

Kevin B Jacobs

List of Publications by Year in descending order

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Version: 2024-02-01

77
papers

28,059
citations

28190

55
h-index

66788

78
g-index

78
all docs

78
docs citations

78
times ranked

34106
citing authors

#	ARTICLE	IF	CITATIONS
1	Reply to "Mosaic loss of chromosome Y in leukocytes matters" TM . <i>Nature Genetics</i> , 2019, 51, 7-9.	9.4	7
2	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.	1.4	50
3	Mosaic loss of chromosome Y is associated with common variation near <i>TCL1A</i> . <i>Nature Genetics</i> , 2016, 48, 563-568.	9.4	134
4	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.	3.0	83
5	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	5.8	86
6	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.	1.5	331
7	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
8	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	13.7	3,823
9	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the <i>TERT-CLPTM1L</i> region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633.	1.4	90
10	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	9.4	1,818
11	Testicular germ cell tumor susceptibility associated with the <i>UCK2</i> locus on chromosome 1q23. <i>Human Molecular Genetics</i> , 2013, 22, 2748-2753.	1.4	59
12	Common Genetic Polymorphisms Modify the Effect of Smoking on Absolute Risk of Bladder Cancer. <i>Cancer Research</i> , 2013, 73, 2211-2220.	0.4	107
13	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.	0.8	114
14	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.	9.4	578
15	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013, 45, 868-876.	9.4	179
16	Meta-analysis identifies four new loci associated with testicular germ cell tumor. <i>Nature Genetics</i> , 2013, 45, 680-685.	9.4	154
17	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.	1.4	116
18	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.	0.8	4

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19	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.	1.5	371
20	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.	0.5	5
21	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.	1.4	58
22	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.	1.3	102
23	Improved imputation of common and uncommon SNPs with a new reference set. <i>Nature Genetics</i> , 2012, 44, 6-7.	9.4	45
24	Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , 2012, 44, 1330-1335.	9.4	286
25	FTO genotype is associated with phenotypic variability of body mass index. <i>Nature</i> , 2012, 490, 267-272.	13.7	383
26	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.	1.4	168
27	Comprehensive resequencing analysis of a 123 kb region of chromosome 11q13 associated with prostate cancer. <i>Prostate</i> , 2012, 72, 476-486.	1.2	5
28	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.	1.4	81
29	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012, 44, 651-658.	9.4	519
30	Y chromosome haplogroups and prostate cancer in populations of European and Ashkenazi Jewish ancestry. <i>Human Genetics</i> , 2012, 131, 1173-1185.	1.8	14
31	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.	2.6	242
32	Fine mapping of 14q24.1 breast cancer susceptibility locus. <i>Human Genetics</i> , 2012, 131, 479-490.	1.8	5
33	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.	1.1	36
34	Genome-Wide Association Study of Circulating Estradiol, Testosterone, and Sex Hormone-Binding Globulin in Postmenopausal Women. <i>PLoS ONE</i> , 2012, 7, e37815.	1.1	61
35	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.	9.4	220
36	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.	1.8	50

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37	Genome-wide association study identifies new prostate cancer susceptibility loci. Human Molecular Genetics, 2011, 20, 3867-3875.	1.4	160
38	Fine mapping of a region of chromosome 11q13 reveals multiple independent loci associated with risk of prostate cancer. Human Molecular Genetics, 2011, 20, 2869-2878.	1.4	43
39	Large-scale fine mapping of the HNF1B locus and prostate cancer risk. Human Molecular Genetics, 2011, 20, 3322-3329.	1.4	28
40	Novel Breast Cancer Susceptibility Locus at 9q31.2: Results of a Genome-Wide Association Study. Journal of the National Cancer Institute, 2011, 103, 425-435.	3.0	225
41	A genome-wide association study of bladder cancer identifies a new susceptibility locus within SLC14A1, a urea transporter gene on chromosome 18q12.3. Human Molecular Genetics, 2011, 20, 4282-4289.	1.4	100
42	Genome-wide association study of circulating retinol levels. Human Molecular Genetics, 2011, 20, 4724-4731.	1.4	93
43	Family history of cancer and risk of pancreatic cancer: A pooled analysis from the Pancreatic Cancer Cohort Consortium (PanScan). International Journal of Cancer, 2010, 127, 1421-1428.	2.3	128
44	Hundreds of variants clustered in genomic loci and biological pathways affect human height. Nature, 2010, 467, 832-838.	13.7	1,789
45	A genome-wide association study identifies pancreatic cancer susceptibility loci on chromosomes 13q22.1, 1q32.1 and 5p15.33. Nature Genetics, 2010, 42, 224-228.	9.4	539
46	Estimation of effect size distribution from genome-wide association studies and implications for future discoveries. Nature Genetics, 2010, 42, 570-575.	9.4	609
47	A shared susceptibility locus in PLCE1 at 10q23 for gastric adenocarcinoma and esophageal squamous cell carcinoma. Nature Genetics, 2010, 42, 764-767.	9.4	453
48	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. Nature Genetics, 2010, 42, 937-948.	9.4	2,634
49	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. Nature Genetics, 2010, 42, 978-984.	9.4	493
50	Refining the Prostate Cancer Genetic Association within the <i>JAZF1</i> Gene on Chromosome 7p15.2. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 1349-1355.	1.1	26
51	Variant ABO Blood Group Alleles, Secretor Status, and Risk of Pancreatic Cancer: Results from the Pancreatic Cancer Cohort Consortium. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 3140-3149.	1.1	78
52	Genome-wide association study of circulating vitamin D levels. Human Molecular Genetics, 2010, 19, 2739-2745.	1.4	700
53	Anthropometric Measures, Body Mass Index, and Pancreatic Cancer. Archives of Internal Medicine, 2010, 170, 791.	4.3	314
54	Performance of Common Genetic Variants in Breast-Cancer Risk Models. New England Journal of Medicine, 2010, 362, 986-993.	13.9	376

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55	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.	3.3	96
56	Comprehensive resequence analysis of a 97Åkb region of chromosome 10q11.2 containing the <i>MSMB</i> gene associated with prostate cancer. <i>Human Genetics</i> , 2009, 126, 743-750.	1.8	21
57	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
58	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.	9.4	487
59	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.	9.4	597
60	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , 2009, 41, 1055-1057.	9.4	218
61	Comprehensive resequence analysis of a 136Åkb region of human chromosome 8q24 associated with prostate and colon cancers. <i>Human Genetics</i> , 2008, 124, 161-170.	1.8	104
62	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
63	Common variants near <i>MC4R</i> are associated with fat mass, weight and risk of obesity. <i>Nature Genetics</i> , 2008, 40, 768-775.	9.4	1,179
64	Multiple loci identified in a genome-wide association study of prostate cancer. <i>Nature Genetics</i> , 2008, 40, 310-315.	9.4	871
65	Genome-wide association study of prostate cancer identifies a second risk locus at 8q24. <i>Nature Genetics</i> , 2007, 39, 645-649.	9.4	1,059
66	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.	9.4	1,370
67	Genome-wide linkage scan for genes affecting longitudinal trends in systolic blood pressure. <i>BMC Genetics</i> , 2003, 4, S82.	2.7	18
68	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.	0.4	91
69	Genome-Wide Linkage Analysis in a General Population Sample Using γ 2A Random Effects (SSARs) Fitted by Gibbs Sampling. <i>Genetic Epidemiology</i> , 2001, 21, S674-9.	0.6	7
70	Pooling Data and Linkage Analysis in the Chromosome 5q Candidate Region for Asthma. <i>Genetic Epidemiology</i> , 2001, 21, S103-8.	0.6	10
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.	0.6	5
72	Haseman and Elston revisited. <i>Genetic Epidemiology</i> , 2000, 19, 1-17.	0.6	324

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73	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.	0.6	30
74	Multilocus Linkage Tests Based on Affected Relative Pairs. <i>American Journal of Human Genetics</i> , 2000, 66, 1273-1286.	2.6	96
75	Model-based and model-free multipoint genome-wide linkage analysis of alcoholism. <i>Genetic Epidemiology</i> , 1999, 17, S175-S180.	0.6	1
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.	0.6	3
77	Exact transmission-disequilibrium tests with multiallelic markers. <i>Genetic Epidemiology</i> , 1997, 14, 337-347.	0.6	74