

Cherni Lotfi

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

733
citations

471509

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552781

26
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39
all docs

39
docs citations

39
times ranked

868
citing authors

#	ARTICLE	IF	CITATIONS
1	Mitochondrial DNA and Alzheimer’s disease: a first case–control study of the Tunisian population. <i>Molecular Biology Reports</i> , 2022, 49, 1687-1700.	2.3	3
2	New Insight into the human genetic diversity in North African populations by genotyping of SNPs in <i>DRD3</i> , <i>CSMD1</i> and <i>NRG1</i> genes. <i>Molecular Genetics & Genomic Medicine</i> , 2022, 10, e1871.	1.2	2
3	North Asian population relationships in a global context. <i>Scientific Reports</i> , 2022, 12, 7214.	3.3	3
4	Genetic diversity of the North African population revealed by the typing of SNPs in the <i>DRD2/ANKK1</i> genomic region. <i>Gene</i> , 2021, 777, 145466.	2.2	2
5	Genetic relationships of Southwest Asian and Mediterranean populations. <i>Forensic Science International: Genetics</i> , 2021, 53, 102528.	3.1	7
6	STAT3 polymorphisms in North Africa and its implication in breast cancer. <i>Molecular Genetics & Genomic Medicine</i> , 2021, 9, e1744.	1.2	1
7	Insights into the Middle Eastern paternal genetic pool in Tunisia: high prevalence of T-M70 haplogroup in an Arab population. <i>Scientific Reports</i> , 2021, 11, 15728.	3.3	1
8	The distinctive geographic patterns of common pigmentation variants at the <i>OCA2</i> gene. <i>Scientific Reports</i> , 2020, 10, 15433.	3.3	8
9	Genetic relationships of European, Mediterranean, and SW Asian populations using a panel of 55 AISNPs. <i>European Journal of Human Genetics</i> , 2019, 27, 1885-1893.	2.8	22
10	Investigation of the genetic structure of Kabyle and Chaouia Algerian populations through the polymorphism of Alu insertion markers. <i>Annals of Human Biology</i> , 2019, 46, 150-159.	1.0	1
11	Usefulness of <i>COMT</i> gene polymorphisms in North African populations. <i>Gene</i> , 2019, 696, 186-196.	2.2	7
12	Expression and polymorphism of micro-RNA according to body mass index and breast cancer presentation in Tunisian patients. <i>Journal of Leukocyte Biology</i> , 2019, 105, 317-327.	3.3	8
13	Population genetics-informed meta-analysis in seven genes associated with risk to dengue fever disease. <i>Infection, Genetics and Evolution</i> , 2018, 62, 60-72.	2.3	16
14	Ancient and recent Middle Eastern maternal genetic contribution to North Africa as viewed by mtDNA diversity in Tunisian Arab populations. <i>American Journal of Human Biology</i> , 2018, 30, e23100.	1.6	13
15	The Orientalisation of North Africa: New hints from the study of autosomal STRs in an Arab population. <i>Annals of Human Biology</i> , 2017, 44, 180-190.	1.0	12
16	Mitochondrial DNA analysis of Tunisians reveals a mosaic genetic structure with recent population expansion. <i>HOMO- Journal of Comparative Human Biology</i> , 2017, 68, 298-315.	0.7	8
17	Reconciling evidence from ancient and contemporary genomes: a major source for the European Neolithic within Mediterranean Europe. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20161976.	2.6	22
18	Distribution of xenobiotic metabolising enzyme genotypes in different Tunisian populations. <i>Annals of Human Biology</i> , 2017, 44, 366-372.	1.0	4

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19	Evaluating a subset of ancestry informative SNPs for discriminating among Southwest Asian and circum-Mediterranean populations. <i>Forensic Science International: Genetics</i> , 2016, 23, 153-158.	3.1	25
20	Genetic variation in Tunisia in the context of human diversity worldwide. <i>American Journal of Physical Anthropology</i> , 2016, 161, 62-71.	2.1	29
21	52 additional reference population samples for the 55 AISNP panel. <i>Forensic Science International: Genetics</i> , 2015, 19, 269-271.	3.1	41
22	An investigation of the genetic diversity of the Kerkennah islands and Mahdia (Tunisia) using biparental markers. <i>Annals of Human Biology</i> , 2014, 41, 53-60.	1.0	2
23	The Arabian Cradle: Mitochondrial Relicts of the First Steps along the Southern Route out of Africa. <i>American Journal of Human Genetics</i> , 2012, 90, 347-355.	6.2	116
24	Un estimateur de distributions ajusté pour une aide à la décision de la neutralité génétique des populations. <i>Irbm</i> , 2012, 33, 24-28.	5.6	0
25	Assessing human genetic diversity in Tunisian Berber populations by Alu insertion polymorphisms. <i>Annals of Human Biology</i> , 2011, 38, 53-58.	1.0	8
26	Genetic analysis of the SNPforID 34-plex ancestry informative SNP panel in Tunisian and Libyan populations. <i>Forensic Science International: Genetics</i> , 2011, 5, e45-e47.	3.1	10
27	Population history of the Red Sea genetic exchanges between the Arabian Peninsula and East Africa signaled in the mitochondrial DNA HV1 haplogroup. <i>American Journal of Physical Anthropology</i> , 2011, 145, 592-598.	2.1	29
28	Internal Diversification of Mitochondrial Haplogroup R0a Reveals Post-Last Glacial Maximum Demographic Expansions in South Arabia. <i>Molecular Biology and Evolution</i> , 2011, 28, 71-78.	8.9	53
29	Ancient Local Evolution of African mtDNA Haplogroups in Tunisian Berber Populations. <i>Human Biology</i> , 2010, 82, 367-384.	0.2	19
30	Data from complete mtDNA sequencing of Tunisian centenarians: Testing haplogroup association and the "golden mean" to longevity. <i>Mechanisms of Ageing and Development</i> , 2009, 130, 222-226.	4.6	26
31	Polymorphisms in one-carbon metabolism pathway genes and risk for bladder cancer in a Tunisian population. <i>Cancer Genetics and Cytogenetics</i> , 2009, 195, 43-53.	1.0	26
32	Post-last glacial maximum expansion from Iberia to North Africa revealed by fine characterization of mtDNA H haplogroup in Tunisia. <i>American Journal of Physical Anthropology</i> , 2009, 139, 253-260.	2.1	54
33	Data for Y-chromosome haplotypes defined by 17 STRs (AmpFLSTR® Yfiler, c) in two Tunisian Berber communities. <i>Forensic Science International</i> , 2006, 160, 80-83.	2.2	31
34	Islands Inside an Island: Reproductive Isolates on Jerba Island. <i>American Journal of Human Biology</i> , 2006, 18, 149-153.	1.6	30
35	Data for 15 autosomal STR markers (Powerplex 16 System) from two Tunisian populations: Kesra (Berber) and Zriba (Arab). <i>Forensic Science International</i> , 2005, 147, 101-106.	2.2	20
36	Y-chromosomal STR haplotypes in three ethnic groups and one cosmopolitan population from Tunisia. <i>Forensic Science International</i> , 2005, 152, 95-99.	2.2	26

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37	Female Gene Pools of Berber and Arab Neighboring Communities in Central Tunisia: Microstructure of mtDNA Variation in North Africa. Human Biology, 2005, 77, 61-70.	0.2	46