Michele Ramsay

List of Publications by Year
in descending order

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181 papers	7,017 citations	87888 38 h-index	76900 74 g-index
227 all docs	227 docs citations	227 times ranked	9954 citing authors

#	Article	IF	Citations
1	Exome-based mutation screening in South African children with primary congenital glaucoma. Eye, 2023, 37, 362-368.	2.1	2
2	African genomic data sharing and the struggle for equitable benefit. Patterns, 2022, 3, 100412.	5.9	17
3	Short- and long-read metagenomics of urban and rural South African gut microbiomes reveal a transitional composition and undescribed taxa. Nature Communications, 2022, 13, 926.	12.8	26
4	Genetic associations with carotid intima-media thickness link to atherosclerosis with sex-specific effects in sub-Saharan Africans. Nature Communications, 2022, 13, 855.	12.8	10
5	Perspectives on returning individual and aggregate genomic research results to study participants and communities in Kenya: a qualitative study. BMC Medical Ethics, 2022, 23, 27.	2.4	3
6	Carotid Atherosclerosis, Microalbuminuria, and Estimated 10-Year Atherosclerotic Cardiovascular Disease Risk in Sub-Saharan Africa. JAMA Network Open, 2022, 5, e227559.	5.9	8
7	Polygenic risk scores for CARDINAL study. Nature Genetics, 2022, 54, 527-530.	21.4	5
8	Meta-analysis of sub-Saharan African studies provides insights into genetic architecture of lipid traits. Nature Communications, 2022, 13, 2578.	12.8	18
9	Apolipoprotein L1 High-Risk Genotypes and Albuminuria in Sub-Saharan African Populations. Clinical Journal of the American Society of Nephrology: CJASN, 2022, 17, 798-808.	4.5	8
10	Genome-wide meta-analysis identifies 127 open-angle glaucoma loci with consistent effect across ancestries. Nature Communications, 2021, 12, 1258.	12.8	196
11	Poor cardiovascular health is associated with subclinical atherosclerosis in apparently healthy sub-Saharan African populations: an H3Africa AWI-Gen study. BMC Medicine, 2021, 19, 30.	5.5	13
12	Admixture/fine-mapping in Brazilians reveals a West African associated potential regulatory variant (rs114066381) with a strong female-specific effect on body mass and fat mass indexes. International Journal of Obesity, 2021, 45, 1017-1029.	3.4	4
13	Association of Rare CYP39A1 Variants With Exfoliation Syndrome Involving the Anterior Chamber of the Eye. JAMA - Journal of the American Medical Association, 2021, 325, 753.	7.4	16
14	Marianne Alberts (1928â€“2020): Caring, compassionate and humble biochemist and pioneer. South African Journal of Science, 2021, 117, .	0.7	1
15	Adiposity Phenotypes and Subclinical Atherosclerosis in Adults from Subâ€“Saharan Africa: An H3Africa AWIâ€“Gen Study. Global Heart, 2021, 16, 19.	2.3	2
16	Community engagement and feedback of results in the H3Africa AWI-Gen project: Experiences from the Navrongo Demographic and Health Surveillance site in Northern Ghana. AAS Open Research, 2021, 4, 15.	1.5	5
17	Potential Impact of DPYD Variation on Fluoropyrimidine Drug Response in sub-Saharan African Populations. Frontiers in Genetics, 2021, 12, 626954.	2.3	10
18	Genetic substructure and complex demographic history of South African Bantu speakers. Nature Communications, 2021, 12, 2080.	12.8	47

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19	POPIA Code of Conduct for Research (with corrigendum). South African Journal of Science, 2021, 117, .	0.7	8
20	Drafting a Code of Conduct for Research under the Protection of Personal Information Act No. 4 of 2013 (with corrigendum). South African Journal of Science, 2021, 117, .	0.7	4
21	Prevalence and socio-demographic correlates of tobacco and alcohol use in four sub-Saharan African countries: a cross-sectional study of middle-aged adults. BMC Public Health, 2021, 21, 1126.	2.9	16
22	Has translational genomics come of age in Africa?. Human Molecular Genetics, 2021, 30, R164-R173.	2.9	11
23	Interleukin-8 genetic diversity, haplotype structure and production differ in two ethnically distinct South African populations. Cytokine, 2021, 143, 155489.	3.2	0
24	G6PD distribution in sub-Saharan Africa and potential risks of using chloroquine/hydroxychloroquine based treatments for COVID-19. Pharmacogenomics Journal, 2021, 21, 649-656.	2.0	11
25	Associations Between CYP17A1 and SERPINA6/A1 Polymorphisms, and Cardiometabolic Risk Factors in Black South Africans. Frontiers in Genetics, 2021, 12, 687335.	2.3	2
26	UGT1A1 regulatory variant with potential effect on efficacy of HIV and cancer drugs commonly prescribed in South Africa. Pharmacogenomics, 2021, 22, 963-972.	1.3	0
27	Estimating the burden of cardiovascular risk in community dwellers over 40 years old in South Africa, Kenya, Burkina Faso and Ghana. BMJ Global Health, 2021, 6, e003499.	4.7	9
28	African genetic diversity and adaptation inform a precision medicine agenda. Nature Reviews Genetics, 2021, 22, 284-306.	16.3	69
29	The Extent and Impact of Variation in ADME Genes in Sub-Saharan African Populations. Frontiers in Pharmacology, 2021, 12, 634016.	3.5	19
30	Genetic Susceptibility to Breast Cancer in Sub-Saharan African Populations. JCO Global Oncology, 2021, 7, 1462-1471.	1.8	3
31	Apolipoprotein E Genetic Variation and Its Association With Cognitive Function in Rural-Dwelling Older South Africans. Frontiers in Genetics, 2021, 12, 689756.	2.3	0
32	Bantu-speaker migration and admixture in southern Africa. Human Molecular Genetics, 2021, 30, R56-R63.	2.9	21
33	The Use of â€˜Omics for Diagnosing and Predicting Progression of Chronic Kidney Disease: A Scoping Review. Frontiers in Genetics, 2021, 12, 682929.	2.3	14
34	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	27.8	353
35	nf-rnaSeqCount: A Nextflow pipeline for obtaining raw read counts from RNA-seq data. South African Computer Journal, 2021, 33, .	0.2	0
36	Building a Platform to Enable NCD Research to Address Population Health in Africa: CVD Working Group Discussion at the Sixth H3Africa Consortium Meeting in Zambia. Global Heart, 2020, 11, 165.	2.3	11

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37	Autozygosity influences cardiometabolic disease-associated traits in the AWI-Gen sub-Saharan African study. Nature Communications, 2020, 11, 5754.	12.8	23
38	Candidate Gene Analysis Reveals Strong Association of CETP Variants With High Density Lipoprotein Cholesterol and PCSK9 Variants With Low Density Lipoprotein Cholesterol in Ghanaian Adults: An AWI-Gen Sub-Study. Frontiers in Genetics, 2020, 11, 456661.	2.3	4
39	High-depth African genomes inform human migration and health. Nature, 2020, 586, 741-748.	27.8	197
40	Blood Pressure Indices and Associated Risk Factors in a Rural West African Adult Population: Insights from an AWI-Gen Substudy in Ghana. International Journal of Hypertension, 2020, 2020, 1-11.	1.3	5
41	Regional Patterns and Association Between Obesity and Hypertension in Africa. Hypertension, 2020, 75, 1167-1178.	2.7	49
42	Genetic associations between serum low LDL-cholesterol levels and variants in LDLR, APOB, PCSK9 andÂLDLRAP1 in African populations. PLoS ONE, 2020, 15, e0229098.	2.5	9
43	Developing a Road Map to Spread Genomic Knowledge in Africa: 10th Conference of the African Society of Human Genetics, Cairo, Egypt. American Journal of Tropical Medicine and Hygiene, 2020, 102, 719-723.	1.4	6
44	Building Skills and Resources for Genomics, Epigenetics, and Bioinformatics Research for Africa: Report of the Joint 11th Conference of the African Society of Human Genetics and 12th H3Africa Consortium, 2018. American Journal of Tropical Medicine and Hygiene, 2020, 102, 1417-1424.	1.4	7
45	Advancing non-communicable diseases research in Ghana: key stakeholdersâ€™ recommendations from a symposium. Ghana Medical Journal, 2020, 54, 121-125.	0.4	3
46	Runs of homozygosity in sub-Saharan African populations provide insights into complex demographic histories. Human Genetics, 2019, 138, 1123-1142.	3.8	20
47	Classical Cardiovascular Risk Factors and HIV are Associated With Carotid Intimaâ€Media Thickness in Adults From Subâ€Saharan Africa: Findings From H3Africa AWIâ€Gen Study. Journal of the American Heart Association, 2019, 8, e011506.	3.7	20
48	Kidney damage and associated risk factors in rural and urban sub-Saharan Africa (AWI-Gen): a cross-sectional population study. The Lancet Global Health, 2019, 7, e1632-e1643.	6.3	56
49	Association of Genetic Variants With Primary Open-Angle Glaucoma Among Individuals With African Ancestry. JAMA - Journal of the American Medical Association, 2019, 322, 1682.	7.4	50
50	Epigenetic modification of the pentose phosphate pathway and the IGF-axis in women with gestational diabetes mellitus. Epigenomics, 2019, 11, 1371-1385.	2.1	12
51	The protective variant rs7173049 at LOXL1 locus impacts on retinoic acid signaling pathway in pseudoexfoliation syndrome. Human Molecular Genetics, 2019, 28, 2531-2548.	2.9	22
52	Sydney Brenner (1927â€“2019): The opening game. South African Journal of Science, 2019, 115, .	0.7	0
53	An optimistic vision for biosciences in South Africa: Reply to Thaldar et al. (2019). South African Journal of Science, 2019, 115, .	0.7	0
54	Leveraging genomic diversity to promote human and animal health. Communications Biology, 2019, 2, 463.	4.4	0

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55	The podocin V260E mutation predicts steroid resistant nephrotic syndrome in black South African children with focal segmental glomerulosclerosis. Communications Biology, 2019, 2, 416.	4.4	3
56	Keratolytic Winter Erythema: An Update. Dermatopathology (Basel, Switzerland), 2019, 6, 126-132.	1.5	6
57	Targeted ultra-deep sequencing of a South African Bantu-speaking cohort to comprehensively map and characterize common and novel variants in 65 pharmacologically-related genes. Pharmacogenetics and Genomics, 2019, 29, 167-178.	1.5	5
58	HLA-DRB1 Amino Acid Positions and Residues Associated with Antibody-positive Rheumatoid Arthritis in Black South Africans. Journal of Rheumatology, 2019, 46, 138-144.	2.0	6
59	Data Resource Profile: Cardiovascular H3Africa Innovation Resource (CHAIR). International Journal of Epidemiology, 2019, 48, 366-367g.	1.9	19
60	Dysregulation of the Wnt signaling pathway in South African patients with diffuse systemic sclerosis. Clinical Rheumatology, 2019, 38, 933-938.	2.2	14
61	Novel and Known Gene-Smoking Interactions With cIMT Identified as Potential Drivers for Atherosclerosis Risk in West-African Populations of the AWI-Gen Study. Frontiers in Genetics, 2019, 10, 1354.	2.3	10
62	Ethical and practical issues to consider in the governance of genomic and human research data and data sharing in South Africa: a meeting report. AAS Open Research, 2019, 2, 15.	1.5	5
63	Integrating environmental health and genomics research in Africa: challenges and opportunities identified during a Human Heredity and Health in Africa (H3Africa) Consortium workshop. AAS Open Research, 2019, 2, 159.	1.5	3
64	Population specific genetic heterogeneity of familial hypercholesterolemia in South Africa. Current Opinion in Lipidology, 2018, 29, 72-79.	2.7	13
65	Runs of homozygosity: windows into population history and trait architecture. Nature Reviews Genetics, 2018, 19, 220-234.	16.3	497
66	Different adiposity indices and their association with blood pressure and hypertension in middle-aged urban black South African men and women: findings from the AWI-GEN South African Soweto Site. BMC Public Health, 2018, 18, 524.	2.9	10
67	Insights into the genetics of blood pressure in black South African individuals: the Birth to Twenty cohort. BMC Medical Genomics, 2018, 11, 2.	1.5	11
68	ASSAf consensus study on the ethical, legal and social implications of genetics and genomics in South Africa. South African Journal of Science, 2018, 114, .	0.7	6
69	Regional and sex-specific variation in BMI distribution in four sub-Saharan African countries: The H3Africa AWI-Gen study. Global Health Action, 2018, 11, 1556561.	1.9	37
70	The burden of dyslipidaemia and factors associated with lipid levels among adults in rural northern Ghana: An AWI-Gen sub-study. PLoS ONE, 2018, 13, e0206326.	2.5	33
71	An upward trajectory of genomic publications from Africa: cautious optimism for a turning tide. Global Health, Epidemiology and Genomics, 2018, 3, e17.	0.8	3
72	Patterns of adult body mass in sub-Saharan Africa. Global Health Action, 2018, 11, 1556497.	1.9	0

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73	Genomic and environmental risk factors for cardiometabolic diseases in Africa: methods used for Phase 1 of the AWI-Gen population cross-sectional study. Global Health Action, 2018, 11, 1507133.	1.9	82
74	Genetic variants in SEC16B are associated with body composition in black South Africans. Nutrition and Diabetes, 2018, 8, 43.	3.2	24
75	The Next Generation Scientist program: capacity-building for future scientific leaders in low- and middle-income countries. BMC Medical Education, 2018, 18, 233.	2.4	17
76	Gender differences in sociodemographic and behavioural factors associated with BMI in an adult population in rural Burkina Faso â€“ an AWI-Gen sub-study. Global Health Action, 2018, 11, 1527557.	1.9	8
77	H3Africa: current perspectives. Pharmacogenomics and Personalized Medicine, 2018, Volume 11, 59-66.	0.7	101
78	Common and Founder Mutations for Monogenic Traits in Sub-Saharan African Populations. Annual Review of Genomics and Human Genetics, 2018, 19, 149-175.	6.2	9
79	Sociodemographic and behavioural factors associated with body mass index among men and women in Nairobi slums: AWI-Gen Project. Global Health Action, 2018, 11, 1470738.	1.9	19
80	Socio-demographic and behavioural determinants of body mass index among an adult population in rural Northern Ghana: the AWI-Gen study. Global Health Action, 2018, 11, 1467588.	1.9	23
81	Demographic, socio-economic and behavioural correlates of BMI in middle-aged black men and women from urban Johannesburg, South Africa. Global Health Action, 2018, 11, 1448250.	1.9	26
82	African genetic diversity provides novel insights into evolutionary history and local adaptations. Human Molecular Genetics, 2018, 27, R209-R218.	2.9	38
83	Assessing runs of Homozygosity: a comparison of SNP Array and whole genome sequence low coverage data. BMC Genomics, 2018, 19, 106.	2.8	93
84	Familial congenital cataract, coloboma, and nystagmus phenotype with variable expression caused by mutation in in a South African family. Molecular Vision, 2018, 24, 407-413.	1.1	12
85	Duplicated Enhancer Region Increases Expression of CTSB and Segregates with Keratolytic Winter Erythema in South African and Norwegian Families. American Journal of Human Genetics, 2017, 100, 737-750.	6.2	35
86	Genetic association study of exfoliation syndrome identifies a protective rare variant at LOXL1 and five new susceptibility loci. Nature Genetics, 2017, 49, 993-1004.	21.4	114
87	Whole-genome sequencing for an enhanced understanding of genetic variation among South Africans. Nature Communications, 2017, 8, 2062.	12.8	88
88	Establishing an academic biobank in a resource-challenged environment. South African Medical Journal, 2017, 107, 486.	0.6	15
89	Regional and Sex Differences in the Prevalence and Awareness of Hypertension: An H3Africa AWI-Gen Study Across 6 Sites in Sub-Saharan Africa. Global Heart, 2017, 12, 81.	2.3	105
90	Implications of direct-to-consumer whole-exome sequencing in South Africa. South African Medical Journal, 2016, 106, 139.	0.6	3

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91	Population Stratification and Underrepresentation of Indian Subcontinent Genetic Diversity in the 1000 Genomes Project Dataset. Genome Biology and Evolution, 2016, 8, 3460-3470.	2.5	33
92	Novel mutation in the CHST6 gene causes macular corneal dystrophy in a black South African family. BMC Medical Genetics, 2016, 17, 47.	2.1	11
93	African partnerships through the H3Africa Consortium bring a genomic dimension to longitudinal population studies on the continent. International Journal of Epidemiology, 2016, 45, 305-308.	1.9	32
94	Tele-ophthalmology: Opportunities for improving diabetes eye care in resource- and specialist-limited Sub-Saharan African countries. Journal of Telemedicine and Telecare, 2016, 22, 311-316.	2.7	38
95	The Genetics of POAG in Black South Africans: A Candidate Gene Association Study. Scientific Reports, 2015, 5, 8378.	3.3	33
96	Growing genomic research on the African continent: The H3Africa Consortium. South African Medical Journal, 2015, 105, 1016.	0.6	14
97	MYOCMutations in Black South African Patients with Primary Open-angle Glaucoma: Genetic Testing and Cascade Screening. Ophthalmic Genetics, 2015, 36, 31-38.	1.2	8
98	The H3Africa policy framework: negotiating fairness in genomics. Trends in Genetics, 2015, 31, 117-119.	6.7	65
99	Reduced DNA methylation at the PEG3 DMR and KvDMR1 loci in children exposed to alcohol in utero: a South African Fetal Alcohol Syndrome cohort study. Frontiers in Genetics, 2015, 6, 85.	2.3	36
100	APOL1 Risk Variants Are Strongly Associated with HIV-Associated Nephropathy in Black South Africans. Journal of the American Society of Nephrology: JASN, 2015, 26, 2882-2890.	6.1	256
101	Epigenetic epidemiology: is there cause for optimism?. Epigenomics, 2015, 7, 683-685.	2.1	4
102	Epigenetics and the burden of noncommunicable disease: a paucity of research in Africa. Epigenomics, 2015, 7, 627-639.	2.1	14
103	Genetic variants and cellular stressors associated with exfoliation syndrome modulate promoter activity of a lncRNA within theLOXL1locus. Human Molecular Genetics, 2015, 24, 6552-6563.	2.9	76
104	The African Genome Variation Project shapes medical genetics in Africa. Nature, 2015, 517, 327-332.	27.8	473
105	African origins and chronic kidney disease susceptibility in the human immunodeficiency virus era. World Journal of Nephrology, 2015, 4, 295.	2.0	25
106	Immunochip Identifies Novel, and Replicates Known, Genetic Risk Loci for Rheumatoid Arthritis in Black South Africans. Molecular Medicine, 2014, 20, 341-349.	4.4	21
107	Enabling the genomic revolution in Africa. Science, 2014, 344, 1346-1348.	12.6	361
108	Population-specific common SNPs reflect demographic histories and highlight regions of genomic plasticity with functional relevance. BMC Genomics, 2014, 15, 437.	2.8	40

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109	Ethical issues in genomic research on the African continent: experiences and challenges to ethics review committees. Human Genomics, 2014, 8, 15.	2.9	50
110	Understanding the rise in cardiovascular diseases in Africa : harmonising H3Africa genomic epidemiological teams and tools : cardiovascular topic. Cardiovascular Journal of Africa, 2014, 25, 134-136.	0.4	46
111	Genetic factors influencing bone mineral content in a black South African population. Journal of Bone and Mineral Metabolism, 2013, 31, 708-716.	2.7	7
112	Genetic diversity in black South Africans from Soweto. BMC Genomics, 2013, 14, 644.	2.8	49
113	The elusive gene for keratolytic winter erythema. South African Medical Journal, 2013, 103, 961.	0.6	6
114	Direct-to-consumer genetic testing: To test or not to test, that is the question. South African Medical Journal, 2013, 103, 510.	0.6	11
115	Professor Trefor Jenkins: A tribute. South African Medical Journal, 2013, 103, 956.	0.6	0
116	Biology and genetics of oculocutaneous albinism and vitiligo â€“ common pigmentation disorders in southern Africa. South African Medical Journal, 2013, 103, 984.	0.6	21
117	In utero alcohol exposure, epigenetic changes, and their consequences. , 2013, 35, 37-46.		47
118	Appetite regulation genes are associated with body mass index in black South African adolescents: a genetic association study. BMJ Open, 2012, 2, e000873.	1.9	28
119	Differential gene expression of MMP-1, TIMP-1 and HGF in clinically involved and uninvolved skin in South Africans with SSc. Rheumatology, 2012, 51, 1049-1052.	1.9	31
120	Africa: Continent of genome contrasts with implications for biomedical research and health. FEBS Letters, 2012, 586, 2813-2819.	2.8	31
121	Exclusion of CTSB and FDFT1 as positional and functional candidate genes for keratolytic winter erythema (KWE). Journal of Dermatological Science, 2012, 65, 58-62.	1.9	6
122	The Effect of Preconception Paternal Alcohol Exposure on Epigenetic Remodeling of the H19 and Rasgrf1 Imprinting Control Regions in Mouse Offspring. Frontiers in Genetics, 2012, 3, 10.	2.3	97
123	No Evidence for a Parent-of-Origin Specific Differentially Methylated Region Linked to RASGRF1. Frontiers in Genetics, 2012, 3, 41.	2.3	6
124	Absence of GJB2 gene mutations, the GJB6 deletion (GJB6-D13S1830) and four common mitochondrial mutations in nonsyndromic genetic hearing loss in a South African population. International Journal of Pediatric Otorhinolaryngology, 2011, 75, 611-617.	1.0	48
125	A computational approach to candidate gene prioritization for X-linked mental retardation using annotation-based binary filtering and motif-based linear discriminatory analysis. Biology Direct, 2011, 6, 30.	4.6	11
126	Africa: the next frontier for human disease gene discovery?. Human Molecular Genetics, 2011, 20, R214-R220.	2.9	25

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127	Genetic variants inCYP(-1A2,-2C9,-2C19,-3A4and-3A5),VKORC1andABCB1genes in a black South African population: a window into diversity. Pharmacogenomics, 2011, 12, 1663-1670.		60
128	Myocilin mutations in black South Africans with POAG. Molecular Vision, 2011, 17, 1064-9.	1.1	16
129	The PTPN22 R620W polymorphism is not associated with systemic rheumatic diseases in South Africans. Rheumatology, 2010, 49, 820-821.	1.9	4
130	Genetic and epigenetic insights into fetal alcohol spectrum disorders. Genome Medicine, 2010, 2, 27.	8.2	82
131	Major LOXL1 risk allele is reversed in exfoliation glaucoma in a black South African population. Molecular Vision, 2010, 16, 705-12.	1.1	86
132	Exposure of Mouse Embryos to Ethanol During Preimplantation Development: Effect on DNA Methylation in the H19 Imprinting Control Region1. Biology of Reproduction, 2009, 81, 618-627.	2.7	140
133	Effect of Alcohol Consumption on CpG Methylation in the Differentially Methylated Regions of H19 and IGâ€DMR in Male Gametesâ€"Implications for Fetal Alcohol Spectrum Disorders. Alcoholism: Clinical and Experimental Research, 2009, 33, 1615-1627.	2.4	224
134	Glucocerebrosidase gene mutations in black South Africans with Gaucher disease. Blood Cells, Molecules, and Diseases, 2009, 43, 129-133.	1.4	6
135	Spectrum of genetic variation at the ABCC6 locus in South Africans: Pseudoxanthoma elasticum patients and healthy individuals. Journal of Dermatological Science, 2009, 54, 198-204.	1.9	10
136	Interleukin-1 receptor antagonist gene polymorphisms are associated with disease severity in Black South Africans with rheumatoid arthritis. Joint Bone Spine, 2008, 75, 422-425.	1.6	13
137	WGA Allows the Molecular Characterization of a Novel Large CFTR Rearrangement in a Black South African Cystic Fibrosis Patient. Journal of Molecular Diagnostics, 2008, 10, 544-548.	2.8	8
138	Mutation detection in the ABCC6 gene and genotype phenotype analysis in a large international case series affected by pseudoxanthoma elasticum. Journal of Medical Genetics, 2007, 44, 621-628.	3.2	161
139	Computational selection and prioritization of candidate genes for Fetal Alcohol Syndrome. BMC Genomics, 2007, 8, 389.	2.8	36
140	Complete Androgen Insensitivity Syndrome in A Black South African Family: A Clinical and Molecular Investigation. Endocrine Practice, 2006, 12, 664-669.	2.1	3
141	Diagnosing cystic fibrosis in South Africa. South African Medical Journal, 2006, 96, 304, 306.	0.6	4
142	DNA Polymorphism and Selection at the Melanocortinâ€1 Receptor Gene in Normally Pigmented Southern African Individuals. Annals of the New York Academy of Sciences, 2003, 994, 299-306.	3.8	72
143	Physical and transcriptional map of the critical region for keratolytic winter erythema (KWE) on chromosome 8p22-p23 between D8S550 and D8S1759. European Journal of Human Genetics, 2002, 10, 17-25.	2.8	14
144	Four novel variants in MC1R in red-haired South African individuals of European descent: S83P, Y152X, A171D, P256S. Human Mutation, 2002, 19, 461-462.	2.5	28

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145	Larger Genetic Differences Within Africans Than Between Africans and Eurasians. Genetics, 2002, 161, 269-274.	2.9	178
146	In Southern Africa, Brown Oculocutaneous Albinism (BOCA) Maps to the OCA2 Locus on Chromosome 15q: P-Gene Mutations Identified. American Journal of Human Genetics, 2001, 68, 782-787.	6.2	34
147	Human DNA Sequence Variation in a 6.6-kb Region Containing the Melanocortin 1 Receptor Promoter. Genetics, 2001, 158, 1253-1268.	2.9	52
148	Identification ofP gene mutations in individuals with oculocutaneous albinism in sub-Saharan Africa. Human Mutation, 2000, 15, 166-172.	2.5	43
149	High Polymorphism at the Human Melanocortin 1 Receptor Locus. Genetics, 1999, 151, 1547-1557.	2.9	258
150	Human pigmentation genetics: the difference is only skin deep. BioEssays, 1998, 20, 712-721.	2.5	156
151	Evidence for a Common Ethnic Origin of Cystic Fibrosis Mutation 3120+1Gât'A in Diverse Populations. American Journal of Human Genetics, 1998, 63, 656-662.	6.2	34
152	Localization of the Gene Causing Keratolytic Winter Erythema to Chromosome 8p22-p23, and Evidence for a Founder Effect in South African Afrikaans-Speakers. American Journal of Human Genetics, 1997, 61, 370-378.	6.2	38
153	Oculocutaneous albinism (OCA2) in sub-Saharan Africa: distribution of the common 2.7-kb P gene deletion mutation. Human Genetics, 1997, 99, 523-527.	3.8	55
154	Protein trafficking violations. Nature Genetics, 1996, 14, 242-245.	21.4	14
155	XX true hermaphroditism in Southern African Blacks: Exclusion of SRY sequences and uniparental disomy of the X chromosome. American Journal of Medical Genetics Part A, 1995, 55, 53-56.	2.4	29
156	Maternal origin of extra haploid set of chromosomes in third trimester triploid fetuses. American Journal of Medical Genetics Part A, 1995, 58, 360-364.	2.4	29
157	Yeast artificial chromosome cloning. Molecular Biotechnology, 1994, 1, 181-201.	2.4	23
158	Analysis of 40 known cystic fibrosis mutations in South African patients. Clinical Genetics, 1994, 46, 398-400.	2.0	2
159	Haplotype analysis to determine the position of a mutation among closely linked DNA markers. Human Molecular Genetics, 1993, 2, 1007-1014.	2.9	19
160	Glucose-6-phosphate dehydrogenase (G6PD) electrophoretic variants and the PvuII polymorphism in Southern African populations. Human Genetics, 1992, 89, 111-113.	3.8	3
161	Mapping the human CAS2 gene, the homologue of the mouse brown (b) locus, to human Chromosome 9p22-pter. Biochemical and Biophysical Research Communications, 1991, 178, 227-235.	2.1	43
162	Analysis of the transgenome of MET transfectant cell lines reveals that MET activation is accompanied by an interstitial insertion. Human Genetics, 1990, 84, 274-8.	3.8	1

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163	A new marker on chromosome 15, 1115S74, detects three RFLPs. Nucleic Acids Research, 1989, 17, 7121-7121.	14.5	2
164	An anonymous single copy probe, D3S153, detects an SstI RFLP. Nucleic Acids Research, 1989, 17, 7122-7122.	14.5	0
165	A new marker on chromosome 4q, D4S138, detects two RFLPs. Nucleic Acids Research, 1989, 17, 7123-7123.	14.5	0
166	An arbitrary single copy sequence VC64 [D1S86] detects a moderate frequency TaqI RFLP on chromosome 1. Nucleic Acids Research, 1989, 17, 4009-4009.	14.5	1
167	Detection of a 3 allele AvaII RFLP by a single copy anonymous DNA sequence VC75 [D7S404] localized to chromosome 7. Nucleic Acids Research, 1989, 17, 5421-5421.	14.5	0
168	An arbitrary single copy sequence VC64 [D1S86] detects a moderate frequency TaqI RFLP on chromosome 1. Nucleic Acids Research, 1989, 17, 5422-5422.	14.5	1
169	Detection of a 3 allele AvaII RFLP by a single copy anonymous DNA sequence VC75 [D7S404] localized to chromosome 7. Nucleic Acids Research, 1989, 17, 4008-4008.	14.5	0
170	A high frequency two allele TaqI RFLP detected by an anonymous sequence VC61 [D2S65] on chromosome 2. Nucleic Acids Research, 1989, 17, 5424-5424.	14.5	0
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#	Article	IF	Citations
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