Loci associated with skin pigmentation identified in Afr

Science 358, DOI: 10.1126/science.aan8433

Citation Report

#	Article	IF	CITATIONS
1	An Unexpectedly Complex Architecture for Skin Pigmentation in Africans. Cell, 2017, 171, 1340-1353.e14.	13.5	134
2	Skin color variation in Africa. Science, 2017, 358, 867-868.	6.0	8
3	How have our clocks evolved? Adaptive and demographic history of the out-of-African dispersal told by polymorphic loci in circadian genes. Chronobiology International, 2018, 35, 511-532.	0.9	7
4	Tales of Human Migration, Admixture, and Selection in Africa. Annual Review of Genomics and Human Genetics, 2018, 19, 405-428.	2.5	78
5	Focus on African diversity confirms complexity of skin pigmentation genetics. Genome Biology, 2018, 19, 13.	3.8	4
6	The HIrisPlex-S system for eye, hair and skin colour prediction from DNA: Introduction and forensic developmental validation. Forensic Science International: Genetics, 2018, 35, 123-135.	1.6	199
7	Genetic architecture and selective sweeps after polygenic adaptation to distant trait optima. PLoS Genetics, 2018, 14, e1007794.	1.5	48
8	The evolutionary history of Southern Africa. Current Opinion in Genetics and Development, 2018, 53, 157-164.	1.5	10
9	Rapid evolution of a skin-lightening allele in southern African KhoeSan. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 13324-13329.	3.3	17
10	Differential gene regulation underlies variation in melanic plumage coloration in the darkâ€eyed junco (<i>Junco hyemalis</i>). Molecular Ecology, 2018, 27, 4501-4515.	2.0	41
11	Frontiers in pigment cell and melanoma research. Pigment Cell and Melanoma Research, 2018, 31, 728-735.	1.5	10
13	Deciphering the Emerging Complexities of Molecular Mechanisms at GWAS Loci. American Journal of Human Genetics, 2018, 103, 637-653.	2.6	93
14	The critical needs and challenges for genetic architecture studies in Africa. Current Opinion in Genetics and Development, 2018, 53, 113-120.	1.5	57
15	Genetic Resistance to Mycobacterium tuberculosis Infection and Disease. Frontiers in Immunology, 2018, 9, 2219.	2.2	29
16	Clinical and Biological Characterization of Skin Pigmentation Diversity and Its Consequences on UV Impact. International Journal of Molecular Sciences, 2018, 19, 2668.	1.8	158
17	The Year In Genetic Anthropology: New Lands, New Technologies, New Questions. American Anthropologist, 2018, 120, 266-277.	0.7	5
18	Darwinian Positive Selection on the Pleiotropic Effects of KITLG Explain Skin Pigmentation and Winter Temperature Adaptation in Eurasians. Molecular Biology and Evolution, 2018, 35, 2272-2283.	3.5	27
19	Selecting microhaplotypes optimized for different purposes. Electrophoresis, 2018, 39, 2815-2823.	1.3	39

#	Article	IF	CITATIONS
20	Genotype-Corrector: improved genotype calls for genetic mapping in F2 and RIL populations. Scientific Reports, 2018, 8, 10088.	1.6	22
21	African genetic diversity provides novel insights into evolutionary history and local adaptations. Human Molecular Genetics, 2018, 27, R209-R218.	1.4	38
23	Cellular localization of the K ⁺ â€dependent Na ⁺ –Ca ²⁺ exchanger <scp>NCKX</scp> 5 and the role of the cytoplasmic loop in its distribution in pigmented cells. Pigment Cell and Melanoma Research, 2019, 32, 55-67.	1.5	15
24	MEK inhibition remodels the active chromatin landscape and induces SOX10 genomic recruitment in BRAF(V600E) mutant melanoma cells. Epigenetics and Chromatin, 2019, 12, 50.	1.8	12
25	LEI: A Novel Allele Frequency-Based Feature Selection Method for Multi-ancestry Admixed Populations. Scientific Reports, 2019, 9, 11103.	1.6	2
26	A four-DNA methylation biomarker is a superior predictor of survival of patients with cutaneous melanoma. ELife, 2019, 8, .	2.8	69
27	Human races are not like dog breeds: refuting a racist analogy. Evolution: Education and Outreach, 2019, 12, .	0.3	12
28	Thinking About the Evolution of Complex Traits in the Era of Genome-Wide Association Studies. Annual Review of Genomics and Human Genetics, 2019, 20, 461-493.	2.5	186
29	Meta-analysis of GWA studies provides new insights on the genetic architecture of skin pigmentation in recently admixed populations. BMC Genetics, 2019, 20, 59.	2.7	32
30	Darker skin color is associated with a lower likelihood of smoking cessation among males but not females. Social Science and Medicine, 2019, 240, 112562.	1.8	4
31	Frameshift Variant in MFSD12 Explains the Mushroom Coat Color Dilution in Shetland Ponies. Genes, 2019, 10, 826.	1.0	14
32	Evaluation of the HIrisPlex-S system in a Brazilian population sample. Forensic Science International: Genetics Supplement Series, 2019, 7, 794-796.	0.1	2
33	Reconstructed lost Native American populations from Eastern Brazil are shaped by differential Jê/Tupi ancestry. Genome Biology and Evolution, 2019, 11, 2593-2604.	1.1	8
34	The Paradox Behind the Pattern of Rapid Adaptive Radiation: How Can the Speciation Process Sustain Itself Through an Early Burst?. Annual Review of Ecology, Evolution, and Systematics, 2019, 50, 569-593.	3.8	67
35	A GWAS in Latin Americans highlights the convergent evolution of lighter skin pigmentation in Eurasia. Nature Communications, 2019, 10, 358.	5.8	130
36	Population structure of human gut bacteria in a diverse cohort from rural Tanzania and Botswana. Genome Biology, 2019, 20, 16.	3.8	66
37	Identification of a Missense Variant in MFSD12 Involved in Dilution of Phaeomelanin Leading to White or Cream Coat Color in Dogs. Genes, 2019, 10, 386.	1.0	20
38	Resolving the Insertion Sites of Polymorphic Duplications Reveals a HERC2 Haplotype under Selection. Genome Biology and Evolution, 2019, 11, 1679-1690.	1.1	6

#	Article	IF	CITATIONS
39	Comparative genomics provides new insights into the remarkable adaptations of the African wild dog (Lycaon pictus). Scientific Reports, 2019, 9, 8329.	1.6	23
40	Population genomics perspectives on convergent adaptation. Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20180236.	1.8	56
42	The Genetics of Human Skin and Hair Pigmentation. Annual Review of Genomics and Human Genetics, 2019, 20, 41-72.	2.5	98
43	Susceptibility Loci for Tanning Ability in the JapaneseÂPopulation Identified by aÂGenome-WideÂAssociation Study from the TohokuÂMedical Megabank Project Cohort Study. Journal of Investigative Dermatology, 2019, 139, 1605-1608.e13.	0.3	14
44	Potential Mutations in Chinese Pathologic Myopic Patients and Contributions to Phenotype. Current Molecular Medicine, 2019, 18, 689-697.	0.6	2
45	Pervasive function and evidence for selection across standing genetic variation in S. cerevisiae. Nature Communications, 2019, 10, 1222.	5.8	10
46	Genetic signatures of gene flow and malaria-driven natural selection in sub-Saharan populations of the "endemic Burkitt Lymphoma belt". PLoS Genetics, 2019, 15, e1008027.	1.5	23
47	A Genome-Wide Association Study of Skin and Iris Pigmentation among Individuals of South Asian Ancestry. Genome Biology and Evolution, 2019, 11, 1066-1076.	1.1	21
48	Prediction of skin color, tanning and freckling from DNA in Polish population: linear regression, random forest and neural network approaches. Human Genetics, 2019, 138, 635-647.	1.8	15
50	The Emergence of Genomic Research in Africa and New Frameworks for Equity in Biomedical Research. Ethnicity and Disease, 2019, 29, 179-186.	1.0	18
51	Toward a more humane genetics education: Learning about the social and quantitative complexities of human genetic variation research could reduce racial bias in adolescent and adult populations. Science Education, 2019, 103, 529-560.	1.8	61
52	Positive selection in Europeans and East-Asians at the ABCA12 gene. Scientific Reports, 2019, 9, 4843.	1.6	1
53	Ancestry-Specific Analyses Reveal Differential Demographic Histories and Opposite Selective Pressures in Modern South Asian Populations. Molecular Biology and Evolution, 2019, 36, 1628-1642.	3.5	20
54	Nutrition and its role in human evolution. Journal of Internal Medicine, 2019, 285, 533-549.	2.7	43
55	Meta-analysis and prioritization of human skin pigmentation-associated GWAS-SNPs using ENCODE data-based web-tools. Archives of Dermatological Research, 2019, 311, 163-171.	1.1	9
56	Insights into malaria susceptibility using genome-wide data on 17,000 individuals from Africa, Asia and Oceania. Nature Communications, 2019, 10, 5732.	5.8	126
57	Genetics of cognitive trajectory in Brazilians: 15 years of follow-up from the BambuÃ-Epigen Cohort Study of Aging. Scientific Reports, 2019, 9, 18085.	1.6	6
58	Precision medicine validation: identifying the <i>MYBPC</i> 3 A31P variant with whole-genome sequencing in two Maine Coon cats with hypertrophic cardiomyopathy. Journal of Feline Medicine and Surgery, 2019, 21, 1086-1093.	0.6	10

ARTICLE IF CITATIONS # Shades of complexity: New perspectives on the evolution and genetic architecture of human skin. 59 2.1 45 American Journal of Physical Anthropology, 2019, 168, 4-26. A curated gene list for expanding the horizons of pigmentation biology. Pigment Cell and Melanoma 1.5 Research, 2019, 32, 348-358. Bioinformatics-based analysis reveals elevated MFSD12 as a key promoter of cell proliferation and a 61 2.6 33 potential therapeutic target in melanoma. Oncogene, 2019, 38, 1876-1891. Reconstructing the History of Polygenic Scores Using Coalescent Trees. Genetics, 2019, 211, 235-262. 1.2 Off-target phenotypes in forensic DNA phenotyping and biogeographic ancestry inference: A resource. 63 1.6 15 Forensic Science International: Genetics, 2019, 38, 93-104. Microhaplotypes in forensic genetics. Forensic Science International: Genetics, 2019, 38, 54-69. 1.6 Population genetic evidence for positive and purifying selection acting at the human IFN- \hat{I}^3 locus in 65 2.2 4 Africa. Genes and Immunity, 2019, 20, 143-157. Genetic-based signatures of the latitudinal differences in chronotype. Biological Rhythm Research, 0.4 66 2019, 50, 255-271. 67 Forensic genetics and the prediction of race: What is the problem?. BioSocieties, 2020, 15, 329-349. 0.8 31 The Evolutionary History of Human Skin Pigmentation. Journal of Molecular Evolution, 2020, 88, 77-87. 0.8 Peptidomic analysis of blastocyst culture medium and the effect of peptide derived from blastocyst culture medium on blastocyst formation and viability. Molecular Reproduction and Development, 69 3 1.0 2020, 87, 191-201. 2019 Curt Stern Award Address. American Journal of Human Genetics, 2020, 106, 297-298. MFSD12 mediates the import of cysteine into melanosomes and lysosomes. Nature, 2020, 588, 699-704. 71 13.7 52 Overcoming Immune Evasion in Melanoma. International Journal of Molecular Sciences, 2020, 21, 8984. 1.8 MicroHapDB: A Portable and Extensible Database of All Published Microhaplotype Marker and 73 1.1 14 Frequency Data. Frontiers in Genetics, 2020, 11, 781. Quantifying and contextualizing the impact of bioRxiv preprints through automated social media 74 audience segmentation. PLoS Biology, 2020, 18, e3000860. SLC45A2 protein stability and regulation of melanosome pH determine melanocyte pigmentation. 75 0.9 49 Molecular Biology of the Cell, 2020, 31, 2687-2702. Genetics and geography of leukocyte telomere length in sub-Saharan Africans. Human Molecular 1.4 Genetics, 2020, 29, 3014-3020.

	Сітатіс	CITATION REPORT	
#	Article	IF	CITATIONS
77	The road ahead in genetics and genomics. Nature Reviews Genetics, 2020, 21, 581-596.	7.7	118
78	Adaptation and coâ€adaptation of skin pigmentation and vitamin D genes in native Americans. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2020, 184, 1060-1077.	0.7	5
79	The genomics of coloration provides insights into adaptive evolution. Nature Reviews Genetics, 2020, 21, 461-475.	7.7	88
80	Quantifying the potential causes of Neanderthal extinction: Abrupt climate change versus competition and interbreeding. Quaternary Science Reviews, 2020, 238, 106331.	1.4	38
81	Whole-genome resequencing of wild and domestic sheep identifies genes associated with morphological and agronomic traits. Nature Communications, 2020, 11, 2815.	5.8	142
82	<p>Melanogenic Difference Consideration in Ethnic Skin Type: A Balance Approach Between Skin Brightening Applications and Beneficial Sun Exposure</p> . Clinical, Cosmetic and Investigational Dermatology, 2020, Volume 13, 215-232.	0.8	13
83	Functional colour genes and signals of selection in colourâ€polymorphic salamanders. Molecular Ecology, 2020, 29, 1284-1299.	2.0	15
84	Studying human and nonhuman primate evolutionary biology with powerful in vitro and in vivo functional genomics tools. Evolutionary Anthropology, 2020, 29, 143-158.	1.7	9
85	The importance of including ethnically diverse populations in studies of quantitative trait evolution. Current Opinion in Genetics and Development, 2020, 62, 30-35.	1.5	5
86	Gene and environment interactions and phenotypes. , 2020, , 79-107.		1
87	Evaluating the promise of inclusion of African ancestry populations in genomics. Npj Genomic Medicine, 2020, 5, 5.	1.7	86
88	Extreme genetic signatures of local adaptation during Lotus japonicus colonization of Japan. Nature Communications, 2020, 11, 253.	5.8	30
89	Research Techniques Made Simple: Cell Biology Methods for the Analysis of Pigmentation. Journal of Investigative Dermatology, 2020, 140, 257-268.e8.	0.3	22
90	Dating genomic variants and shared ancestry in population-scale sequencing data. PLoS Biology, 2020, 18, e3000586.	2.6	127
91	The 4 D's of Pellagra and Progress. International Journal of Tryptophan Research, 2020, 13, 117864692091015.	1.0	8
92	Membrane transport proteins in melanosomes: Regulation of ions for pigmentation. Biochimica Et Biophysica Acta - Biomembranes, 2020, 1862, 183318.	1.4	46
93	Evaluation of fetal medicine foundation algorithm in predicting small-for-gestational-age neonates. Journal of Maternal-Fetal and Neonatal Medicine, 2021, 34, 876-882.	0.7	0
94	A linkage disequilibrium-based statistical test for Genome-Wide Epistatic Selection Scans in structured populations. Heredity, 2021, 126, 77-91.	1.2	5

#	Article	IF	CITATIONS
95	Identifying adaptive alleles in the human genome: from selection mapping to functional validation. Human Genetics, 2021, 140, 241-276.	1.8	13
96	Translational genetics: a challenging but important path. British Journal of Dermatology, 2021, 184, 800-801.	1.4	0
99	The influence of evolutionary history on human health and disease. Nature Reviews Genetics, 2021, 22, 269-283.	7.7	133
100	Persistence of mature dendritic cells, T _H 2A, and Tc2 cells characterize clinically resolved atopic dermatitis under IL-4Rα blockade. Science Immunology, 2021, 6, .	5.6	76
101	Melanin Based Classification of Skin Types and Their Susceptibility to UV-Induced Cancer. , 2021, , 41-67.		1
102	Evolution of Pigment Pattern Formation in Teleosts. , 2021, , 309-342.		4
103	Melanins in Vertebrates. , 2021, , 45-89.		4
104	Dissecting dynamics and differences of selective pressures in the evolution of human pigmentation. Biology Open, 2021, 10, .	0.6	4
105	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.	1.6	4
106	Genetic loci associated with skin pigmentation in African Americans and their effects on vitamin D deficiency. PLoS Genetics, 2021, 17, e1009319.	1.5	10
107	Vitamin-related phenotypic adaptation to exposomal factors: The folate-vitamin D-exposome triad. Molecular Aspects of Medicine, 2022, 87, 100944.	2.7	5
109	Spontaneously Resolved Atopic Dermatitis Shows Melanocyte and Immune Cell Activation Distinct From Healthy Control Skin. Frontiers in Immunology, 2021, 12, 630892.	2.2	21
110	Scanning the human genome for "signatures―of positive selection: Transformative opportunities and ethical obligations. Evolutionary Anthropology, 2021, 30, 113-121.	1.7	6
111	Genetic diversity of the Sudanese: insights on origin and implications for health. Human Molecular Genetics, 2021, 30, R37-R41.	1.4	3
112	The constraints of racialization: How classification and valuation hinder scientific research on human variation. American Journal of Physical Anthropology, 2021, 175, 376-386.	2.1	9
113	Chemical and biochemical control of skin pigmentation with special emphasis on mixed melanogenesis. Pigment Cell and Melanoma Research, 2021, 34, 730-747.	1.5	38
114	Feather Gene Expression Elucidates the Developmental Basis of Plumage Iridescence in African Starlings. Journal of Heredity, 2021, 112, 417-429.	1.0	15
115	Identification of Three Novel Susceptibility Loci for Inflammatory Bowel Disease in Koreans in an Extended Genome-Wide Association Study. Journal of Crohn's and Colitis, 2021, 15, 1898-1907.	0.6	13

#	Article	IF	CITATIONS
116	Prediction of eye and hair pigmentation phenotypes using the HIrisPlex system in a Brazilian admixed population sample. International Journal of Legal Medicine, 2021, 135, 1329-1339.	1.2	8
117	Genome-wide DNA methylation and RNA expression differences correlate with invasiveness in melanoma cell lines. Epigenomics, 2021, 13, 577-598.	1.0	6
118	Efficient mixed model approach for large-scale genome-wide association studies of ordinal categorical phenotypes. American Journal of Human Genetics, 2021, 108, 825-839.	2.6	25
119	How a membrane transporter keeps melanocytes in the red. Pigment Cell and Melanoma Research, 2021, 34, 666-669.	1.5	0
120	Comparative transcriptomics of albino and warninglyâ€coloured caterpillars. Ecology and Evolution, 2021, 11, 7507-7517.	0.8	6
121	Cultural evolution of genetic heritability. Behavioral and Brain Sciences, 2022, 45, 1-147.	0.4	26
122	The evolution of human skin pigmentation involved the interactions of genetic, environmental, and cultural variables. Pigment Cell and Melanoma Research, 2021, 34, 707-729.	1.5	32
123	Five genetic variants explain over 70% of hair coat pheomelanin intensity variation in purebred and mixed breed domestic dogs. PLoS ONE, 2021, 16, e0250579.	1.1	6
124	Melanosome Biogenesis in the Pigmentation of Mammalian Skin. Integrative and Comparative Biology, 2021, 61, 1517-1545.	0.9	44
125	The quagmire of race, genetic ancestry, and health disparities. Journal of Clinical Investigation, 2021, 131, .	3.9	14
126	Genome of PeÅŸtera Muierii skull shows high diversity and low mutational load in pre-glacial Europe. Current Biology, 2021, 31, 2973-2983.e9.	1.8	18
127	Prediction of eye, hair and skin colour in Latin Americans. Forensic Science International: Genetics, 2021, 53, 102517.	1.6	6
128	Association of pigmentation related-genes polymorphisms and geographic environmental variables in the Chinese population. Hereditas, 2021, 158, 24.	0.5	4
129	Vitamin D and the risk for cancer: A molecular analysis. Biochemical Pharmacology, 2022, 196, 114735.	2.0	36
130	NNT mediates redox-dependent pigmentation via a UVB- and MITF-independent mechanism. Cell, 2021, 184, 4268-4283.e20.	13.5	35
131	A custom capture sequence approach for oculocutaneous albinism identifies structural variant alleles at the <i>OCA2</i> locus. Human Mutation, 2021, 42, 1239-1253.	1.1	7
132	Human biological variation and the "normal― American Journal of Human Biology, 2021, 33, e23658.	0.8	14
133	Phylogenetic Analysis of Core Melanin Synthesis Genes Provides Novel Insights Into the Molecular Basis of Albinism in Fish. Frontiers in Genetics, 2021, 12, 707228.	1.1	16

#	Article	IF	CITATIONS
134	The critical role of glutathione redox homeostasis towards oxidation in ermanin-induced melanogenesis. Free Radical Biology and Medicine, 2021, 176, 392-405.	1.3	7
135	Human melanocyte development and melanoma dedifferentiation at single-cell resolution. Nature Cell Biology, 2021, 23, 1035-1047.	4.6	59
136	Local adaptation and archaic introgression shape global diversity at human structural variant loci. ELife, 2021, 10, .	2.8	33
137	Identification of the Role of Wnt/Ĵ²-Catenin Pathway Through Integrated Analyses and in vivo Experiments in Vitiligo. Clinical, Cosmetic and Investigational Dermatology, 2021, Volume 14, 1089-1103.	0.8	8
139	Advances and challenges in quantitative delineation of the genetic architecture of complex traits. Quantitative Biology, 2021, 9, 168-184.	0.3	0
140	The evolution of group differences in changing environments. PLoS Biology, 2021, 19, e3001072.	2.6	37
141	Evolutionary genetics of skin pigmentation in African populations. Human Molecular Genetics, 2021, 30, R88-R97.	1.4	23
142	Cross-cultural perception of female facial appearance: A multi-ethnic and multi-centre study. PLoS ONE, 2021, 16, e0245998.	1.1	21
143	The evolution of skin pigmentation-associated variation in West Eurasia. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	50
144	Genome-Wide Natural Selection Signatures Are Linked to Genetic Risk of Modern Phenotypes in the Japanese Population. Molecular Biology and Evolution, 2020, 37, 1306-1316.	3.5	22
155	Terahertz spectroscopy of human skin tissue models with different melanin content. Biomedical Optics Express, 2019, 10, 2942.	1.5	28
157	Performance of Fetal Medicine Foundation Software for Pre-Eclampsia Prediction Upon Marker Customization: Cross-Sectional Study. Journal of Medical Internet Research, 2019, 21, e14738.	2.1	3
159	GWAS Identifies Multiple Genetic Loci for Skin Color in Korean Women. Journal of Investigative Dermatology, 2022, 142, 1077-1084.	0.3	10
160	Forensic DNA Phenotyping: starting point to prediction model in Pernambuco population, Brazil. Research, Society and Development, 2021, 10, e262101320955.	0.0	0
164	Skin cancer: an overview regarding treatment and its cosmetic repair. Journal of Dermatology & Cosmetology, 2018, 2, .	0.1	0
169	Cellular and Molecular Changes. , 2019, , 70-86.		0
172	Human Biodiversity and Close Encounters. , 2020, , 11-26.		0
173	Similarity-Based Analysis of Allele Frequency Distribution among Multiple Populations Identifies Adaptive Genomic Structural Variants. Molecular Biology and Evolution, 2022, 39, .	3.5	6

#	Article	IF	CITATIONS
174	The color of normal: How a Eurocentric focus erases pigmentation complexity. American Journal of Human Biology, 2021, 33, e23554.	0.8	9
175	Interpreting the Meaning in Our Genomes: Perspectives from Biochemistry, Genetics, Infectious Disease, and Dance. , 2020, , 213-228.		0
180	Canine coat pigmentation genetics: a review. Animal Genetics, 2022, 53, 3-34.	0.6	15
181	A large Canadian cohort provides insights into the genetic architecture of human hair colour. Communications Biology, 2021, 4, 1253.	2.0	11
182	Advances in integrative African genomics. Trends in Genetics, 2022, 38, 152-168.	2.9	6
185	Toward Interactively Balancing the Screen Time of Actors Based on Observable Phenotypic Traits in Live Telecast. Proceedings of the ACM on Human-Computer Interaction, 2020, 4, 1-18.	2.5	1
187	MiRNA-mRNA Integration Analysis Reveals the Regulatory Roles of MiRNAs in Shell Pigmentation of the Manila clam (Ruditapes philippinarum). Marine Biotechnology, 2021, 23, 976-993.	1.1	6
188	Race, Ethnicity, and the Scarr-Rowe Hypothesis: A Cautionary Example of Fringe Science Entering the Mainstream. Perspectives on Psychological Science, 2022, 17, 696-710.	5.2	6
191	Human population genomics approach in food metabolism. , 2022, , 433-449.		0
192	Predicting Physical Appearance from DNA Data—Towards Genomic Solutions. Genes, 2022, 13, 121.	1.0	8
193	Integrative analysis of 3604 GWAS reveals multiple novel cell type-specific regulatory associations. Genome Biology, 2022, 23, 13.	3.8	19
194	Vitamin D and Pigmented Skin. Nutrients, 2022, 14, 325.	1.7	4
195	Uniparental markers reveal new insights on subcontinental ancestry and sex-biased admixture in Brazil. Molecular Genetics and Genomics, 2022, 297, 419.	1.0	0
196	Genetic Connections and Convergent Evolution of Tropical Indigenous Peoples in Asia. Molecular Biology and Evolution, 2022, 39, .	3.5	2
198	Genetic ancestry in precision medicine is reshaping the race debate. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2203033119.	3.3	17
199	Whole-genome sequencing of 1,171 elderly admixed individuals from Brazil. Nature Communications, 2022, 13, 1004.	5.8	35
200	Multidisciplinary investigation reveals an individual of West African origin buried in a Portuguese Mesolithic shell midden four centuries ago. Journal of Archaeological Science: Reports, 2022, 42, 103370.	0.2	3
201	Gene Geographic Atlas of DNA Markers Controlling Human Eye and Hair Color. Russian Journal of Genetics, 2021, 57, 1370-1388.	0.2	2

#	Article	IF	CITATIONS
203	An overview of SNP-SNP microhaplotypes in the 26 populations of the 1000 Genomes Project. International Journal of Legal Medicine, 2022, 136, 1211-1226.	1.2	6
204	Forensic DNA phenotyping: Inferring phenotypic traits from crime scene DNA. Journal of Clinical Forensic and Legal Medicine, 2022, 88, 102351.	0.5	3
205	Ablation of Proton/Glucose Exporter SLC45A2 Enhances Melanosomal Glycolysis to Inhibit Melanin Biosynthesis and Promote Melanoma Metastasis. Journal of Investigative Dermatology, 2022, 142, 2744-2755.e9.	0.3	5
206	Gene expression in male and female stickleback from populations with convergent and divergent throat coloration. Ecology and Evolution, 2022, 12, e8860.	0.8	2
207	Importance of Including Non-European Populations in Large Human Genetic Studies to Enhance Precision Medicine. Annual Review of Biomedical Data Science, 2022, 5, 321-339.	2.8	17
208	Books on Human Evolution. American Biology Teacher, 2022, 84, 318-319.	0.1	0
209	The evolution of human skin pigmentation: A changing medley of vitamins, genetic variability, and <scp>UV</scp> radiation during human expansion. American Journal of Biological Anthropology, 2023, 180, 252-271.	0.6	7
210	Including diverse and admixed populations in genetic epidemiology research. Genetic Epidemiology, 2022, 46, 347-371.	0.6	11
212	A systematic review of skin ageing genes: gene pleiotropy and genes on the chromosomal band 16q24.3 may drive skin ageing. Scientific Reports, 2022, 12, .	1.6	3
213	MFSD12 affects glycosphingolipid metabolism by modulating lysosome homeostasis. Protein and Cell, 0, , .	4.8	2
214	Comprehensive in Silico Analyses of Single Nucleotide Variants of the Human Orthologues of 171 Murine Loci to Seek Novel Insights into the Genetics of Human Pigmentation. Proceedings of the Zoological Society, 0, , .	0.4	0
215	A Comprehensive Study to Explore Tyrosinase Inhibitory Medicinal Plants and Respective Phytochemicals for Hyperpigmentation; Molecular Approach and Future Perspectives. Current Pharmaceutical Biotechnology, 2023, 24, 780-813.	0.9	2
216	Novel genetic associations with five aesthetic facial traits: A genome-wide association study in the Chinese population. Frontiers in Genetics, 0, 13, .	1.1	0
217	Signatures of Convergent Evolution and Natural Selection at the Alcohol Dehydrogenase Gene Region are Correlated with Agriculture in Ethnically Diverse Africans. Molecular Biology and Evolution, 2022, 39, .	3.5	5
218	Genetic adaptation of skin pigmentation in highland Tibetans. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	9
219	Evaluation of Personalized Skincare Through in-silico Gene Interactive Networks and Cellular Responses to UVR and Oxidative Stress. Clinical, Cosmetic and Investigational Dermatology, 0, Volume 15, 2221-2243.	0.8	0
220	Redundancy analysis, genomeâ€wide association studies and the pigmentation of brown trout (<i>Salmo) Tj ETC</i>	Qq0 0 0 rgl 0.7	BT /Overlock

221	Eliminating Disparities in Cardiovascular Disease for Black Women. Journal of the American College of Cardiology, 2022, 80, 1762-1771.	1.2	9
-----	--	-----	---

#	Article	IF	CITATIONS
222	Melanin and Neuromelanin: Linking Skin Pigmentation and Parkinson's Disease. Movement Disorders, 2023, 38, 185-195.	2.2	5
224	Définition et physiologie des peaux fortement pigmentées et/ou dites « noires ». , 2022, , 31-35.		0
225	The potential impact of melanosomal pH and metabolism on melanoma. Frontiers in Oncology, 0, 12, .	1.3	1
226	Prediction and association analyses of skin phenotypes in Japanese females using genetic, environmental, and physical features. Skin Research and Technology, 2023, 29, .	0.8	2
227	Skin Complexion and the Blush. Emotion Review, 0, , 175407392211502.	2.1	0
230	Analysis of the genetic loci of pigment pattern evolution in vertebrates. Biological Reviews, 2023, 98, 1250-1277.	4.7	6
231	Nutrigenomics in the context of evolution. Redox Biology, 2023, 62, 102656.	3.9	3
233	The genetic and evolutionary basis of gene expression variation in East Africans. Genome Biology, 2023, 24, .	3.8	3
234	Whole-genome sequencing reveals a complex African population demographic history and signatures of local adaptation. Cell, 2023, 186, 923-939.e14.	13.5	34
236	Evolutionary Genetics and Admixture in African Populations. Genome Biology and Evolution, 2023, 15, .	1.1	7
238	Recent advances in Forensic DNA Phenotyping of appearance, ancestry and age. Forensic Science International: Genetics, 2023, 65, 102870.	1.6	9
239	A gene-level test for directional selection on gene expression. Genetics, 2023, 224, .	1.2	2
240	A frame-shift mutation in COMTD1 is associated with impaired pheomelanin pigmentation in chicken. PLoS Genetics, 2023, 19, e1010724.	1.5	1
251	Evolution of human skin pigmentation and vitamin D. , 2024, , 9-25.		0
257	Forensic DNA phenotyping in the next-generation sequencing era. , 2024, , 311-336.		0