

# Rashed H Alghafri

## List of Publications by Year in descending order

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Version: 2024-02-01

35  
papers

364  
citations

1162367

8  
h-index

839053

18  
g-index

36  
all docs

36  
docs citations

36  
times ranked

564  
citing authors

#	ARTICLE	IF	CITATIONS
1	Metagenomic Sequencing and Reverse Transcriptase PCR Reveal That Mobile Phones and Environmental Surfaces Are Reservoirs of Multidrug-Resistant Superbugs and SARS-CoV-2. <i>Frontiers in Cellular and Infection Microbiology</i> , 2022, 12, 806077.	1.8	6
2	Mobile phones are hazardous microbial platforms warranting robust public health and biosecurity protocols. <i>Scientific Reports</i> , 2022, 12, .	1.6	7
3	First confirmed detection of SARS-COV-2 in untreated municipal and aircraft wastewater in Dubai, UAE: The use of wastewater based epidemiology as an early warning tool to monitor the prevalence of COVID-19. <i>Science of the Total Environment</i> , 2021, 760, 143350.	3.9	97
4	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
5	Mobile phones of paediatric hospital staff are never cleaned and commonly used in toilets with implications for healthcare nosocomial diseases. <i>Scientific Reports</i> , 2021, 11, 12999.	1.6	16
6	A pilot metagenomic study reveals that community derived mobile phones are reservoirs of viable pathogenic microbes. <i>Scientific Reports</i> , 2021, 11, 14102.	1.6	10
7	The role of mobile phones as a possible pathway for pathogen movement, a cross-sectional microbial analysis. <i>Travel Medicine and Infectious Disease</i> , 2021, 43, 102095.	1.5	9
8	Study of 27 Y-STR markers in United Arab Emirates population. <i>Forensic Science International: Reports</i> , 2020, 2, 100057.	0.4	4
9	Mobile phones represent a pathway for microbial transmission: A scoping review. <i>Travel Medicine and Infectious Disease</i> , 2020, 35, 101704.	1.5	58
10	Y Chromosome Short Tandem Repeats Typing. , 2020, , 277-300.		0
11	Forensic and population genetic analysis of Serbian population using 21 STR loci of GlobalFilerâ„¢ PCR amplification kit. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 47-49.	0.1	0
12	Forensic evaluation of VeriFilerâ„¢ Plus 6-dye chemistry kit composed of 23 loci with casework samples. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 892-896.	0.1	3
13	Population genetics data of 23 autosomal STR loci for three Populations in United Arab Emirates. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 187-188.	0.1	1
14	STRmixâ„¢ collaborative exercise on DNA mixture interpretation. <i>Forensic Science International: Genetics</i> , 2019, 40, 1-8.	1.6	39
15	Evaluation of rapidly mutating Y-STRs in Pakistani population. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 245-247.	0.1	0
16	Whole mtGenome analysis in United Arab Emirates populations. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 408-410.	0.1	0
17	Evaluation of 13 rapidly mutating Y-STRs on a Dravidian pedigree. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 216-217.	0.1	0
18	Investigation on rapidly mutating Y-STRs multiplex in Indian population: A pilot study. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 805-806.	0.1	0

#	ARTICLE	IF	CITATIONS
19	DNA typing from skeletal remains using GlobalFiler <sup>®</sup> , <sup>®</sup> PCR amplification and Investigator <sup>®</sup> 24plex QS kits. Forensic Science International: Genetics Supplement Series, 2019, 7, 50-52.	0.1	0
20	Rapidly mutating Y-STR analyses of compromised forensic samples. International Journal of Legal Medicine, 2018, 132, 397-403.	1.2	9
21	Reduced volume for direct PCR amplification of blood reference samples using Identifiler <sup>®</sup> Direct and GlobalFiler <sup>®</sup> , <sup>®</sup> Express assays. Forensic Science International: Genetics Supplement Series, 2017, 6, e340-e341.	0.1	2
22	Mutation rate at 13 rapidly mutating Y-STR loci in the population of Serbia. Forensic Science International: Genetics Supplement Series, 2017, 6, e377-e379.	0.1	5
23	Rapidly mutating Y-STRs population data in the population of Serbia and haplotype probability assessment for forensic purposes. Forensic Science International: Genetics Supplement Series, 2017, 6, e383-e384.	0.1	3
24	DNA analysis from human skeletal remains in forensic casework. Forensic Science International: Genetics Supplement Series, 2017, 6, e342-e345.	0.1	6
25	Genetic characterization of 27 Y-STR loci with the Yfiler <sup>®</sup> Plus kit in the population of Serbia. Forensic Science International: Genetics, 2017, 31, e48-e49.	1.6	13
26	Y-chromosome polymorphisms in the United Arab Emirates population. Forensic Science International: Genetics Supplement Series, 2017, 6, e397-e398.	0.1	0
27	Rapid amplification of the RM <sup>®</sup> Yplex assay. Electrophoresis, 2016, 37, 2817-2821.	1.3	5
28	Population data for SE33 locus in United Arab Emirates Arab population. Forensic Science International: Genetics Supplement Series, 2015, 5, e238-e239.	0.1	2
29	An evaluation of rapidly mutating Y-STR multi-allelic markers. Forensic Science International: Genetics Supplement Series, 2015, 5, e647-e649.	0.1	1
30	A comparison between Yfiler <sup>®</sup> and RM Y-STRs in United Arab Emirates population. Forensic Science International: Genetics Supplement Series, 2015, 5, e650-e652.	0.1	0
31	An evaluation of miniSTR markers for casework applications. Forensic Science International: Genetics Supplement Series, 2015, 5, e512-e514.	0.1	0
32	Development and validation of an allelic frequency database for Qatari population using 13 rapidly mutating Y-STRs multiplex assay. Forensic Science International: Genetics Supplement Series, 2015, 5, e365-e367.	0.1	0
33	A novel multiplex assay for simultaneously analysing 13 rapidly mutating Y-STRs. Forensic Science International: Genetics, 2015, 17, 91-98.	1.6	55
34	An investigation of 21 insertion deletion markers in United Arab Emirates population. Forensic Science International: Genetics Supplement Series, 2015, 5, e261-e263.	0.1	0
35	Rapidly mutating Y-STRs multiplex genotyping panel to investigate UAE population. Forensic Science International: Genetics Supplement Series, 2013, 4, e200-e201.	0.1	6