variants in the <i>ABO</i> locus associated with susceptibility to pancreatic cancer. <i>Nature Genetics</i> , 2009, 41, 986-990.	21.4	597
13	Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. <i>Nature Genetics</i> , 2013, 45, 501-512.	21.4	578
14	A genome-wide association study identifies pancreatic cancer susceptibility loci on chromosomes 13q22.1, 1q32.1 and 5p15.33. <i>Nature Genetics</i> , 2010, 42, 224-228.	21.4	539
15	Identification of ten loci associated with height highlights new biological pathways in human growth. <i>Nature Genetics</i> , 2008, 40, 584-591.	21.4	537
16	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012, 44, 651-658.	21.4	519
17	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010, 42, 978-984.	21.4	493
18	A multistage genome-wide association study in breast cancer identifies two new risk alleles at 1p11.2 and 14q24.1 ( <i>RAD51L1</i> ). <i>Nature Genetics</i> , 2009, 41, 579-584.	21.4	487

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19	A shared susceptibility locus in PLCE1 at 10q23 for gastric adenocarcinoma and esophageal squamous cell carcinoma. <i>Nature Genetics</i> , 2010, 42, 764-767.	21.4	453
20	FTO genotype is associated with phenotypic variability of body mass index. <i>Nature</i> , 2012, 490, 267-272.	27.8	383
21	Performance of Common Genetic Variants in Breast-Cancer Risk Models. <i>New England Journal of Medicine</i> , 2010, 362, 986-993.	27.0	376
22	Sex-stratified Genome-wide Association Studies Including 270,000 Individuals Show Sexual Dimorphism in Genetic Loci for Anthropometric Traits. <i>PLoS Genetics</i> , 2013, 9, e1003500.	3.5	371
23	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015, 11, e1005378.	3.5	331
24	Haseman and Elston revisited. <i>Genetic Epidemiology</i> , 2000, 19, 1-17.	1.3	324
25	Anthropometric Measures, Body Mass Index, and Pancreatic Cancer. <i>Archives of Internal Medicine</i> , 2010, 170, 791.	3.8	314
26	Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , 2012, 44, 1330-1335.	21.4	286
27	A Subset-Based Approach Improves Power and Interpretation for the Combined Analysis of Genetic Association Studies of Heterogeneous Traits. <i>American Journal of Human Genetics</i> , 2012, 90, 821-835.	6.2	242
28	Novel Breast Cancer Susceptibility Locus at 9q31.2: Results of a Genome-Wide Association Study. <i>Journal of the National Cancer Institute</i> , 2011, 103, 425-435.	6.3	225
29	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , 2011, 43, 60-65.	21.4	220
30	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , 2009, 41, 1055-1057.	21.4	218
31	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013, 45, 868-876.	21.4	179
32	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , 2012, 21, 5373-5384.	2.9	168
33	Genome-wide association study identifies new prostate cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2011, 20, 3867-3875.	2.9	160
34	Meta-analysis identifies four new loci associated with testicular germ cell tumor. <i>Nature Genetics</i> , 2013, 45, 680-685.	21.4	154
35	Mosaic loss of chromosome Y is associated with common variation near TCL1A. <i>Nature Genetics</i> , 2016, 48, 563-568.	21.4	134
36	Family history of cancer and risk of pancreatic cancer: A pooled analysis from the Pancreatic Cancer Cohort Consortium (PanScan). <i>International Journal of Cancer</i> , 2010, 127, 1421-1428.	5.1	128

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37	Genome-wide analysis of BMI in adolescents and young adults reveals additional insight into the effects of genetic loci over the life course. <i>Human Molecular Genetics</i> , 2013, 22, 3597-3607.	2.9	116
38	Diabetes and risk of pancreatic cancer: a pooled analysis from the pancreatic cancer cohort consortium. <i>Cancer Causes and Control</i> , 2013, 24, 13-25.	1.8	114
39	Common Genetic Polymorphisms Modify the Effect of Smoking on Absolute Risk of Bladder Cancer. <i>Cancer Research</i> , 2013, 73, 2211-2220.	0.9	107
40	Comprehensive resequencing analysis of a 136 kb region of human chromosome 8q24 associated with prostate and colon cancers. <i>Human Genetics</i> , 2008, 124, 161-170.	3.8	104
41	Pathway analysis of genome-wide association study data highlights pancreatic development genes as susceptibility factors for pancreatic cancer. <i>Carcinogenesis</i> , 2012, 33, 1384-1390.	2.8	102
42	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	6.2	101
43	A genome-wide association study of bladder cancer identifies a new susceptibility locus within SLC14A1, a urea transporter gene on chromosome 18q12.3. <i>Human Molecular Genetics</i> , 2011, 20, 4282-4289.	2.9	100
44	Multilocus Linkage Tests Based on Affected Relative Pairs. <i>American Journal of Human Genetics</i> , 2000, 66, 1273-1286.	6.2	96
45	Fine mapping and functional analysis of a common variant in <i>MSMB</i> on chromosome 10q11.2 associated with prostate cancer susceptibility. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 7933-7938.	7.1	96
46	Genome-wide association study of circulating retinol levels. <i>Human Molecular Genetics</i> , 2011, 20, 4724-4731.	2.9	93
47	Adding Further Power to the Haseman and Elston Method for Detecting Linkage in Larger Sibships: Weighting Sums and Differences. <i>Human Heredity</i> , 2003, 55, 79-85.	0.8	91
48	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633.	2.9	90
49	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	12.8	86
50	Genomewide meta-analysis identifies loci associated with IGF and IGFBP levels with impact on age-related traits. <i>Aging Cell</i> , 2016, 15, 811-824.	6.7	83
51	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , 2012, 21, 456-462.	2.9	81
52	Variant ABO Blood Group Alleles, Secretor Status, and Risk of Pancreatic Cancer: Results from the Pancreatic Cancer Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 3140-3149.	2.5	78
53	Exact transmission-disequilibrium tests with multiallelic markers. <i>Genetic Epidemiology</i> , 1997, 14, 337-347.	1.3	74
54	Genome-Wide Association Study of Circulating Estradiol, Testosterone, and Sex Hormone-Binding Globulin in Postmenopausal Women. <i>PLoS ONE</i> , 2012, 7, e37815.	2.5	61

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55	Testicular germ cell tumor susceptibility associated with the UCK2 locus on chromosome 1q23. <i>Human Molecular Genetics</i> , 2013, 22, 2748-2753.	2.9	59
56	Genotypic variants at 2q33 and risk of esophageal squamous cell carcinoma in China: a meta-analysis of genome-wide association studies. <i>Human Molecular Genetics</i> , 2012, 21, 2132-2141.	2.9	58
57	Fine mapping the KLK3 locus on chromosome 19q13.33 associated with prostate cancer susceptibility and PSA levels. <i>Human Genetics</i> , 2011, 129, 675-685.	3.8	50
58	Association between GWAS-identified lung adenocarcinoma susceptibility loci and EGFR mutations in never-smoking Asian women, and comparison with findings from Western populations. <i>Human Molecular Genetics</i> , 2016, 26, ddw414.	2.9	50
59	Improved imputation of common and uncommon SNPs with a new reference set. <i>Nature Genetics</i> , 2012, 44, 6-7.	21.4	45
60	Fine mapping of a region of chromosome 11q13 reveals multiple independent loci associated with risk of prostate cancer. <i>Human Molecular Genetics</i> , 2011, 20, 2869-2878.	2.9	43
61	Large-Scale Pathway-Based Analysis of Bladder Cancer Genome-Wide Association Data from Five Studies of European Background. <i>PLoS ONE</i> , 2012, 7, e29396.	2.5	36
62	Haseman and Elston revisited: The effects of ascertainment and residual familial correlations on power to detect linkage. <i>Genetic Epidemiology</i> , 2000, 19, 456-460.	1.3	30
63	Large-scale fine mapping of the HNF1B locus and prostate cancer risk. <i>Human Molecular Genetics</i> , 2011, 20, 3322-3329.	2.9	28
64	Refining the Prostate Cancer Genetic Association within the <i>JAZF1</i> Gene on Chromosome 7p15.2. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 1349-1355.	2.5	26
65	Comprehensive resequence analysis of a 97 kb region of chromosome 10q11.2 containing the MSMB gene associated with prostate cancer. <i>Human Genetics</i> , 2009, 126, 743-750.	3.8	21
66	Genome-wide linkage scan for genes affecting longitudinal trends in systolic blood pressure. <i>BMC Genetics</i> , 2003, 4, S82.	2.7	18
67	Y chromosome haplogroups and prostate cancer in populations of European and Ashkenazi Jewish ancestry. <i>Human Genetics</i> , 2012, 131, 1173-1185.	3.8	14
68	Pooling Data and Linkage Analysis in the Chromosome 5q Candidate Region for Asthma. <i>Genetic Epidemiology</i> , 2001, 21, S103-8.	1.3	10
69	Genome-Wide Linkage Analysis in a General Population Sample Using $\lambda$ 2A Random Effects (SSARs) Fitted by Gibbs Sampling. <i>Genetic Epidemiology</i> , 2001, 21, S674-9.	1.3	7
70	Reply to "Mosaic loss of chromosome Y in leukocytes matters". <i>Nature Genetics</i> , 2019, 51, 7-9.	21.4	7
71	Improved Evidence for Linkage on 6p and 5p with Retrospective Pooling of Data from Three Asthma Genome Screens. <i>Genetic Epidemiology</i> , 2001, 21, S130-5.	1.3	5
72	Comprehensive resequence analysis of a 123 kb region of chromosome 11q13 associated with prostate cancer. <i>Prostate</i> , 2012, 72, 476-486.	2.3	5

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73	Fine mapping of 14q24.1 breast cancer susceptibility locus. <i>Human Genetics</i> , 2012, 131, 479-490.	3.8	5
74	A Resequencing Analysis of Genomic Loci on Chromosomes 1q32.1, 5p15.33, and 13q22.1 Associated With Pancreatic Cancer Risk. <i>Pancreas</i> , 2013, 42, 209-215.	1.1	5
75	Polymorphisms in genes related to one-carbon metabolism are not related to pancreatic cancer in PanScan and PanC4. <i>Cancer Causes and Control</i> , 2013, 24, 595-602.	1.8	4
76	Improving the power for disease locus detection in affected sib-pair studies by using two-locus analysis and multiple regression methods. <i>Genetic Epidemiology</i> , 1999, 17, S521-6.	1.3	3
77	Model-based and model-free multipoint genome-wide linkage analysis of alcoholism. <i>Genetic Epidemiology</i> , 1999, 17, S175-S180.	1.3	1