## Valerio Onofri

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A global analysis of Y-chromosomal haplotype diversity for 23 STR loci. Forensic Science International: Genetics, 2014, 12, 12-23.	1.6	214
2	Toward Male Individualization with Rapidly Mutating Y-Chromosomal Short Tandem Repeats. Human Mutation, 2014, 35, 1021-1032.	1.1	151
3	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
4	J1-M267 Y lineage marks climate-driven pre-historical human displacements. European Journal of Human Genetics, 2009, 17, 1520-1524.	1.4	54
5	Y chromosome genetic variation in the Italian peninsula is clinal and supports an admixture model for the Mesolithic–Neolithic encounter. Molecular Phylogenetics and Evolution, 2007, 44, 228-239.	1.2	49
6	Moors and Saracens in Europe: estimating the medieval North African male legacy in southern Europe. European Journal of Human Genetics, 2009, 17, 848-852.	1.4	37
7	Development of an Italian RM Y-STR haplotype database: Results of the 2013 GEFI collaborative exercise. Forensic Science International: Genetics, 2015, 15, 56-63.	1.6	35
8	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
9	The Etruscan timeline: a recent Anatolian connection. European Journal of Human Genetics, 2009, 17, 693-696.	1.4	32
10	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
11	MtDNA analysis for genetic identification of forensically important insects. Forensic Science International: Genetics Supplement Series, 2008, 1, 584-585.	0.1	24
12	Molecular characterisation and population genetics of the DYS458 .2 allelic variant. Forensic Science International: Genetics Supplement Series, 2008, 1, 203-205.	0.1	20
13	Assessment of the Precision ID Identity Panel kit on challenging forensic samples. Forensic Science International: Genetics, 2020, 49, 102400.	1.6	19
14	Multiplex PCR Development of Y-chromosomal Biallelic Polymorphisms for Forensic Application. Journal of Forensic Sciences, 2005, 50, 1-7.	0.9	11
15	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
16	Analysis of uni and bi-parental markers in mixture samples: Lessons from the 22nd GHEP-ISFG Intercomparison Exercise. Forensic Science International: Genetics, 2016, 25, 63-72.	1.6	7
17	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
18	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

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19	Unusual Association of Three Rare Alleles and a Mismatch in a Case of Paternity Testing. Journal of Forensic Sciences, 2004, 49, 1-3.	0.9	5
20	Microgeographic variation of Y-chromosome haplotypes in Italy. Forensic Science International: Genetics Supplement Series, 2008, 1, 239-241.	0.1	4
21	Evaluating Y-chromosome STRs mutation rates: A collaborative study of the Ge.F.IISFG Italian Group. Forensic Science International: Genetics Supplement Series, 2009, 2, 419-420.	0.1	3
22	X-chromosome in Italy: A database of 29 STR markers. Forensic Science International: Genetics Supplement Series, 2011, 3, e37-e38.	0.1	3
23	Development of a forensic DNA phenotyping panel using massive parallel sequencing. Forensic Science International: Genetics Supplement Series, 2019, 7, 177-179.	0.1	3
24	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
25	Multiplex genotyping of 22 autosomal SNPs and its application in the forensic field. International Congress Series, 2006, 1288, 40-42.	0.2	2
26	Population data for D10S1248, D14S1434, and D22S1045 miniSTRs loci from the Marches region (Central) Tj E	ГQ <sub>8</sub> 000г	gBT /Overloc
27	The Ge.F.I. DNA Proficiency Test: Year-one experience. Forensic Science International: Genetics Supplement Series, 2013, 4, e27-e28.	0.1	2

	Supplement Series, 2013, 4, e27-e28.		
28	Multiplex PCR development of Y-chromosomal biallelic polymorphisms for forensic application. Journal of Forensic Sciences, 2005, 50, 519-25.	0.9	2
29	Past, Present and Future in Forensic Human Identification. , 2020, , 81-92.		1
30	Multiplex PCR development of Y-chromosomal biallelic polymorphisms for forensic applications. International Congress Series, 2004, 1261, 363-365.	0.2	0
31	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
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	Y chromosome J2 subtyping in an Italian sample: Population and forensic implications. Forensic Science International: Genetics Supplement Series, 2008, 1, 233-234.	0.1	0
34	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