

Scott M Williams

List of Publications by Year in descending order

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136
papers

8,806
citations

71102

41
h-index

46799

89
g-index

140
all docs

140
docs citations

140
times ranked

15234
citing authors

#	ARTICLE	IF	CITATIONS
1	Expanding human variation at PLOS Genetics. PLoS Genetics, 2022, 18, e1010070.	3.5	0
2	Liberal-arts education helps scientists think and communicate. Nature, 2022, 603, 578-578.	27.8	2
3	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. American Journal of Human Genetics, 2021, 108, 564-582.	6.2	18
4	Global variation in sequencing impedes SARS-CoV-2 surveillance. PLoS Genetics, 2021, 17, e1009620.	3.5	18
5	Helicobacter pylori infection causes both protective and deleterious effects in human health and disease. Genes and Immunity, 2021, 22, 218-226.	4.1	25
6	Ornithine decarboxylase (ODC1) gene variant (rs2302615) is associated with gastric cancer independently of Helicobacter pylori CagA serostatus. Oncogene, 2021, 40, 5963-5969.	5.9	2
7	Resistance to TST/IGRA conversion in Uganda: Heritability and Genome-Wide Association Study. EBioMedicine, 2021, 74, 103727.	6.1	9
8	Estimating prevalence of human traits among populations from polygenic risk scores. Human Genomics, 2021, 15, 70.	2.9	5
9	Dissecting maternal and fetal genetic effects underlying the associations between maternal phenotypes, birth outcomes, and adult phenotypes: A mendelian-randomization and haplotype-based genetic score analysis in 10,734 motherâ€“infant pairs. PLoS Medicine, 2020, 17, e1003305.	8.4	37
10	CLEC4E (Mincle) genetic variation associates with pulmonary tuberculosis in Guinea-Bissau (West) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	2.3	5
11	Genetics and evolution of tuberculosis pathogenesis: New perspectives and approaches. Infection, Genetics and Evolution, 2020, 81, 104204.	2.3	24
12	Interaction between host genes and Mycobacterium tuberculosis lineage can affect tuberculosis severity: Evidence for coevolution?. PLoS Genetics, 2020, 16, e1008728.	3.5	40
13	A Novel Mapping Strategy Utilizing Mouse Chromosome Substitution Strains Identifies Multiple Epistatic Interactions That Regulate Complex Traits. G3: Genes, Genomes, Genetics, 2020, 10, 4553-4563.	1.8	4
14	A comparison of two workflows for regulome and transcriptomeâ€“based prioritization of genetic variants associated with myocardial mass. Genetic Epidemiology, 2019, 43, 717-726.	1.3	1
15	The Missing Diversity in Human Genetic Studies. Cell, 2019, 177, 26-31.	28.9	838
16	The Plight of Muntaser Ibrahim. PLoS Genetics, 2019, 15, e1008100.	3.5	1
17	Testing the assumptions of parametric linear models: the need for biological data mining in disciplines such as human genetics. BioData Mining, 2019, 12, 6.	4.0	3
18	Evaluating the strength of genetic results: Risks and responsibilities. PLoS Genetics, 2019, 15, e1008437.	3.5	1

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19	Research to achieve a reduction in the global rate of preterm birth needs attention: Preface to the special issue by the preterm Birth International Collaborative (PREBIC). <i>Placenta</i> , 2019, 79, 1-2.	1.5	8
20	Cytochrome P450 epoxygenases and cancer: A genetic and a molecular perspective. , 2019, 196, 183-194.		23
21	The ubiquity of pleiotropy in human disease. <i>Human Genetics</i> , 2018, 137, 39-44.	3.8	81
22	Doubling down on forensic twin studies. <i>PLoS Genetics</i> , 2018, 14, e1007831.	3.5	0
23	Tipping the Scale Towards Gastric Disease: a Host-Pathogen Genomic Mismatch?. <i>Current Genetic Medicine Reports</i> , 2018, 6, 199-207.	1.9	7
24	The premature infant gut microbiome during the first 6 weeks of life differs based on gestational maturity at birth. <i>Pediatric Research</i> , 2018, 84, 71-79.	2.3	101
25	Leveraging epigenomics and contactomics data to investigate SNP pairs in GWAS. <i>Human Genetics</i> , 2018, 137, 413-425.	3.8	8
26	The Cytochrome P450 Slow Metabolizers CYP2C9*2 and CYP2C9*3 Directly Regulate Tumorigenesis via Reduced Epoxyeicosatrienoic Acid Production. <i>Cancer Research</i> , 2018, 78, 4865-4877.	0.9	27
27	Whole exome sequencing reveals HSPA1L as a genetic risk factor for spontaneous preterm birth. <i>PLoS Genetics</i> , 2018, 14, e1007394.	3.5	35
28	HS3ST1 genotype regulates antithrombin's inflammomodulatory tone and associates with atherosclerosis. <i>Matrix Biology</i> , 2017, 63, 69-90.	3.6	19
29	Draft Genome Sequences of 13 Colombian <i>Helicobacter pylori</i> Strains Isolated from Pacific Coast and Andean Residents. <i>Genome Announcements</i> , 2017, 5, .	0.8	1
30	Evolutionary Triangulation to Refine Genetic Association Studies of Spontaneous Preterm Birth. <i>American Journal of Perinatology</i> , 2017, 34, 1041-1047.	1.4	1
31	Evolutionarily derived networks to inform disease pathways. <i>Genetic Epidemiology</i> , 2017, 41, 866-875.	1.3	1
32	Height associated variants demonstrate assortative mating in human populations. <i>Scientific Reports</i> , 2017, 7, 15689.	3.3	15
33	Genomics of Human Pulmonary Tuberculosis: from Genes to Pathways. <i>Current Genetic Medicine Reports</i> , 2017, 5, 149-166.	1.9	30
34	Genetic variation in the eicosanoid pathway is associated with non-small-cell lung cancer (NSCLC) survival. <i>PLoS ONE</i> , 2017, 12, e0180471.	2.5	8
35	Up For A Challenge (U4C): Stimulating innovation in breast cancer genetic epidemiology. <i>PLoS Genetics</i> , 2017, 13, e1006945.	3.5	3
36	Genetic Effects on the Correlation Structure of CVD Risk Factors: Exome-Wide Data From a Ghanaian Population. <i>Global Heart</i> , 2017, 12, 133.	2.3	4

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37	A chromosome 5q31.1 locus associates with tuberculin skin test reactivity in HIV-positive individuals from tuberculosis hyper-endemic regions in east Africa. <i>PLoS Genetics</i> , 2017, 13, e1006710.	3.5	28
38	Widespread epistasis regulates glucose homeostasis and gene expression. <i>PLoS Genetics</i> , 2017, 13, e1007025.	3.5	13
39	GEneSTATION 1.0: a synthetic resource of diverse evolutionary and functional genomic data for studying the evolution of pregnancy-associated tissues and phenotypes. <i>Nucleic Acids Research</i> , 2016, 44, D908-D916.	14.5	6
40	Is Isolated Low High-Density Lipoprotein Cholesterol a Cardiovascular Disease Risk Factor?. <i>Circulation: Cardiovascular Quality and Outcomes</i> , 2016, 9, 206-212.	2.2	71
41	Identifying significant gene-environment interactions using a combination of screening testing and hierarchical false discovery rate control. <i>Genetic Epidemiology</i> , 2016, 40, 544-557.	1.3	20
42	Complex Patterns of Association between Pleiotropy and Transcription Factor Evolution. <i>Genome Biology and Evolution</i> , 2016, 8, 3159-3170.	2.5	17
43	Plasminogen Activator Inhibitor-1 and Diagnosis of the Metabolic Syndrome in a West African Population. <i>Journal of the American Heart Association</i> , 2016, 5, .	3.7	21
44	Epigenetic and genetic variation in GATA5 is associated with gastric disease risk. <i>Human Genetics</i> , 2016, 135, 895-906.	3.8	9
45	A Locus at 5q33.3 Confers Resistance to Tuberculosis in Highly Susceptible Individuals. <i>American Journal of Human Genetics</i> , 2016, 98, 514-524.	6.2	78
46	Molecular analyses of circadian gene variants reveal sex-dependent links between depression and clocks. <i>Translational Psychiatry</i> , 2016, 6, e748-e748.	4.8	65
47	Cardiovascular Disease Risk Factors in Ghana during the Rural-to-Urban Transition: A Cross-Sectional Study. <i>PLoS ONE</i> , 2016, 11, e0162753.	2.5	41
48	Using agent-based simulation to understand population dynamics and coevolution in host-pathogen relationships. , 2015, , .		0
49	Genetic Variation and Insulin Resistance in Middle-Aged Chinese Men. <i>Annals of Human Genetics</i> , 2015, 79, 357-365.	0.8	2
50	A Systems Genetics Approach to Dyslipidemia in Children and Adolescents. <i>OMICS A Journal of Integrative Biology</i> , 2015, 19, 248-259.	2.0	5
51	Meta-analysis of Randomized Controlled Trials of Genotype-Guided vs Standard Dosing of Warfarin. <i>Chest</i> , 2015, 148, 701-710.	0.8	26
52	Sex-Specific Parental Effects on Offspring Lipid Levels. <i>Journal of the American Heart Association</i> , 2015, 4, .	3.7	8
53	Pleiotropic Effects of Immune Responses Explain Variation in the Prevalence of Fibroproliferative Diseases. <i>PLoS Genetics</i> , 2015, 11, e1005568.	3.5	17
54	PLOS Genetics Data Sharing Policy: In Pursuit of Functional Utility. <i>PLoS Genetics</i> , 2015, 11, e1005716.	3.5	10

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55	Genetics of Plasminogen Activator Inhibitor-1 (PAI-1) in a Ghanaian Population. <i>PLoS ONE</i> , 2015, 10, e0136379.	2.5	8
56	A Dietary-Wide Association Study (DWAS) of Environmental Metal Exposure in US Children and Adults. <i>PLoS ONE</i> , 2014, 9, e104768.	2.5	43
57	Association between lifestyle-related disorders and visceral fat mass in Japanese males: a hospital based cross-sectional study. <i>Environmental Health and Preventive Medicine</i> , 2014, 19, 429-435.	3.4	1
58	Epiregulin (EREG) and human V-ATPase (TCIRG1): genetic variation, ethnicity and pulmonary tuberculosis susceptibility in Guinea-Bissau and The Gambia. <i>Genes and Immunity</i> , 2014, 15, 370-377.	4.1	11
59	The Association of the Vanin-1 N131S Variant with Blood Pressure Is Mediated by Endoplasmic Reticulum-Associated Degradation and Loss of Function. <i>PLoS Genetics</i> , 2014, 10, e1004641.	3.5	16
60	Genome-Wide Association Study for Circulating Tissue Plasminogen Activator Levels and Functional Follow-Up Implicates Endothelial <i>STXBP5</i> and <i>STX2</i> . <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, 1093-1101.	2.4	43
61	Disrupted human- <i>Yersinia enterocolitica</i> pathogen co-evolution: a model for disease. <i>Frontiers in Genetics</i> , 2014, 5, 290.	2.3	50
62	A Single Nucleotide Polymorphism in SLC7A5 Is Associated with Gastrointestinal Toxicity after High-Dose Melphalan and Autologous Stem Cell Transplantation for Multiple Myeloma. <i>Biology of Blood and Marrow Transplantation</i> , 2014, 20, 1014-1020.	2.0	20
63	The multiscale backbone of the human phenotype network based on biological pathways. <i>BioData Mining</i> , 2014, 7, 1.	4.0	32
64	Human and <i>Helicobacter pylori</i> coevolution shapes the risk of gastric disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 1455-1460.	7.1	198
65	Genetic Variation in the Peroxisome Proliferator-Activated Receptor (PPAR) and Peroxisome Proliferator-Activated Receptor Gamma Co-activator 1 (PGC1) Gene Families and Type 2 Diabetes. <i>Annals of Human Genetics</i> , 2014, 78, 23-32.	0.8	26
66	Genome-wide Association Analysis of Blood-Pressure Traits in African-Ancestry Individuals Reveals Common Associated Genes in African and Non-African Populations. <i>American Journal of Human Genetics</i> , 2013, 93, 545-554.	6.2	189
67	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. <i>Nature Genetics</i> , 2013, 45, 690-696.	21.4	232
68	An information-gain approach to detecting three-way epistatic interactions in genetic association studies. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2013, 20, 630-636.	4.4	69
69	Recurrent Tissue-Specific mtDNA Mutations Are Common in Humans. <i>PLoS Genetics</i> , 2013, 9, e1003929.	3.5	130
70	Admixture Mapping in Lupus Identifies Multiple Functional Variants within IFIH1 Associated with Apoptosis, Inflammation, and Autoantibody Production. <i>PLoS Genetics</i> , 2013, 9, e1003222.	3.5	107
71	Preterm Birth Genome Project (PGP) – validation of resources for preterm birth genome-wide studies. <i>Journal of Perinatal Medicine</i> , 2013, 41, 45-9.	1.4	10
72	A Simple and Computationally Efficient Approach to Multifactor Dimensionality Reduction Analysis of Gene-Gene Interactions for Quantitative Traits. <i>PLoS ONE</i> , 2013, 8, e66545.	2.5	82

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73	Guidelines for Genome-Wide Association Studies. <i>PLoS Genetics</i> , 2012, 8, e1002812.	3.5	88
74	HTR1B, ADIPOR1, PPARGC1A, and CYP19A1 and Obesity in a Cohort of Caucasians and African Americans: An Evaluation of Gene-Environment Interactions and Candidate Genes. <i>American Journal of Epidemiology</i> , 2012, 175, 11-21.	3.4	42
75	High Body Mass Index is an Important Risk Factor for the Development of Type 2 Diabetes. <i>Internal Medicine</i> , 2012, 51, 1821-1826.	0.7	47
76	Joint Effect of Genetic and Lifestyle Risk Factors on Type 2 Diabetes Risk among Chinese Men and Women. <i>PLoS ONE</i> , 2012, 7, e49464.	2.5	16
77	Neighborhood socio-economic characteristics, African ancestry, and <i>Helicobacter pylori</i> sero-prevalence. <i>Cancer Causes and Control</i> , 2012, 23, 897-906.	1.8	19
78	Common Variation in Vitamin D Pathway Genes Predicts Circulating 25-Hydroxyvitamin D Levels among African Americans. <i>PLoS ONE</i> , 2011, 6, e28623.	2.5	103
79	Peroxisome Proliferator-Activated Receptor Delta (PPARD) Genetic Variation and Type 2 Diabetes in Middle-Aged Chinese Women. <i>Annals of Human Genetics</i> , 2011, 75, 621-629.	0.8	16
80	<i>ADIPOQ</i> , <i>ADIPOR1</i> , and <i>ADIPOR2</i> Polymorphisms in Relation to Serum Adiponectin Levels and BMI in Black and White Women. <i>Obesity</i> , 2011, 19, 2053-2062.	3.0	39
81	Race, African Ancestry, and <i>Helicobacter pylori</i> Infection in a Low-Income United States Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 826-834.	2.5	76
82	Epistatic Interactions in Genetic Regulation of t-PA and PAI-1 Levels in a Ghanaian Population. <i>PLoS ONE</i> , 2011, 6, e16639.	2.5	4
83	10.1023/A:1003512607608. , 2011, , .		0
84	Desensitization of vascular response in vivo: contribution of genetic variation in the β_2 -adrenergic receptor subtype. <i>Journal of Hypertension</i> , 2010, 28, 278-284.	0.5	23
85	Genetic Diversity of the Fragile X Syndrome Gene (<i>FMR1</i>) in a Large Sub-Saharan West African Population. <i>Annals of Human Genetics</i> , 2010, 74, 316-325.	0.8	14
86	Blood Vitamin D Levels in Relation to Genetic Estimation of African Ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2325-2331.	2.5	56
87	HbS and HbC associate with malaria transmission: Human genetics links to vaccinology?. <i>Vaccine</i> , 2010, 28, 6403.	3.8	0
88	AGT M235T Genotype/Anxiety Interaction and Gender in the HyperGEN Study. <i>PLoS ONE</i> , 2010, 5, e13353.	2.5	4
89	Failure to Replicate a Genetic Association May Provide Important Clues About Genetic Architecture. <i>PLoS ONE</i> , 2009, 4, e5639.	2.5	227
90	Genetic Population Structure Analysis in New Hampshire Reveals Eastern European Ancestry. <i>PLoS ONE</i> , 2009, 4, e6928.	2.5	4

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91	Shadows of complexity: what biological networks reveal about epistasis and pleiotropy. <i>BioEssays</i> , 2009, 31, 220-227.	2.5	162
92	Genetic variants of GSNOR and ADRB2 influence response to albuterol in African-American children with severe asthma. <i>Pediatric Pulmonology</i> , 2009, 44, 649-654.	2.0	61
93	The Genetic Structure and History of Africans and African Americans. <i>Science</i> , 2009, 324, 1035-1044.	12.6	1,267
94	Epistasis and Its Implications for Personal Genetics. <i>American Journal of Human Genetics</i> , 2009, 85, 309-320.	6.2	326
95	Male-female differences in the genetic regulation of t-PA and PAI-1 levels in a Ghanaian population. <i>Human Genetics</i> , 2008, 124, 479-488.	3.8	21
96	Genetic regulation of cervical antiinflammatory cytokine concentrations during pregnancy. <i>American Journal of Obstetrics and Gynecology</i> , 2008, 199, 163.e1-163.e11.	1.3	12
97	β_2 -Adrenergic Receptor Promoter Haplotype Influences Spirometric Response During an Acute Asthma Exacerbation. <i>Clinical and Translational Science</i> , 2008, 1, 155-161.	3.1	5
98	Calculation and Use of the Hardy-Weinberg Model in Association Studies. <i>Current Protocols in Human Genetics</i> , 2008, 57, Unit 1.18.	3.5	33
99	Genetic Differences in Human Circadian Clock Genes among Worldwide Populations. <i>Journal of Biological Rhythms</i> , 2008, 23, 330-340.	2.6	108
100	Cytokine polymorphisms and gastric cancer risk: An evolving view. <i>Cancer Biology and Therapy</i> , 2008, 7, 157-162.	3.4	21
101	Beta-1-adrenoceptor genetic variants and ethnicity independently affect response to beta-blockade. <i>Pharmacogenetics and Genomics</i> , 2008, 18, 895-902.	1.5	48
102	Lumping, splitting and mapping: assessing linkage in different ethnic groups for albuminuria and glomerular filtration rate in the HyperGen study. <i>Nephrology Dialysis Transplantation</i> , 2007, 22, 687-689.	0.7	0
103	Epistatic effects of polymorphisms in genes from the renin-angiotensin, bradykinin, and fibrinolytic systems on plasma t-PA and PAI-1 levels. <i>Genomics</i> , 2007, 89, 362-369.	2.9	30
104	Gender-specific correlations of plasminogen activator inhibitor-1 and tissue plasminogen activator levels with cardiovascular disease-related traits. <i>Journal of Thrombosis and Haemostasis</i> , 2007, 5, 313-320.	3.8	44
105	The effects of polymorphisms in genes from the renin-angiotensin, bradykinin, and fibrinolytic systems on plasma t-PA and PAI-1 levels are dependent on environmental context. <i>Human Genetics</i> , 2007, 122, 275-281.	3.8	16
106	A population-based study in Ghana to investigate inter-individual variation in plasma t-PA and PAI-1. <i>Ethnicity and Disease</i> , 2007, 17, 492-7.	2.3	10
107	Genomics, Nutrition, Obesity, and Diabetes. <i>Journal of Nursing Scholarship</i> , 2006, 38, 11-18.	2.4	11
108	Variations in the β_2 -adrenergic receptor gene and their functional effects. <i>Clinical Pharmacology and Therapeutics</i> , 2006, 79, 173-185.	4.7	28

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109	Elevated male European and female African contributions to the genomes of African American individuals. <i>Human Genetics</i> , 2006, 120, 713-722.	3.8	84
110	Single-Nucleotide Polymorphisms for Diagnosis of Salt-Sensitive Hypertension. <i>Clinical Chemistry</i> , 2006, 52, 352-360.	3.2	103
111	Variation in the β_2 -adrenergic receptor gene (ADRA2B) and its relationship to vascular response in vivo. <i>Pharmacogenetics and Genomics</i> , 2005, 15, 407-414.	1.5	31
112	Traversing the conceptual divide between biological and statistical epistasis: systems biology and a more modern synthesis. <i>BioEssays</i> , 2005, 27, 637-646.	2.5	301
113	Endothelial NO Synthase Polymorphisms and Postural Tachycardia Syndrome. <i>Hypertension</i> , 2005, 46, 1103-1110.	2.7	36
114	Reporting of model validation procedures in human studies of genetic interactions. <i>Nutrition</i> , 2004, 20, 69-73.	2.4	24
115	The use of animal models in the study of complex disease: all else is never equal or why do so many human studies fail to replicate animal findings?. <i>BioEssays</i> , 2004, 26, 170-179.	2.5	81
116	Multilocus Analysis of Hypertension: A Hierarchical Approach. <i>Human Heredity</i> , 2004, 57, 28-38.	0.8	146
117	Genetic markers associated with resistance to infectious hematopoietic necrosis in rainbow and steelhead trout (<i>Oncorhynchus mykiss</i>) backcrosses. <i>Aquaculture</i> , 2004, 241, 93-115.	3.5	63
118	A High-Density Admixture Map for Disease Gene Discovery in African Americans. <i>American Journal of Human Genetics</i> , 2004, 74, 1001-1013.	6.2	416
119	Methyl-group dietary intake and risk of breast cancer among African-American women: a case-control study by methylation status of the estrogen receptor alpha genes. <i>Cancer Causes and Control</i> , 2003, 14, 827-836.	1.8	48
120	Ethnic diversity in a critical gene responsible for glutathione synthesis. <i>Free Radical Biology and Medicine</i> , 2003, 34, 72-76.	2.9	19
121	Common single nucleotide polymorphisms in the promoter region of the human factor XI gene. <i>Journal of Thrombosis and Haemostasis</i> , 2003, 1, 1854-1856.	3.8	7
122	5' Flanking Variants of Resistin Are Associated With Obesity. <i>Diabetes</i> , 2002, 51, 1629-1634.	0.6	158
123	New strategies for identifying gene-gene interactions in hypertension. <i>Annals of Medicine</i> , 2002, 34, 88-95.	3.8	377
124	Phylogeny of the tropical tree family Dipterocarpaceae based on nucleotide sequences of the chloroplast RBCL gene. <i>American Journal of Botany</i> , 1999, 86, 1182-1190.	1.7	125
125	Methyl-deficient diets, methylated ER genes and breast cancer: an hypothesized association. , 1998, 9, 615-620.		22
126	Estrogen receptor status of breast cancer: a marker of different stages of tumor or different entities of the disease?. <i>Medical Hypotheses</i> , 1997, 49, 69-75.	1.5	25

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127	High Density of an SAR-Associated Motif Differentiates Heterochromatin From Euchromatin. <i>Journal of Theoretical Biology</i> , 1996, 183, 159-167.	1.7	13
128	Comparative reproductive success of communally breeding burying beetles as assessed by PCR with randomly amplified polymorphic DNA.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1993, 90, 2242-2245.	7.1	66
129	Parentage analysis using RAPD PCR. <i>Nucleic Acids Research</i> , 1992, 20, 5493-5493.	14.5	74
130	Molecular genetic analysis of <i>Drosophila</i> rDNA arrays. <i>Trends in Genetics</i> , 1992, 8, 335-340.	6.7	37
131	Superstructure of the <i>Drosophila</i> ribosomal gene family.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1990, 87, 3156-3160.	7.1	18
132	Chloroplast DNA polymorphisms in lodgepole and jack pines and their hybrids.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1987, 84, 2097-2100.	7.1	409
133	Differences in life history traits between alcohol dehydrogenase genotypes of <i>Drosophila mercatorum</i> : background and maternal genotype effects. <i>Genetica</i> , 1987, 74, 149-153.	1.1	0
134	Sister chromatid exchange and the evolution of rDNA spacer length. <i>Journal of Theoretical Biology</i> , 1985, 116, 625-636.	1.7	25
135	The maintenance of polymorphism owing to differences in developmental time and competition. <i>Genome</i> , 1985, 27, 328-333.	0.7	4
136	On the applicability of game theory to evolution: A response. <i>Journal of Theoretical Biology</i> , 1981, 91, 603-605.	1.7	1