

# Mauro Pesaresi

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/345090/publications.pdf>

Version: 2024-02-01

18  
papers

241  
citations

1162367

8  
h-index

996533

15  
g-index

18  
all docs

18  
docs citations

18  
times ranked

322  
citing authors

#	ARTICLE	IF	CITATIONS
1	Evaluation of a microhaplotypes panel for forensic genetics using massive parallel sequencing technology. <i>Forensic Science International: Genetics</i> , 2019, 41, 120-127.	1.6	57
2	Development of multiplex PCRs for evolutionary and forensic applications of 37 human Y chromosome SNPs. <i>Forensic Science International</i> , 2006, 157, 23-35.	1.3	55
3	Y-chromosome genetic structure in sub-Apennine populations of Central Italy by SNP and STR analysis. <i>International Journal of Legal Medicine</i> , 2007, 121, 234-237.	1.2	33
4	Visceral fat inflammation and fat embolism are associated with lung's lipidic hyaline membranes in subjects with COVID-19. <i>International Journal of Obesity</i> , 2022, 46, 1009-1017.	1.6	22
5	Identification and functional analysis of a new putative caveolin-3 variant found in a patient with sudden unexplained death. <i>Journal of Biomedical Science</i> , 2014, 21, 58.	2.6	11
6	Multiplex PCR Development of Y-chromosomal Biallelic Polymorphisms for Forensic Application. <i>Journal of Forensic Sciences</i> , 2005, 50, 1-7.	0.9	11
7	Investigation on genetic thrombophilic factors in FFPE autopsy tissue from subjects who died from pulmonary embolism. <i>International Journal of Legal Medicine</i> , 2017, 131, 447-458.	1.2	10
8	Y-chromosome markers distribution in Northern Africa: High-resolution SNP and STR analysis in Tunisia and Morocco populations. <i>Forensic Science International: Genetics Supplement Series</i> , 2008, 1, 235-236.	0.1	8
9	A microhaplotypes panel for forensic genetics using massive parallel sequencing. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e117-e118.	0.1	7
10	Searching the undetected mtDNA variants in forensic MPS data. <i>Forensic Science International: Genetics</i> , 2020, 49, 102399.	1.6	7
11	Post-mortem DNA damage: A comparative study of STRs and SNPs typing efficiency in simulated forensic samples. <i>International Congress Series</i> , 2006, 1288, 510-512.	0.2	5
12	D16S539 microvariant or D2S1338 off-ladder allele? A case report about a range overlapping between two loci. <i>Forensic Science International: Genetics Supplement Series</i> , 2008, 1, 123-124.	0.1	5
13	Performance of a massive parallel sequencing microhaplotypes assay on degraded DNA. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 782-783.	0.1	5
14	Multiplex genotyping of 22 autosomal SNPs and its application in the forensic field. <i>International Congress Series</i> , 2006, 1288, 40-42.	0.2	2
15	Multiplex PCR development of Y-chromosomal biallelic polymorphisms for forensic application. <i>Journal of Forensic Sciences</i> , 2005, 50, 519-25.	0.9	2
16	Past, Present and Future in Forensic Human Identification. , 2020, , 81-92.		1
17	Y-chromosome genetic structure in a sub-Apennine population of the Marches (central Italy): Analysis by SNP and STR polymorphisms. <i>International Congress Series</i> , 2006, 1288, 168-170.	0.2	0
18	Exploring the usefulness of microhaplotypes in forensic identification using massive parallel sequencing technology. <i>Minerva Medicolegale; Archivio Di Antropologia Criminale, Psichiatria, E Medicina Legale</i> , 2020, 140, .	0.0	0