

# Scott M Williams

## List of Publications by Year in descending order

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135  
papers

8,806  
citations

70961

41  
h-index

46693

89  
g-index

140  
all docs

140  
docs citations

140  
times ranked

15234  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Genetic Structure and History of Africans and African Americans. <i>Science</i> , 2009, 324, 1035-1044.	6.0	1,267
2	The Missing Diversity in Human Genetic Studies. <i>Cell</i> , 2019, 177, 26-31.	13.5	838
3	A High-Density Admixture Map for Disease Gene Discovery in African Americans. <i>American Journal of Human Genetics</i> , 2004, 74, 1001-1013.	2.6	416
4	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.	3.3	409
5	New strategies for identifying gene-gene interactions in hypertension. <i>Annals of Medicine</i> , 2002, 34, 88-95.	1.5	377
6	Epistasis and Its Implications for Personal Genetics. <i>American Journal of Human Genetics</i> , 2009, 85, 309-320.	2.6	326
7	Traversing the conceptual divide between biological and statistical epistasis: systems biology and a more modern synthesis. <i>BioEssays</i> , 2005, 27, 637-646.	1.2	301
8	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. <i>Nature Genetics</i> , 2013, 45, 690-696.	9.4	232
9	Failure to Replicate a Genetic Association May Provide Important Clues About Genetic Architecture. <i>PLoS ONE</i> , 2009, 4, e5639.	1.1	227
10	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.	3.3	198
11	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.	2.6	189
12	Shadows of complexity: what biological networks reveal about epistasis and pleiotropy. <i>BioEssays</i> , 2009, 31, 220-227.	1.2	162
13	5' Flanking Variants of Resistin Are Associated With Obesity. <i>Diabetes</i> , 2002, 51, 1629-1634.	0.3	158
14	Multilocus Analysis of Hypertension: A Hierarchical Approach. <i>Human Heredity</i> , 2004, 57, 28-38.	0.4	146
15	Recurrent Tissue-Specific mtDNA Mutations Are Common in Humans. <i>PLoS Genetics</i> , 2013, 9, e1003929.	1.5	130
16	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.	0.8	125
17	Genetic Differences in Human Circadian Clock Genes among Worldwide Populations. <i>Journal of Biological Rhythms</i> , 2008, 23, 330-340.	1.4	108
18	Admixture Mapping in Lupus Identifies Multiple Functional Variants within IFIH1 Associated with Apoptosis, Inflammation, and Autoantibody Production. <i>PLoS Genetics</i> , 2013, 9, e1003222.	1.5	107

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19	Single-Nucleotide Polymorphisms for Diagnosis of Salt-Sensitive Hypertension. <i>Clinical Chemistry</i> , 2006, 52, 352-360.	1.5	103
20	Common Variation in Vitamin D Pathway Genes Predicts Circulating 25-Hydroxyvitamin D Levels among African Americans. <i>PLoS ONE</i> , 2011, 6, e28623.	1.1	103
21	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.	1.1	101
22	Guidelines for Genome-Wide Association Studies. <i>PLoS Genetics</i> , 2012, 8, e1002812.	1.5	88
23	Elevated male European and female African contributions to the genomes of African American individuals. <i>Human Genetics</i> , 2006, 120, 713-722.	1.8	84
24	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.	1.1	82
25	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.	1.2	81
26	The ubiquity of pleiotropy in human disease. <i>Human Genetics</i> , 2018, 137, 39-44.	1.8	81
27	A Locus at 5q33.3 Confers Resistance to Tuberculosis in Highly Susceptible Individuals. <i>American Journal of Human Genetics</i> , 2016, 98, 514-524.	2.6	78
28	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.	1.1	76
29	Parentage analysis using RAPD PCR. <i>Nucleic Acids Research</i> , 1992, 20, 5493-5493.	6.5	74
30	Is Isolated Low High-Density Lipoprotein Cholesterol a Cardiovascular Disease Risk Factor?. <i>Circulation: Cardiovascular Quality and Outcomes</i> , 2016, 9, 206-212.	0.9	71
31	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.	2.2	69
32	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.	3.3	66
33	Molecular analyses of circadian gene variants reveal sex-dependent links between depression and clocks. <i>Translational Psychiatry</i> , 2016, 6, e748-e748.	2.4	65
34	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.	1.7	63
35	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.	1.0	61
36	Blood Vitamin D Levels in Relation to Genetic Estimation of African Ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2325-2331.	1.1	56

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37	Disrupted human- <i>ŕ</i> pathogen co-evolution: a model for disease. <i>Frontiers in Genetics</i> , 2014, 5, 290.	1.1	50
38	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.	0.8	48
39	Beta-1-adrenoceptor genetic variants and ethnicity independently affect response to beta-blockade. <i>Pharmacogenetics and Genomics</i> , 2008, 18, 895-902.	0.7	48
40	High Body Mass Index is an Important Risk Factor for the Development of Type 2 Diabetes. <i>Internal Medicine</i> , 2012, 51, 1821-1826.	0.3	47
41	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.	1.9	44
42	A Dietary-Wide Association Study (DWAS) of Environmental Metal Exposure in US Children and Adults. <i>PLoS ONE</i> , 2014, 9, e104768.	1.1	43
43	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.	1.1	43
44	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.	1.6	42
45	Cardiovascular Disease Risk Factors in Ghana during the Rural-to-Urban Transition: A Cross-Sectional Study. <i>PLoS ONE</i> , 2016, 11, e0162753.	1.1	41
46	Interaction between host genes and <i>Mycobacterium tuberculosis</i> lineage can affect tuberculosis severity: Evidence for coevolution?. <i>PLoS Genetics</i> , 2020, 16, e1008728.	1.5	40
47	<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.	1.5	39
48	Molecular genetic analysis of <i>Drosophila</i> rDNA arrays. <i>Trends in Genetics</i> , 1992, 8, 335-340.	2.9	37
49	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 mother-infant pairs. <i>PLoS Medicine</i> , 2020, 17, e1003305.	3.9	37
50	Endothelial NO Synthase Polymorphisms and Postural Tachycardia Syndrome. <i>Hypertension</i> , 2005, 46, 1103-1110.	1.3	36
51	Whole exome sequencing reveals HSPA1L as a genetic risk factor for spontaneous preterm birth. <i>PLoS Genetics</i> , 2018, 14, e1007394.	1.5	35
52	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
53	The multiscale backbone of the human phenotype network based on biological pathways. <i>BioData Mining</i> , 2014, 7, 1.	2.2	32
54	Variation in the $\beta$ -adrenergic receptor gene ( <i>ADRA2B</i> ) and its relationship to vascular response in vivo. <i>Pharmacogenetics and Genomics</i> , 2005, 15, 407-414.	0.7	31

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55	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.	1.3	30
56	Genomics of Human Pulmonary Tuberculosis: from Genes to Pathways. <i>Current Genetic Medicine Reports</i> , 2017, 5, 149-166.	1.9	30
57	Variations in the $\beta$ 2A-adrenergic receptor gene and their functional effects. <i>Clinical Pharmacology and Therapeutics</i> , 2006, 79, 173-185.	2.3	28
58	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.	1.5	28
59	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.	0.4	27
60	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.	0.3	26
61	Meta-analysis of Randomized Controlled Trials of Genotype-Guided vs Standard Dosing of Warfarin. <i>Chest</i> , 2015, 148, 701-710.	0.4	26
62	Sister chromatid exchange and the evolution of rDNA spacer length. <i>Journal of Theoretical Biology</i> , 1985, 116, 625-636.	0.8	25
63	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.	0.8	25
64	<i>Helicobacter pylori</i> infection causes both protective and deleterious effects in human health and disease. <i>Genes and Immunity</i> , 2021, 22, 218-226.	2.2	25
65	Reporting of model validation procedures in human studies of genetic interactions. <i>Nutrition</i> , 2004, 20, 69-73.	1.1	24
66	Genetics and evolution of tuberculosis pathogenesis: New perspectives and approaches. <i>Infection, Genetics and Evolution</i> , 2020, 81, 104204.	1.0	24
67	Desensitization of vascular response in vivo: contribution of genetic variation in the $\beta$ 2B-adrenergic receptor subtype. <i>Journal of Hypertension</i> , 2010, 28, 278-284.	0.3	23
68	Cytochrome P450 epoxygenases and cancer: A genetic and a molecular perspective. , 2019, 196, 183-194.		23
69	Methyl-deficient diets, methylated ER genes and breast cancer: an hypothesized association. , 1998, 9, 615-620.		22
70	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.	1.8	21
71	Cytokine polymorphisms and gastric cancer risk: An evolving view. <i>Cancer Biology and Therapy</i> , 2008, 7, 157-162.	1.5	21
72	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, .	1.6	21

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73	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
74	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.	0.6	20
75	Ethnic diversity in a critical gene responsible for glutathione synthesis. <i>Free Radical Biology and Medicine</i> , 2003, 34, 72-76.	1.3	19
76	Neighborhood socio-economic characteristics, African ancestry, and <i>Helicobacter pylori</i> sero-prevalence. <i>Cancer Causes and Control</i> , 2012, 23, 897-906.	0.8	19
77	HS3ST1 genotype regulates antithrombin's inflammomodulatory tone and associates with atherosclerosis. <i>Matrix Biology</i> , 2017, 63, 69-90.	1.5	19
78	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.	3.3	18
79	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021, 108, 564-582.	2.6	18
80	Global variation in sequencing impedes SARS-CoV-2 surveillance. <i>PLoS Genetics</i> , 2021, 17, e1009620.	1.5	18
81	Complex Patterns of Association between Pleiotropy and Transcription Factor Evolution. <i>Genome Biology and Evolution</i> , 2016, 8, 3159-3170.	1.1	17
82	Pleiotropic Effects of Immune Responses Explain Variation in the Prevalence of Fibroproliferative Diseases. <i>PLoS Genetics</i> , 2015, 11, e1005568.	1.5	17
83	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.	1.8	16
84	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.	0.3	16
85	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.	1.1	16
86	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.	1.5	16
87	Height associated variants demonstrate assortative mating in human populations. <i>Scientific Reports</i> , 2017, 7, 15689.	1.6	15
88	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.	0.3	14
89	High Density of an SAR-Associated Motif Differentiates Heterochromatin From Euchromatin. <i>Journal of Theoretical Biology</i> , 1996, 183, 159-167.	0.8	13
90	Widespread epistasis regulates glucose homeostasis and gene expression. <i>PLoS Genetics</i> , 2017, 13, e1007025.	1.5	13

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91	Genetic regulation of cervical antiinflammatory cytokine concentrations during pregnancy. <i>American Journal of Obstetrics and Gynecology</i> , 2008, 199, 163.e1-163.e11.	0.7	12
92	Genomics, Nutrition, Obesity, and Diabetes. <i>Journal of Nursing Scholarship</i> , 2006, 38, 11-18.	1.1	11
93	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.	2.2	11
94	Preterm Birth Genome Project (PGP) – validation of resources for preterm birth genome-wide studies. <i>Journal of Perinatal Medicine</i> , 2013, 41, 45-9.	0.6	10
95	PLOS Genetics Data Sharing Policy: In Pursuit of Functional Utility. <i>PLoS Genetics</i> , 2015, 11, e1005716.	1.5	10
96	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.	1.0	10
97	Epigenetic and genetic variation in GATA5 is associated with gastric disease risk. <i>Human Genetics</i> , 2016, 135, 895-906.	1.8	9
98	Resistance to TST/IGRA conversion in Uganda: Heritability and Genome-Wide Association Study. <i>EBioMedicine</i> , 2021, 74, 103727.	2.7	9
99	Sex-specific Parental Effects on Offspring Lipid Levels. <i>Journal of the American Heart Association</i> , 2015, 4, .	1.6	8
100	Genetic variation in the eicosanoid pathway is associated with non-small-cell lung cancer (NSCLC) survival. <i>PLoS ONE</i> , 2017, 12, e0180471.	1.1	8
101	Leveraging epigenomics and contactomics data to investigate SNP pairs in GWAS. <i>Human Genetics</i> , 2018, 137, 413-425.	1.8	8
102	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.	0.7	8
103	Genetics of Plasminogen Activator Inhibitor-1 (PAI-1) in a Ghanaian Population. <i>PLoS ONE</i> , 2015, 10, e0136379.	1.1	8
104	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.	1.9	7
105	Tipping the Scale Towards Gastric Disease: a Host-Pathogen Genomic Mismatch?. <i>Current Genetic Medicine Reports</i> , 2018, 6, 199-207.	1.9	7
106	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.	6.5	6
107	$\beta_2$ -Adrenergic Receptor Promoter Haplotype Influences Spirometric Response During an Acute Asthma Exacerbation. <i>Clinical and Translational Science</i> , 2008, 1, 155-161.	1.5	5
108	A Systems Genetics Approach to Dyslipidemia in Children and Adolescents. <i>OMICS A Journal of Integrative Biology</i> , 2015, 19, 248-259.	1.0	5

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109	CLEC4E (Mincle) genetic variation associates with pulmonary tuberculosis in Guinea-Bissau (West) Tj ETQq1 1 0.784314 rgBT <sub>5</sub> /Overlock	1.0	5
110	Estimating prevalence of human traits among populations from polygenic risk scores. Human Genomics, 2021, 15, 70.	1.4	5
111	The maintenance of polymorphism owing to differences in developmental time and competition. Genome, 1985, 27, 328-333.	0.7	4
112	Genetic Population Structure Analysis in New Hampshire Reveals Eastern European Ancestry. PLoS ONE, 2009, 4, e6928.	1.1	4
113	Genetic Effects on the Correlation Structure of CVD Risk Factors: Exome-Wide Data From a Ghanaian Population. Global Heart, 2017, 12, 133.	0.9	4
114	AGT M235T Genotype/Anxiety Interaction and Gender in the HyperGEN Study. PLoS ONE, 2010, 5, e13353.	1.1	4
115	Epistatic Interactions in Genetic Regulation of t-PA and PAI-1 Levels in a Ghanaian Population. PLoS ONE, 2011, 6, e16639.	1.1	4
116	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.	0.8	4
117	Up For A Challenge (U4C): Stimulating innovation in breast cancer genetic epidemiology. PLoS Genetics, 2017, 13, e1006945.	1.5	3
118	Testing the assumptions of parametric linear models: the need for biological data mining in disciplines such as human genetics. BioData Mining, 2019, 12, 6.	2.2	3
119	Genetic Variation and Insulin Resistance in Middle-aged Chinese Men. Annals of Human Genetics, 2015, 79, 357-365.	0.3	2
120	Ornithine decarboxylase (ODC1) gene variant (rs2302615) is associated with gastric cancer independently of Helicobacter pylori CagA serostatus. Oncogene, 2021, 40, 5963-5969.	2.6	2
121	Liberal-arts education helps scientists think and communicate. Nature, 2022, 603, 578-578.	13.7	2
122	On the applicability of game theory to evolution: A response. Journal of Theoretical Biology, 1981, 91, 603-605.	0.8	1
123	Association between lifestyle-related disorders and visceral fat mass in Japanese males: a hospital based cross-sectional study. Environmental Health and Preventive Medicine, 2014, 19, 429-435.	1.4	1
124	Draft Genome Sequences of 13 Colombian Helicobacter pylori Strains Isolated from Pacific Coast and Andean Residents. Genome Announcements, 2017, 5, .	0.8	1
125	Evolutionary Triangulation to Refine Genetic Association Studies of Spontaneous Preterm Birth. American Journal of Perinatology, 2017, 34, 1041-1047.	0.6	1
126	Evolutionarily derived networks to inform disease pathways. Genetic Epidemiology, 2017, 41, 866-875.	0.6	1



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127	A comparison of two workflows for regulome and transcriptome-based prioritization of genetic variants associated with myocardial mass. <i>Genetic Epidemiology</i> , 2019, 43, 717-726.	0.6	1
128	The Plight of Muntaser Ibrahim. <i>PLoS Genetics</i> , 2019, 15, e1008100.	1.5	1
129	Evaluating the strength of genetic results: Risks and responsibilities. <i>PLoS Genetics</i> , 2019, 15, e1008437.	1.5	1
130	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.	0.5	0
131	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.	0.4	0
132	HbS and HbC associate with malaria transmission: Human genetics links to vaccinology?. <i>Vaccine</i> , 2010, 28, 6403.	1.7	0
133	Using agent-based simulation to understand population dynamics and coevolution in host-pathogen relationships. , 2015, , .		0
134	Doubling down on forensic twin studies. <i>PLoS Genetics</i> , 2018, 14, e1007831.	1.5	0
135	Expanding human variation at PLOS Genetics. <i>PLoS Genetics</i> , 2022, 18, e1010070.	1.5	0