

# Karima Fadhlaoui-Zid

## List of Publications by Citations

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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	Genomic ancestry of North Africans supports back-to-Africa migrations. <i>PLoS Genetics</i> , <b>2012</b> , 8, e1002397	7.2	219
27	Mitochondrial DNA heterogeneity in Tunisian Berbers. <i>Annals of Human Genetics</i> , <b>2004</b> , 68, 222-33	2.2	77
26	Recent Historical Migrations Have Shaped the Gene Pool of Arabs and Berbers in North Africa. <i>Molecular Biology and Evolution</i> , <b>2017</b> , 34, 318-329	8.3	39
25	Genome-wide and paternal diversity reveal a recent origin of human populations in North Africa. <i>PLoS ONE</i> , <b>2013</b> , 8, e80293	3.7	36
24	Mitochondrial DNA structure in North Africa reveals a genetic discontinuity in the Nile Valley. <i>American Journal of Physical Anthropology</i> , <b>2011</b> , 145, 107-17	2.5	34
23	Genetic structure of Tunisian ethnic groups revealed by paternal lineages. <i>American Journal of Physical Anthropology</i> , <b>2011</b> , 146, 271-80	2.5	33
22	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> , <b>2019</b> , 20, 77	18.3	29
21	Biogenic amine production by bacteria isolated from ice-preserved sardine and mackerel. <i>Food Control</i> , <b>2012</b> , 25, 89-95	6.2	26
20	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> , <b>2012</b> , 335, 625-36	1.4	19
19	Ancient local evolution of African mtDNA haplogroups in Tunisian Berber populations. <i>Human Biology</i> , <b>2010</b> , 82, 367-84	1.2	17
18	Substructure of a Tunisian Berber population as inferred from 15 autosomal short tandem repeat loci. <i>Human Biology</i> , <b>2008</b> , 80, 435-48	1.2	17
17	Sousse: extreme genetic heterogeneity in North Africa. <i>Journal of Human Genetics</i> , <b>2015</b> , 60, 41-9	4.3	16
16	Whole Y-chromosome sequences reveal an extremely recent origin of the most common North African paternal lineage E-M183 (M81). <i>Scientific Reports</i> , <b>2017</b> , 7, 15941	4.9	15
15	Allele frequencies for 15 autosomal STR markers in the Libyan population. <i>Annals of Human Biology</i> , <b>2012</b> , 39, 80-3	1.7	15
14	Identification of the CCR5-Delta32 HIV resistance allele and new mutations of the CCR5 gene in different Tunisian populations. <i>Human Immunology</i> , <b>2007</b> , 68, 993-1000	2.3	14
13	Existence of two widespread semi-isolated genetic entities within Mediterranean anchovies. <i>Marine Biology</i> , <b>2014</b> , 161, 1063-1071	2.5	12
12	Heterogeneity in Palaeolithic Population Continuity and Neolithic Expansion in North Africa. <i>Current Biology</i> , <b>2019</b> , 29, 3953-3959.e4	6.3	11

11	The genetic landscape of Mediterranean North African populations through complete mtDNA sequences. <i>Annals of Human Biology</i> , <b>2018</b> , 45, 98-104	1.7	10
10	Sousse, Tunisia: tumultuous history and high Y-STR diversity. <i>Electrophoresis</i> , <b>2012</b> , 33, 3555-63	3.6	10
9	Genetic analysis of the SNPforID 34-plex ancestry informative SNP panel in Tunisian and Libyan populations. <i>Forensic Science International: Genetics</i> , <b>2011</b> , 5, e45-7	4.3	10
8	Genetic diversity in Tunisia: a study based on the GM polymorphism of human immunoglobulins. <i>Human Biology</i> , <b>2004</b> , 76, 559-67	1.2	10
7	From Arabia to Iberia: A Y chromosome perspective. <i>Gene</i> , <b>2015</b> , 564, 141-52	3.8	9
6	Polymorphism of HLA class II genes in Berbers from Southern Tunisia. <i>Tissue Antigens</i> , <b>2010</b> , 76, 416-20		9
5	Genetic variation of 17 X-chromosome STR loci in Tunisian population of Nabeul. <i>International Journal of Legal Medicine</i> , <b>2019</b> , 133, 85-88	3.1	5
4	The Marquesans at the fringes of the Austronesian expansion. <i>European Journal of Human Genetics</i> , <b>2019</b> , 27, 801-810	5.3	2
3	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 &amp; Genomic Medicine</i> , <b>2021</b> , 9, e1810	2.3	2
2	HLA Class II Allele and Haplotype Diversity in Libyans and Their Genetic Relationships with Other Populations. <i>Immunological Investigations</i> , <b>2019</b> , 48, 875-892	2.9	1
1	Afghanistan: conduits of human migrations identified using AmpFI STR markers. <i>International Journal of Legal Medicine</i> , <b>2019</b> , 133, 1659-1666	3.1	1