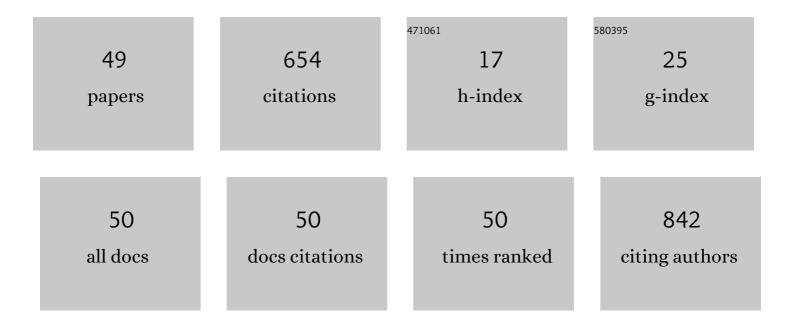
## Chiara Turchi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2017821/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Circulating SARS-CoV-2 variants in Italy, October 2020–March 2021. Virology Journal, 2021, 18, 168.	1.4	36
2	Searching the undetected mtDNA variants in forensic MPS data. Forensic Science International: Genetics, 2020, 49, 102399.	1.6	7
3	mtDNA exploitation in forensics. , 2020, , 145-169.		0
4	Evaluation of the Ion AmpliSeq SARS-CoV-2 Research Panel by Massive Parallel Sequencing. Genes, 2020, 11, 929.	1.0	27
5	Assessment of the Precision ID Identity Panel kit on challenging forensic samples. Forensic Science International: Genetics, 2020, 49, 102400.	1.6	19
6	Past, Present and Future in Forensic Human Identification. , 2020, , 81-92.		1
7	Exploring the usefulness of microhaplotypes in forensic identification using massive parallel sequencing technology. Minerva Medicolegale; Archivio Di Antropologia Criminale, Psichiatria, E Medicina Legale, 2020, 140, .	0.0	0
8	Development of a forensic DNA phenotyping panel using massive parallel sequencing. Forensic Science International: Genetics Supplement Series, 2019, 7, 177-179.	0.1	3
9	Evaluation of a microhaplotypes panel for forensic genetics using massive parallel sequencing technology. Forensic Science International: Genetics, 2019, 41, 120-127.	1.6	57
10	Dealing with low amounts of degraded DNA: Evaluation of SNP typing of challenging forensic samples by using massive parallel sequencing. Forensic Science International: Genetics Supplement Series, 2019, 7, 83-84.	0.1	3
11	Performance of a massive parallel sequencing microhaplotypes assay on degraded DNA. Forensic Science International: Genetics Supplement Series, 2019, 7, 782-783.	0.1	5
12	Analysis of recombination and mutation events for 12 X-Chr STR loci: A collaborative family study of the Italian Speaking Working Group Ge.F.I. Forensic Science International: Genetics Supplement Series, 2019, 7, 398-400.	0.1	5
13	Massive parallel sequencing and osteogenesis imperfecta: An essential tool for forensic investigation over child abuse. Forensic Science International: Genetics Supplement Series, 2019, 7, 103-104.	0.1	0
14	A microhaplotypes panel for forensic genetics using massive parallel sequencing. Forensic Science International: Genetics Supplement Series, 2017, 6, e117-e118.	0.1	7
15	The mitochondrial DNA makeup of Romanians: A forensic mtDNA control region database and phylogenetic characterization. Forensic Science International: Genetics, 2016, 24, 136-142.	1.6	20
16	The molecular characterization of a depurinated trial DNA sample can be a model to understand the reliability of the results in forensic genetics. Electrophoresis, 2014, 35, 3134-3144.	1.3	12
17	Heroin addictions in Italians: Evaluation of OPRM1 genetic variants by case–control association study. Forensic Science International: Genetics Supplement Series, 2013, 4, e57-e58.	0.1	1
18	Role of 5-HTTLPR Polymorphism in the Development of the Inward/Outward Personality Organization: A Genetic Association Study. PLoS ONE, 2013, 8, e82192.	1.1	8

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19	ADH4 intronic variations are associated with alcohol dependence. Pharmacogenetics and Genomics, 2012, 22, 79-94.	0.7	7
20	HTR2A gene polymorphisms and Inward and Outward Personal Meaning Organisations. Acta Neuropsychiatrica, 2012, 24, 336-343.	1.0	6
21	Genetic factors in inward vs outward personality Organizations: focus on HTR2A polymorphisms. Quaderni Italiani Di Psichiatria, 2011, 30, 83-88.	0.1	2
22	An overview of the genetic susceptibility to alcoholism. Medicine, Science and the Law, 2011, 51, 2-6.	0.6	25
23	Searching for a relationship between the serotonin receptor 2A gene variations and the development of Inward and Outward Personal Meaning Organizations. Psychiatric Genetics, 2011, 21, 269-270.	0.6	6
24	GABRA2 and Alcohol Use Disorders: No Evidence of an Association in an Italian Case–Control Study. Alcoholism: Clinical and Experimental Research, 2010, 34, 659-668.	1.4	19
25	Polymorphisms of mtDNA control region in Tunisian and Moroccan populations: An enrichment of forensic mtDNA databases with Northern Africa data. Forensic Science International: Genetics, 2009, 3, 166-172.	1.6	27
26	Multiplex mtDNA coding region SNP assays for molecular dissection of haplogroups U/K and J/T. Forensic Science International: Genetics, 2009, 4, 21-25.	1.6	20
27	Genetic susceptibility for addiction: Searching of risk loci for the widespread drugs of abuse. Forensic Science International: Genetics Supplement Series, 2009, 2, 487-488.	0.1	1
28	Italian mitochondrial DNA database: results of a collaborative exercise and proficiency testing. International Journal of Legal Medicine, 2008, 122, 199-204.	1.2	48
29	Y-chromosome markers distribution in Northern Africa: High-resolution SNP and STR analysis in Tunisia and Morocco populations. Forensic Science International: Genetics Supplement Series, 2008, 1, 235-236.	0.1	8
30	D16S539 microvariant or D2S1338 off-ladder allele? A case report about a range overlapping between two loci. Forensic Science International: Genetics Supplement Series, 2008, 1, 123-124.	0.1	5
31	Population data for D10S1248, D14S1434, and D22S1045 miniSTRs loci from the Marches region (Central) Tj	ETQg110	.784314 rgB1
32	Association of genetic variations in alcohol dehydrogenase 4 with alcohol dependence in Italian population sample. Forensic Science International: Genetics Supplement Series, 2008, 1, 580-581.	0.1	0
33	MtDNA analysis for genetic identification of forensically important insects. Forensic Science International: Genetics Supplement Series, 2008, 1, 584-585.	0.1	24
34	A missense germline mutation in exon 7 of the MSH2 gene in a HNPCC family from center-Italy. Familial Cancer, 2007, 6, 97-102.	0.9	4
35	Y-chromosome genetic structure in sub-Apennine populations of Central Italy by SNP and STR analysis. International Journal of Legal Medicine, 2007, 121, 234-237.	1.2	33
36	Multiplex genotyping of 22 autosomal SNPs and its application in the forensic field. International Congress Series, 2006, 1288, 40-42.	0.2	2

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37	Post-mortem DNA damage: A comparative study of STRs and SNPs typing efficiency in simulated forensic samples. International Congress Series, 2006, 1288, 510-512.	0.2	5
38	Y-chromosome genetic structure in a sub-Apennine population of the Marches (central Italy): Analysis by SNP and STR polymorphisms. International Congress Series, 2006, 1288, 168-170.	0.2	0
39	Development of multiplex PCRs for evolutionary and forensic applications of 37 human Y chromosome SNPs. Forensic Science International, 2006, 157, 23-35.	1.3	55
40	Subtyping mtDNA haplogroup H by SNaPshot minisequencing and its application in forensic individual identification. International Journal of Legal Medicine, 2006, 120, 151-156.	1.2	36
41	Development of a heptaplex PCR system to analyse X-chromosome STR loci from five Italian population samples. Forensic Science International, 2005, 153, 231-236.	1.3	26
42	Multiplex PCR Development of Y-chromosomal Biallelic Polymorphisms for Forensic Application. Journal of Forensic Sciences, 2005, 50, 1-7.	0.9	11
43	Multiplex PCR development of Y-chromosomal biallelic polymorphisms for forensic application. Journal of Forensic Sciences, 2005, 50, 519-25.	0.9	2
44	Development and forensic applications of multiplex PCR of autosomal biallele polymorphisms. International Congress Series, 2004, 1261, 213-215.	0.2	2
45	Development of a heptaplex PCR system to analyse X-chromosome STR loci from five Italian population samples. A collaborative study. International Congress Series, 2004, 1261, 272-274.	0.2	3
46	Occurrence of heteroplasmy in related individuals. International Congress Series, 2003, 1239, 553-556.	0.2	1
47	A multicentric study of SE33 allele frequencies in the Italian population. International Congress Series, 2003, 1239, 83-86.	0.2	Ο
48	Fingerprints as evidence for a genetic profile: morphological study on fingerprints and analysis of exogenous and individual factors affecting DNA typing. Journal of Forensic Sciences, 2003, 48, 586-92.	0.9	18
49	Polymorphism of the mitochondrial DNA control region in Italians. International Journal of Legal Medicine, 2001, 114, 224-228.	1.2	45