Adebowale A Adeyemo

List of Publications by Year
in descending order

195 papers	19,487 citations	26567 56 h-index	12233 133 g-index
207 all docs	207 docs citations	207 times ranked	26826 citing authors

#	Article	IF	Citations
1	Polygenic Prediction of Type 2 Diabetes in Africa. Diabetes Care, 2022, 45, 717-723.	4.3	12
2	Beta-cell dysfunction and insulin resistance in relation to abnormal glucose tolerance in African populations: can we afford to ignore the diversity within African populations?. BMJ Open Diabetes Research and Care, 2022, 10, e002685.	1.2	0
3	The Lancet Nigeria Commission: investing in health and the future of the nation. Lancet, The, 2022, 399, 1155-1200.	6.3	87
4	Polygenic risk scores for CARDINAL study. Nature Genetics, 2022, 54, 527-530.	9.4	5
5	Additive genetic effect of GCKR, G6PC2, and SLC30A8 variants on fasting glucose levels and risk of type 2 diabetes. PLoS ONE, 2022, 17, e0269378.	1.1	4
6	An epigenome-wide association study of insulin resistance in African Americans. Clinical Epigenetics, 2022, 14, .	1.8	3
7	Endemic Burkitt Lymphoma in second-degree relatives in Northern Uganda: in-depth genome-wide analysis suggests clues about genetic susceptibility. Leukemia, 2021, 35, 1209-1213.	3.3	5
8	From one human genome to a complex tapestry of ancestry. Nature, 2021, 590, 220-221.	13.7	10
9	Exome Sequencing and Congenital Heart Disease in Sub-Saharan Africa. Circulation Genomic and Precision Medicine, 2021, 14, e003108.	1.6	16
10	Genetic risk scores for cardiometabolic traits in sub-Saharan African populations. International Journal of Epidemiology, 2021, 50, 1283-1296.	0.9	10
11	Evolutionary forces in diabetes and hypertension pathogenesis in Africans. Human Molecular Genetics, 2021, 30, R110-R118.	1.4	6
12	Plasma cell immunoglobulin heavy chain repertoire dynamics before and after tetanus booster vaccination. Immunogenetics, 2021, 73, 321-332.	1.2	0
13	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.	2.6	18
14	DNA methylation as the link between migration and theÂmajor noncommunicable diseases: the RODAM study. Epigenomics, 2021, 13, 653-666.	1.0	5
15	A UGT1A1 variant is associated with serum total bilirubin levels, which are causal for hypertension in African-ancestry individuals. Npj Genomic Medicine, 2021, 6, 44.	1.7	6
16	Genome-wide DNA methylation analysis on C-reactive protein among Ghanaians suggests molecular links to the emerging risk of cardiovascular diseases. Npj Genomic Medicine, 2021, 6, 46.	1.7	4
17	Serum fructosamine and glycemic status in the presence of the sickle cell mutation. Diabetes Research and Clinical Practice, 2021, 177, 108918.	1.1	4
18	GWAS in Africans identifies novel lipids loci and demonstrates heterogenous association within Africa. Human Molecular Genetics, 2021, 30, 2205-2214.	1.4	6

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19	Genome-Wide Scan for Parent-of-Origin Effects in a sub-Saharan African Cohort With Nonsyndromic Cleft Lip and/or Cleft Palate (CL/P). Cleft Palate-Craniofacial Journal, 2021, , 105566562110363.	0.5	1
20	Evolutionary genetics and acclimatization in nephrology. Nature Reviews Nephrology, 2021, 17, 827-839.	4.1	5
21	Genome-wide analyses of multiple obesity-related cytokines and hormones informs biology of cardiometabolic traits. Genome Medicine, 2021, 13, 156.	3.6	6
22	Responsible use of polygenic risk scores in the clinic: potential benefits, risks and gaps. Nature Medicine, 2021, 27, 1876-1884.	15.2	214
23	Epigenome-wide association study of serum urate reveals insights into urate co-regulation and the SLC2A9 locus. Nature Communications, 2021, 12, 7173.	5.8	8
24	Meta-analyses identify DNA methylation associated with kidney function and damage. Nature Communications, 2021, 12, 7174.	5.8	30
25	Refining genome-wide associated loci for serum uric acid in individuals with African ancestry. Human Molecular Genetics, 2020, 29, 506-514.	1.4	6
26	Prevalence of vitamin D deficiency in Africa: a systematic review and meta-analysis. The Lancet Global Health, 2020, 8, e134-e142.	2.9	150
27	High-depth African genomes inform human migration and health. Nature, 2020, 586, 741-748.	13.7	197
28	The Genetics of Circulating Resistin Level, A Biomarker for Cardiovascular Diseases, Is Informed by Mendelian Randomization and the Unique Characteristics of African Genomes. Circulation Genomic and Precision Medicine, 2020, 13, 488-503.	1.6	1
29	Genetic modifiers of longâ€term survival in sickle cell anemia. Clinical and Translational Medicine, 2020, 10, e152.	1.7	21
30	Common risk variants in NPHS1 and TNFSF15 are associated with childhood steroid-sensitive nephrotic syndrome. Kidney International, 2020, 98, 1308-1322.	2.6	39
31	Nonâ€random distribution of deleterious mutations in the DNA and proteinâ€binding domains of IRF6 are associated with Van Der Woude syndrome. Molecular Genetics & Genomic Medicine, 2020, 8, e1355.	0.6	13
32	Epigenome-wide association study for perceived discrimination among sub-Saharan African migrants in Europe - the RODAM study. Scientific Reports, 2020, 10, 4919.	1.6	7
33	HLA and autoantibodies define scleroderma subtypes and risk in African and European Americans and suggest a role for molecular mimicry. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 552-562.	3.3	52
34	Time-to-event modeling of hypertension reveals the nonexistence of true controls. ELife, 2020, 9, .	2.8	1
35	Epigenome-wide association study in whole blood on type 2 diabetes among sub-Saharan African individuals: findings from the RODAM study. International Journal of Epidemiology, 2019, 48, 58-70.	0.9	62
36	ZRANB3 is an African-specific type 2 diabetes locus associated with beta-cell mass and insulin response. Nature Communications, 2019, 10, 3195.	5.8	69

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37	Genetic Identification of Two Novel Loci Associated with Steroid-Sensitive Nephrotic Syndrome. Journal of the American Society of Nephrology: JASN, 2019, 30, 1375-1384.	3.0	40
38	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. Cell, 2019, 179, 984-1002.e36.	13.5	152
39	Type 2 diabetes complications and comorbidity in Sub-Saharan Africans. EClinicalMedicine, 2019, 16, 30-41.	3.2	58
40	Missense Pathogenic variants in KIF4A Affect Dental Morphogenesis Resulting in X-linked Taurodontism, Microdontia and Dens-Invaginatus. Frontiers in Genetics, 2019, 10, 800.	1.1	7
41	Genome-wide association study for proliferative diabetic retinopathy in Africans. Npj Genomic Medicine, 2019, 4, 20.	1.7	18
42	Genetic Basis of Obesity and Type 2 Diabetes in Africans: Impact on Precision Medicine. Current Diabetes Reports, 2019, 19, 105.	1.7	9
43	Studying the effects of diet on DNA methylation: challenges, pitfalls and a way forward. British Journal of Nutrition, 2019, 122, 717-718.	1.2	0
44	Tuberous sclerosis in a patient from Nigeria. American Journal of Medical Genetics, Part A, 2019, 179, 1423-1425.	0.7	4
45	Classical HLA alleles are associated with prevalent and persistent cervical high-risk HPV infection in African women. Human Immunology, 2019, 80, 723-730.	1.2	9
46	Genome-wide association study of type 2 diabetes in Africa. Diabetologia, 2019, 62, 1204-1211.	2.9	56
47	Multi-ancestry genome-wide geneâ€“smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. Nature Genetics, 2019, 51, 636-648.	9.4	112
48	Echocardiographic screening of 4107 Nigerian school children for rheumatic heart disease. Tropical Medicine and International Health, 2019, 24, 757-765.	1.0	11
49	Genomics of Cardiometabolic Disorders in Sub-Saharan Africa. , 2019, , 168-198.		0
50	Genomic analyses in African populations identify novel risk loci for cleft palate. Human Molecular Genetics, 2019, 28, 1038-1051.	1.4	61
51	HLA-DQA1 and APOL1 as Risk Loci for Childhood-Onset Steroid-Sensitive and Steroid-Resistant Nephrotic Syndrome. American Journal of Kidney Diseases, 2018, 71, 399-406.	2.1	41
52	Genetic Ancestry of Hadza and Sandawe Peoples Reveals Ancient Population Structure in Africa. Genome Biology and Evolution, 2018, 10, 875-882.	1.1	6
53	Clinical epidemiology of congenital heart disease in Nigerian children, 2012â€“2017. Birth Defects Research, 2018, 110, 1233-1240.	0.8	15
54	Identification of paternal uniparental disomy on chromosome 22 and a de novo deletion on chromosome 18 in individuals with orofacial clefts. Molecular Genetics & Genomic Medicine, 2018, 6, 924-932.	0.6	4

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55	Circulating MiR-374a-5p is a potential modulator of the inflammatory process in obesity. Scientific Reports, 2018, 8, 7680.	1.6	28
56	Genetic Variants Associated with Clinicopathological Profiles in Sporadic Breast Cancer in Sri Lankan Women. Journal of Breast Cancer, 2018, 21, 165.	0.8	16
57	Brief Report: Wholeâ€Exome Sequencing to Identify Rare Variants and Gene Networks That Increase Susceptibility to Scleroderma in African Americans. Arthritis and Rheumatology, 2018, 70, 1654-1660.	2.9	10
58	Medical genetics and genomic medicine in Nigeria. Molecular Genetics & Genomic Medicine, 2018, 6, 314-321.	0.6	13
59	Analyses of genome wide association data, cytokines, and gene expression in African-Americans with benign ethnic neutropenia. PLoS ONE, 2018, 13, e0194400.	1.1	36
60	Transferability of genome-wide associated loci for asthma in African Americans. Journal of Asthma, 2017, 54, 1-8.	0.9	11
61	Peripheral insulin resistance rather than beta cell dysfunction accounts for geographical differences in impaired fasting blood glucose among sub-Saharan African individuals: findings from the RODAM study. Diabetologia, 2017, 60, 854-864.	2.9	22
62	NFAT5 and SLC4A10 Loci Associate with Plasma Osmolality. Journal of the American Society of Nephrology: JASN, 2017, 28, 2311-2321.	3.0	24
63	22q11.2 deletion syndrome in diverse populations. American Journal of Medical Genetics, Part A, 2017, 173, 879-888.	0.7	103
64	Genomeâ€wide analysis identifies an africanâ€specific variant in SEMA4D associated with body mass index. Obesity, 2017, 25, 794-800.	1.5	30
65	Down syndrome in diverse populations. American Journal of Medical Genetics, Part A, 2017, 173, 42-53.	0.7	75
66	Assessing the spectrum of germline variation in Fanconi anemia genes among patients with head and neck carcinoma before age 50. Cancer, 2017, 123, 3943-3954.	2.0	37
67	The genomic landscape of African populations in health and disease. Human Molecular Genetics, 2017, 26, R225-R236.	1.4	64
68	An epigenome-wide association study in whole blood of measures of adiposity among Ghanaians: the RODAM study. Clinical Epigenetics, 2017, 9, 103.	1.8	55
69	Genomics of Cardiometabolic Disorders in Sub-Saharan Africa. Public Health Genomics, 2017, 20, 9-26.	0.6	17
70	Susceptibility to Cryptococcal Meningoencephalitis Associated With Idiopathic CD4+ Lymphopenia and Secondary Germline or Acquired Defects. Open Forum Infectious Diseases, 2017, 4, ofx082.	0.4	25
71	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. PLoS Genetics, 2017, 13, e1006719.	1.5	98
72	Single-trait and multi-trait genome-wide association analyses identify novel loci for blood pressure in African-ancestry populations. PLoS Genetics, 2017, 13, e1006728.	1.5	88

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73	Common and rare exonic MUC5B variants associated with type 2 diabetes in Han Chinese. PLoS ONE, 2017, 12, e0173784.	1.1	10
74	Impact of Type 2 Diabetes on Impaired Kidney Function in Sub-Saharan African Populations. Frontiers in Endocrinology, 2016, 7, 50.	1.5	9
75	Disparities in type 2 diabetes prevalence among ethnic minority groups resident in Europe: a systematic review and meta-analysis. Internal and Emergency Medicine, 2016, 11, 327-340.	1.0	171
76	Proinflammatory and lipid biomarkers mediate metabolically healthy obesity: A proteomics study. Obesity, 2016, 24, 1257-1265.	1.5	32
77	Obesity and type 2 diabetes in sub-Saharan Africans â€“ Is the burden in todayâ€™s Africa similar to African migrants in Europe? The RODAM study. BMC Medicine, 2016, 14, 166.	2.3	132
78	Estimation of F_{ST} and the Impact of de novo Mutation. Human Heredity, 2016, 82, 37-49.	0.4	1
79	Trans-ethnic Meta-analysis and Functional Annotation Illuminates theÂGenetic Architecture of Fasting Glucose and Insulin. American Journal of Human Genetics, 2016, 99, 56-75.	2.6	55
80	Towards a more representative morphology: clinical and ethical considerations for including diverse populations in diagnostic genetic atlases. Genetics in Medicine, 2016, 18, 1069-1074.	1.1	27
81	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. Nature Communications, 2016, 7, 10023.	5.8	412
82	An electronic atlas of human malformation syndromes in diverse populations. Genetics in Medicine, 2016, 18, 1085-1087.	1.1	44
83	Rare variants in tenascin genes in a cohort of children with primary vesicoureteric reflux. Pediatric Nephrology, 2016, 31, 247-253.	0.9	10
84	The Afro-Cardiac Study: Cardiovascular Disease Risk and Acculturation in West African Immigrants in the United States: Rationale and Study Design. Journal of Immigrant and Minority Health, 2016, 18, 1301-1308.	0.8	10
85	Evolutionary context for the association of Î³-globin, serum uric acid, and hypertension in African Americans. BMC Medical Genetics, 2015, 16, 103.	2.1	7
86	Phenotypic variance explained by local ancestry in admixed African Americans. Frontiers in Genetics, 2015, 6, 324.	1.1	13
87	Evaluation of Genome Wide Association Study Associated Type 2 Diabetes Susceptibility Loci in Sub Saharan Africans. Frontiers in Genetics, 2015, 6, 335.	1.1	50
88	An Improved Fst Estimator. PLoS ONE, 2015, 10, e0135368.	1.1	6
89	Prevalence of type 2 diabetes and its association with measures of body composition among African residents in the Netherlands â€“ The HELIUS study. Diabetes Research and Clinical Practice, 2015, 110, 137-146.	1.1	20
90	APOL1 G1 genotype modifies the association between HDLC and kidney function in African Americans. BMC Genomics, 2015, 16, 421.	1.2	9

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91	Genome-wide association study identifies African-ancestry specific variants for metabolic syndrome. Molecular Genetics and Metabolism, 2015, 116, 305-313.	0.5	41
92	The African Genome Variation Project shapes medical genetics in Africa. Nature, 2015, 517, 327-332.	13.7	473
93	Clinical and pharmacogenomic implications of genetic variation in a Southern Ethiopian population. Pharmacogenomics Journal, 2015, 15, 101-108.	0.9	15
94	HLA-DQA1 and PLCG2 Are Candidate Risk Loci for Childhood-Onset Steroid-Sensitive Nephrotic Syndrome. Journal of the American Society of Nephrology: JASN, 2015, 26, 1701-1710.	3.0	118
95	Novel genomic signals of recent selection in an Ethiopian population. European Journal of Human Genetics, 2015, 23, 1085-1092.	1.4	25
96	Global Gene Expression Profiling in Omental Adipose Tissue of Morbidly Obese Diabetic African Americans. Journal of Endocrinology and Metabolism, 2015, 5, 199-210.	0.1	21
97	Rare hereditary COL4A3/COL4A4 variants may be mistaken for familial focal segmental glomerulosclerosis. Kidney International, 2014, 86, 1253-1259.	2.6	195
98	Gene-Based Sequencing Identifies Lipid-Influencing Variants with Ethnicity-Specific Effects in African Americans. PLoS Genetics, 2014, 10, e1004190.	1.5	34
99	Meta-Analysis of Genome-Wide Association Studies in African Americans Provides Insights into the Genetic Architecture of Type 2 Diabetes. PLoS Genetics, 2014, 10, e1004517.	1.5	191
100	What does genomic medicine mean for diverse populations?. Molecular Genetics & Genomic Medicine, 2014, 2, 3-6.	0.6	17
101	Novel IRF6 mutations in families with Van Der Woude syndrome and popliteal pterygium syndrome from subâ€Saharan Africa. Molecular Genetics & Genomic Medicine, 2014, 2, 254-260.	0.6	24
102	Informed consent and ethical re-use of African genomic data. Human Genomics, 2014, 8, 18.	1.4	14
103	Mutations in the Gene That Encodes the F-Actin Binding Protein Anillin Cause FSGS. Journal of the American Society of Nephrology: JASN, 2014, 25, 1991-2002.	3.0	124
104	Evolution of the primate trypanolytic factor APOL1. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E2130-9.	3.3	183
105	High sensitivity C-reactive protein (Hs-CRP) remains highly stable in long-term archived human serum. Clinical Biochemistry, 2014, 47, 315-318.	0.8	66
106	Reconciling clinical importance and statistical significance. European Journal of Human Genetics, 2014, 22, 158-159.	1.4	1
107	Enabling the genomic revolution in Africa. Science, 2014, 344, 1346-1348.	6.0	361
108	Rare functional variants in genomeâ€“wide association identified candidate genes for nonsyndromic clefts in the African population. American Journal of Medical Genetics, Part A, 2014, 164, 2567-2571.	0.7	35

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109	Childhood Family Living Arrangements and Blood Pressure in Black Men. Hypertension, 2014, 63, 48-53.	1.3	9
110	Ethical and legal implications of whole genome and whole exome sequencing in African populations. BMC Medical Ethics, 2013, 14, 21.	1.0	52
111	Genetic Epidemiology of Type 2 Diabetes and Cardiovascular Diseases in Africa. Progress in Cardiovascular Diseases, 2013, 56, 251-260.	1.6	28
112	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. Nature Genetics, 2013, 45, 690-696.	9.4	232
113	Genome-Wide Association of Body Fat Distribution in African Ancestry Populations Suggests New Loci. PLoS Genetics, 2013, 9, e1003681.	1.5	109
114	TNXB Mutations Can Cause Vesicoureteral Reflux. Journal of the American Society of Nephrology: JASN, 2013, 24, 1313-1322.	3.0	60
115	RPS19 and TYMS SNPs and Prevalent High Risk Human Papilloma Virus Infection in Nigerian Women. PLoS ONE, 2013, 8, e66930.	1.1	17
116	Variation inAPOL1Contributes to Ancestry-Level Differences in HDLc-Kidney Function Association. International Journal of Nephrology, 2012, 2012, 1-10.	0.7	28
117	C-reactive protein (CRP) promoter polymorphisms influence circulating CRP levels in a genome-wide association study of African Americans. Human Molecular Genetics, 2012, 21, 3063-3072.	1.4	32
118	A founder mutation in LEPRE1 carried by 1.5% of West Africans and 0.4% of African Americans causes lethal recessive osteogenesis imperfecta. Genetics in Medicine, 2012, 14, 543-551.	1.1	49
119	HLA Class II Locus and Susceptibility to Podoconiosis. New England Journal of Medicine, 2012, 366, 1200-1208.	13.9	125
120	UGT1A1 is a major locus influencing bilirubin levels in African Americans. European Journal of Human Genetics, 2012, 20, 463-468.	1.4	63
121	Simple F Test Reveals Gene-Gene Interactions in Case-Control Studies. Bioinformatics and Biology Insights, 2012, 6, BBI.S9867.	1.0	1
122	Genome-wide association study identifies novel loci association with fasting insulin and insulin resistance in African Americans. Human Molecular Genetics, 2012, 21, 4530-4536.	1.4	80
123	Transferability and Fine Mapping of genome-wide associated loci for lipids in African Americans. BMC Medical Genetics, 2012, 13, 88.	2.1	33
124	A Genome-Wide Association Search for Type 2 Diabetes Genes in African Americans. PLoS ONE, 2012, 7, e29202.	1.1	197
125	Adrenergic Alpha-1 Pathway Is Associated with Hypertension among Nigerians in a Pathway-focused Analysis. PLoS ONE, 2012, 7, e37145.	1.1	20
126	Prediction of HLA Class II Alleles Using SNPs in an African Population. PLoS ONE, 2012, 7, e40206.	1.1	10

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127	Using a â€œgenomics toolâ€•to develop disease prevention strategy in a low-income setting: lessons from the podoconiosis research project. Journal of Community Genetics, 2012, 3, 303-309.	0.5	13
128	Genome-wide associated loci influencing interleukin (IL)-10, IL-1Ra, and IL-6 levels in African Americans. Immunogenetics, 2012, 64, 351-359.	1.2	31
129	Multiple Loci Associated with Renal Function in African Americans. PLoS ONE, 2012, 7, e45112.	1.1	7
130	Paradoxical Hyperadiponectinemia is Associated With the Metabolically Healthy Obese (MHO) Phenotype in African Americans. Journal of Endocrinology and Metabolism, 2012, 2, 51-65.	0.1	61
131	Mapping of disease-associated variants in admixed populations. Genome Biology, 2011, 12, 223.	13.9	53
132	The Roles of IL-6, IL-10, and IL-1RA in Obesity and Insulin Resistance in African-Americans. Journal of Clinical Endocrinology and Metabolism, 2011, 96, E2018-E2022.	1.8	59
133	A genome-wide association study of serum uric acid in African Americans. BMC Medical Genomics, 2011, 4, 17.	0.7	82
134	Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. Nature, 2011, 478, 103-109.	13.7	1,855
135	Resequencing and Analysis of Variation in the TCF7L2 Gene in African Americans Suggests That SNP rs7903146 Is the Causal Diabetes Susceptibility Variant. Diabetes, 2011, 60, 662-668.	0.3	74
136	Association of ATP1B1, RGS5 and SELE polymorphisms with hypertension and blood pressure in Africanâ€“Americans. Journal of Hypertension, 2011, 29, 1906-1912.	0.3	28
137	Genome-wide Comparison of African-Ancestry Populations from CARe and Other Cohorts Reveals Signals of Natural Selection. American Journal of Human Genetics, 2011, 89, 368-381.	2.6	79
138	Genome-wide association study for serum urate concentrations and gout among African Americans identifies genomic risk loci and a novel URAT1 loss-of-function allele. Human Molecular Genetics, 2011, 20, 4056-4068.	1.4	101
139	Combined admixture mapping and association analysis identifies a novel blood pressure genetic locus on 5p13: contributions from the CARe consortium. Human Molecular Genetics, 2011, 20, 2285-2295.	1.4	77
140	Association of genetic variation with systolic and diastolic blood pressure among African Americans: the Candidate Gene Association Resource study. Human Molecular Genetics, 2011, 20, 2273-2284.	1.4	168
141	Joint Ancestry and Association Testing in Admixed Individuals. PLoS Computational Biology, 2011, 7, e1002325.	1.5	88
142	Genetic Association for Renal Traits among Participants of African Ancestry Reveals New Loci for Renal Function. PLoS Genetics, 2011, 7, e1002264.	1.5	109
143	Identification, Replication, and Fine-Mapping of Loci Associated with Adult Height in Individuals of African Ancestry. PLoS Genetics, 2011, 7, e1002298.	1.5	93
144	Genetic Susceptibility to Acute Rheumatic Fever: A Systematic Review and Meta-Analysis of Twin Studies. PLoS ONE, 2011, 6, e25326.	1.1	102

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145	Practical considerations for imputation of untyped markers in admixed populations. Genetic Epidemiology, 2010, 34, 258-265.	0.6	32
146	Genetic variants on chromosome 5p12 are associated with risk of breast cancer in African American women: the Black Womenâ€™s Health Study. Breast Cancer Research and Treatment, 2010, 123, 525-530.	1.1	25
147	Development of admixture mapping panels for African Americans from commercial high-density SNP arrays. BMC Genomics, 2010, 11, 417.	1.2	15
148	Genome-wide association of anthropometric traits in African- and African-derived populations. Human Molecular Genetics, 2010, 19, 2725-2738.	1.4	90
149	FTO Genetic Variation and Association With Obesity in West Africans and African Americans. Diabetes, 2010, 59, 1549-1554.	0.3	94
150	Circulating Adiponectin Is Associated with Obesity and Serum Lipids in West Africans. Journal of Clinical Endocrinology and Metabolism, 2010, 95, 3517-3521.	1.8	37
151	Genetic Variation at Selected SNPs in the Leptin Gene and Association of Alleles with Markers of Kidney Disease in a Xhosa Population of South Africa. PLoS ONE, 2010, 5, e9086.	1.1	14
152	Tailoring Consent to Context: Designing an Appropriate Consent Process for a Biomedical Study in a Low Income Setting. PLoS Neglected Tropical Diseases, 2009, 3, e482.	1.3	85
153	Energy expenditure does not predict weight change in either Nigerian or African American women. American Journal of Clinical Nutrition, 2009, 89, 169-176.	2.2	50
154	Association of Regions on Chromosomes 6 and 7 With Blood Pressure in Nigerian Families. Circulation: Cardiovascular Genetics, 2009, 2, 38-45.	5.1	17
155	A Genome-Wide Association Study of Hypertension and Blood Pressure in African Americans. PLoS Genetics, 2009, 5, e1000564.	1.5	348
156	Transferability and Fine-Mapping of Genome-Wide Associated Loci for Adult Height across Human Populations. PLoS ONE, 2009, 4, e8398.	1.1	47
157	Beta thalassaemia trait in western Nigeria. African Health Sciences, 2009, 9, 46-8.	0.3	12
158	Genetic studies of African populations: an overview on disease susceptibility and response to vaccines and therapeutics. Human Genetics, 2008, 123, 557-598.	1.8	79
159	Epidemiology, Heritability, and Genetic Linkage of C-Reactive Protein in African Americans (from the) Tj ETQq1 1 0.784314 rgBT /Over	0.7	48
160	Meta-Analysis of Genome-Wide Linkage Studies of Quantitative Lipid Traits in Families Ascertained for Type 2 Diabetes. Diabetes, 2007, 56, 890-896.	0.3	33
161	Genome-wide search for susceptibility genes to type 2 diabetes in West Africans: Potential role of C-peptide. Diabetes Research and Clinical Practice, 2007, 78, e1-e6.	1.1	20
162	Refining the impact of TCF7L2 gene variants on type 2 diabetes and adaptive evolution. Nature Genetics, 2007, 39, 218-225.	9.4	485

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163	A variant in CDKAL1 influences insulin response and risk of type 2 diabetes. Nature Genetics, 2007, 39, 770-775.	9.4	966
164	Two variants on chromosome 17 confer prostate cancer risk, and the one in TCF2 protects against type 2 diabetes. Nature Genetics, 2007, 39, 977-983.	9.4	670
165	A Genomeâ€wide Scan of Loci Linked to Serum Adiponectin in Two Populations of African Descent. Obesity, 2007, 15, 1207-1214.	1.5	11
166	A Genome-Wide Search for Linkage to Renal Function Phenotypes in West Africans With Type 2 Diabetes. American Journal of Kidney Diseases, 2007, 49, 394-400.	2.1	48
167	Positive association between resting energy expenditure and weight gain in a lean adult population. American Journal of Clinical Nutrition, 2006, 83, 1076-1081.	2.2	50
168	Computational disease gene identification: a concert of methods prioritizes type 2 diabetes and obesity candidate genes. Nucleic Acids Research, 2006, 34, 3067-3081.	6.5	134
169	Genomewide Scan and Fine Mapping of Quantitative Trait Loci for Intraocular Pressure on 5q and 14q in West Africans. , 2006, 47, 3262.		48
170	Genetic effects on blood pressure localized to chromosomes 6 and 7. Journal of Hypertension, 2005, 23, 1367-1373.	0.3	14
171	Angiotensinâ€Converting Enzyme Gene Polymorphisms and Obesity: An Examination of Three Black Populations. Obesity, 2005, 13, 823-828.	4.0	26
172	Genetic structure in four West African population groups. BMC Genetics, 2005, 6, 38.	2.7	36
173	Concurrent bacteraemia and malaria in febrile Nigerian infants. Tropical Doctor, 2005, 35, 34-36.	0.2	18
174	Haplotypes produced from rare variants in the promoter and coding regions of angiotensinogen contribute to variation in angiotensinogen levels. Human Molecular Genetics, 2005, 14, 639-643.	1.4	29
175	A genome wide quantitative trait linkage analysis for serum lipids in type 2 diabetes in an African population. Atherosclerosis, 2005, 181, 389-397.	0.4	35
176	Calpain-10 gene polymorphisms and type 2 diabetes in West Africans: the Africa America Diabetes Mellitus (AADM) Study. Annals of Epidemiology, 2005, 15, 153-159.	0.9	33
177	Association Between Blood Pressure and Resting Energy Expenditure Independent of Body Size. Hypertension, 2004, 43, 555-560.	1.3	50
178	A Genome-Wide Search for Type 2 Diabetes Susceptibility Genes in West Africans: The Africa America Diabetes Mellitus (AADM) Study. Diabetes, 2004, 53, 838-841.	0.3	88
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