

# Atif Adnan

## List of Publications by Year in descending order

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33  
papers

359  
citations

932766

10  
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887659

17  
g-index

41  
all docs

41  
docs citations

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times ranked

272  
citing authors

#	ARTICLE	IF	CITATIONS
1	Improving empirical evidence on differentiating closely related men with RM Y-STRs: A comprehensive pedigree study from Pakistan. <i>Forensic Science International: Genetics</i> , 2016, 25, 45-51.	1.6	62
2	A comprehensive exploration of the genetic legacy and forensic features of Afghanistan and Pakistan Mongolian-descent Hazara. <i>Forensic Science International: Genetics</i> , 2019, 42, e1-e12.	1.6	28
3	Forensic characterization of 15 autosomal STRs in four populations from Xinjiang, China, and genetic relationships with neighboring populations. <i>Scientific Reports</i> , 2018, 8, 4673.	1.6	26
4	Genetic characterization of Y-chromosomal STRs in Hazara ethnic group of Pakistan and confirmation of DYS448 null allele. <i>International Journal of Legal Medicine</i> , 2019, 133, 789-793.	1.2	26
5	Population data of 17 Y-STRs (Yfiler) from Punjabis and Kashmiris of Pakistan. <i>International Journal of Legal Medicine</i> , 2018, 132, 137-138.	1.2	22
6	Mutation analysis at 17 Y-STR loci (Yfiler) in father-son pairs of male pedigrees from Pakistan. <i>Forensic Science International: Genetics</i> , 2018, 36, e17-e18.	1.6	21
7	Population data and phylogenetic structure of Han population from Jiangsu province of China on GlobalFiler STR loci. <i>International Journal of Legal Medicine</i> , 2018, 132, 1301-1304.	1.2	18
8	EMPOP-quality mtDNA control region sequences from Kashmiri of Azad Jammu & Kashmir, Pakistan. <i>Forensic Science International: Genetics</i> , 2016, 25, 125-131.	1.6	16
9	Comprehensive biomechanical analysis of three clinically used fixation constructs for posterior malleolar fractures using cadaveric and finite element analysis. <i>Scientific Reports</i> , 2020, 10, 18639.	1.6	13
10	Tracing the Genetic Legacy of the Tibetan Empire in the Balti. <i>Molecular Biology and Evolution</i> , 2021, 38, 1529-1536.	3.5	13
11	Discriminating power of rapidly mutating Y-STRs in deep rooted endogamous pedigrees from Sindhi population of Pakistan. <i>Legal Medicine</i> , 2018, 34, 17-20.	0.6	12
12	Population data of 23 Y STRs from Manchu population of Liaoning Province, Northeast China. <i>International Journal of Legal Medicine</i> , 2019, 133, 785-788.	1.2	12
13	Genetic analysis of 12 X-STRs for forensic purposes in Liaoning Manchu population from China. <i>Gene</i> , 2019, 683, 153-158.	1.0	10
14	Title: Developmental validation of Y-SNP pedigree tagging system: A panel via quick ARMS PCR. <i>Forensic Science International: Genetics</i> , 2020, 46, 102271.	1.6	10
15	mtDNA sequence diversity of Hazara ethnic group from Pakistan. <i>Forensic Science International: Genetics</i> , 2017, 30, e1-e5.	1.6	8
16	Association analysis of <i>NUCKS1</i> and <i>INPP5K</i> polymorphism with Parkinson's disease. <i>Genes and Genetic Systems</i> , 2018, 93, 59-64.	0.2	8
17	Forensic features and genetic legacy of the Baloch population of Pakistan and the Hazara population across Durand line revealed by Y-chromosomal STRs. <i>International Journal of Legal Medicine</i> , 2021, 135, 1777-1784.	1.2	7
18	No association between polymorphisms in the promoter region of dopamine receptor D2 gene and schizophrenia in the northern Chinese Han population: A case-control study. <i>Brain and Behavior</i> , 2019, 9, e01193.	1.0	6

#	ARTICLE	IF	CITATIONS
19	Systematic sequence analysis of the FUT3 gene identifies 11 novel alleles in the Sindhi and Punjabi populations from Pakistan. <i>Scientific Reports</i> , 2020, 10, 5543.	1.6	6
20	Evaluation of 13 rapidly mutating Y-STRs in endogamous Punjabi and Sindhi ethnic groups from Pakistan. <i>International Journal of Legal Medicine</i> , 2019, 133, 799-802.	1.2	4
21	Phylogenetic relationship and genetic history of Central Asian Kazakhs inferred from Y-chromosome and autosomal variations. <i>Molecular Genetics and Genomics</i> , 2020, 295, 221-231.	1.0	4
22	The Heart of Silk Road "Xinjiang," Its Genetic Portray, and Forensic Parameters Inferred From Autosomal STRs. <i>Frontiers in Genetics</i> , 2021, 12, 760760.	1.1	4
23	Population background exploration and genetic distribution analysis of Pakistan Hazara via 23 autosomal STRs. <i>Annals of Human Biology</i> , 2019, 46, 514-518.	0.4	3
24	Genetic structure and forensic characteristics of Saraiki population from Southern Punjab, Pakistan, revealed by 20 Y-chromosomal STRs. <i>International Journal of Legal Medicine</i> , 2020, 134, 977-979.	1.2	3
25	Genetic characterization of 15 autosomal STRs in the interior Sindhi population of Pakistan and their phylogenetic relationship with other populations. <i>International Journal of Immunogenetics</i> , 2020, 47, 149-157.	0.8	2
26	Are Roma People Descended from the Punjab Region of Pakistan: A Y-Chromosomal Perspective. <i>Genes</i> , 2022, 13, 532.	1.0	2
27	Genetic polymorphisms of 20 autosomal STR loci in the Han population of Putian City, Southeastern China. <i>Annals of Human Biology</i> , 2019, 46, 509-513.	0.4	1
28	Population genetics of 19 Y-STR loci in Yanbian Korean samples from China. <i>Annals of Human Genetics</i> , 2019, 83, 134-140.	0.3	1
29	Genetic structure and forensic characteristics of the Korean population revealed by GoldenEye 20A. <i>Annals of Human Biology</i> , 2020, 47, 560-563.	0.4	1
30	Assessing the viability of carious lesions in human identification using STR typing. <i>Science Progress</i> , 2021, 104, 003685042110080.	1.0	1
31	Editorial: Role of Y Chromosome in Molecular Anthropology, Forensics, and Genetic Genealogy. <i>Frontiers in Genetics</i> , 0, 13, .	1.1	1
32	Genetic characterisation and forensic importance of 20 Y-STRs in Han population from Anshan, Northeast of China. <i>Annals of Human Biology</i> , 2020, 47, 478-482.	0.4	0
33	Genetic characteristics and forensic features of Xibe ethnic group revealed via extended set of Y-STRs. <i>Annals of Human Biology</i> , 0, , 1-17.	0.4	0