

Karima Fadhlaoui-Zid

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

28
papers

698
citations

14
h-index

26
g-index

29
ext. papers

783
ext. citations

3.8
avg, IF

3.34
L-index

#	Paper	IF	Citations
28	Deep analysis of the LRTOMTc.242G>A variant in non-syndromic hearing loss North African patients and the Berber population: Implications for genetic diagnosis and genealogical studies. <i>Molecular Genetics & Genomic Medicine</i> , 2021 , 9, e1810	2.3	2
27	The Marquesans at the fringes of the Austronesian expansion. <i>European Journal of Human Genetics</i> , 2019 , 27, 801-810	5.3	2
26	HLA Class II Allele and Haplotype Diversity in Libyans and Their Genetic Relationships with Other Populations. <i>Immunological Investigations</i> , 2019 , 48, 875-892	2.9	1
25	Whole-genome sequence analysis of a Pan African set of samples reveals archaic gene flow from an extinct basal population of modern humans into sub-Saharan populations. <i>Genome Biology</i> , 2019 , 20, 77	18.3	29
24	Afghanistan: conduits of human migrations identified using AmpFI STR markers. <i>International Journal of Legal Medicine</i> , 2019 , 133, 1659-1666	3.1	1
23	Heterogeneity in Palaeolithic Population Continuity and Neolithic Expansion in North Africa. <i>Current Biology</i> , 2019 , 29, 3953-3959.e4	6.3	11
22	Genetic variation of 17 X-chromosome STR loci in Tunisian population of Nabeul. <i>International Journal of Legal Medicine</i> , 2019 , 133, 85-88	3.1	5
21	The genetic landscape of Mediterranean North African populations through complete mtDNA sequences. <i>Annals of Human Biology</i> , 2018 , 45, 98-104	1.7	10
20	Recent Historical Migrations Have Shaped the Gene Pool of Arabs and Berbers in North Africa. <i>Molecular Biology and Evolution</i> , 2017 , 34, 318-329	8.3	39
19	Whole Y-chromosome sequences reveal an extremely recent origin of the most common North African paternal lineage E-M183 (M81). <i>Scientific Reports</i> , 2017 , 7, 15941	4.9	15
18	Sousse: extreme genetic heterogeneity in North Africa. <i>Journal of Human Genetics</i> , 2015 , 60, 41-9	4.3	16
17	From Arabia to Iberia: A Y chromosome perspective. <i>Gene</i> , 2015 , 564, 141-52	3.8	9
16	Existence of two widespread semi-isolated genetic entities within Mediterranean anchovies. <i>Marine Biology</i> , 2014 , 161, 1063-1071	2.5	12
15	Genome-wide and paternal diversity reveal a recent origin of human populations in North Africa. <i>PLoS ONE</i> , 2013 , 8, e80293	3.7	36
14	Sousse, Tunisia: tumultuous history and high Y-STR diversity. <i>Electrophoresis</i> , 2012 , 33, 3555-63	3.6	10
13	Genetic structure of <i>Octopus vulgaris</i> (Cephalopoda, Octopodidae) in the central Mediterranean Sea inferred from the mitochondrial COIII gene. <i>Comptes Rendus - Biologies</i> , 2012 , 335, 625-36	1.4	19
12	Allele frequencies for 15 autosomal STR markers in the Libyan population. <i>Annals of Human Biology</i> , 2012 , 39, 80-3	1.7	15

11	Biogenic amine production by bacteria isolated from ice-preserved sardine and mackerel. <i>Food Control</i> , 2012 , 25, 89-95	6.2	26
10	Genomic ancestry of North Africans supports back-to-Africa migrations. <i>PLoS Genetics</i> , 2012 , 8, e1002397		219
9	Genetic analysis of the SNPforID 34-plex ancestry informative SNP panel in Tunisian and Libyan populations. <i>Forensic Science International: Genetics</i> , 2011 , 5, e45-7	4.3	10
8	Mitochondrial DNA structure in North Africa reveals a genetic discontinuity in the Nile Valley. <i>American Journal of Physical Anthropology</i> , 2011 , 145, 107-17	2.5	34
7	Genetic structure of Tunisian ethnic groups revealed by paternal lineages. <i>American Journal of Physical Anthropology</i> , 2011 , 146, 271-80	2.5	33
6	Polymorphism of HLA class II genes in Berbers from Southern Tunisia. <i>Tissue Antigens</i> , 2010 , 76, 416-20		9
5	Ancient local evolution of African mtDNA haplogroups in Tunisian Berber populations. <i>Human Biology</i> , 2010 , 82, 367-84	1.2	17
4	Substructure of a Tunisian Berber population as inferred from 15 autosomal short tandem repeat loci. <i>Human Biology</i> , 2008 , 80, 435-48	1.2	17
3	Identification of the CCR5-Delta32 HIV resistance allele and new mutations of the CCR5 gene in different Tunisian populations. <i>Human Immunology</i> , 2007 , 68, 993-1000	2.3	14
2	Genetic diversity in Tunisia: a study based on the GM polymorphism of human immunoglobulins. <i>Human Biology</i> , 2004 , 76, 559-67	1.2	10
1	Mitochondrial DNA heterogeneity in Tunisian Berbers. <i>Annals of Human Genetics</i> , 2004 , 68, 222-33	2.2	77