

# Marion Nagy

## List of Publications by Year in descending order

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Version: 2024-02-01

26  
papers

1,700  
citations

687363

13  
h-index

580821

25  
g-index

26  
all docs

26  
docs citations

26  
times ranked

2316  
citing authors

#	ARTICLE	IF	CITATIONS
1	Sarcoidosis is associated with a truncating splice site mutation in BTNL2. <i>Nature Genetics</i> , 2005, 37, 357-364.	21.4	451
2	Characteristics and Frequency of Germline Mutations at Microsatellite Loci from the Human Y Chromosome, as Revealed by Direct Observation in Father/Son Pairs. <i>American Journal of Human Genetics</i> , 2000, 66, 1580-1588.	6.2	334
3	A global analysis of Y-chromosomal haplotype diversity for 23 STR loci. <i>Forensic Science International: Genetics</i> , 2014, 12, 12-23.	3.1	214
4	Analysis of molecular variance (AMOVA) of Y-chromosome-specific microsatellites in two closely related human populations [published erratum appears in <i>Hum Mol Genet</i> 1997 May;6(5):828]. <i>Human Molecular Genetics</i> , 1996, 5, 1029-1033.	2.9	173
5	Continent-Wide Decoupling of Y-Chromosomal Genetic Variation from Language and Geography in Native South Americans. <i>PLoS Genetics</i> , 2013, 9, e1003460.	3.5	89
6	Applications of microsatellite-based Y chromosome haplotyping. <i>Electrophoresis</i> , 1997, 18, 1602-1607.	2.4	63
7	Comparison of Chimerism and Minimal Residual Disease Monitoring for Relapse Prediction after Allogeneic Stem Cell Transplantation for Adult Acute Lymphoblastic Leukemia. <i>Biology of Blood and Marrow Transplantation</i> , 2014, 20, 1522-1529.	2.0	57
8	Analysis of Y chromosome STR haplotypes in the European part of Russia reveals high diversities but non-significant genetic distances between populations. <i>International Journal of Legal Medicine</i> , 2008, 122, 219-223.	2.2	53
9	Graft versus host disease after liver transplantation – a single center experience and review of literature. <i>Transplant International</i> , 2008, 21, 441-451.	1.6	47
10	Validation of a combined autosomal/Y-chromosomal STR approach for analyzing typical biological stains in sexual-assault cases. <i>Forensic Science International: Genetics</i> , 2015, 19, 238-242.	3.1	47
11	Hierarchical Y-SNP assay to study the hidden diversity and phylogenetic relationship of native populations in South America. <i>Forensic Science International: Genetics</i> , 2011, 5, 100-104.	3.1	36
12	Y-chromosomal STR haplotypes in Kalmyk population samples. <i>Forensic Science International</i> , 2007, 173, 204-209.	2.2	31
13	Comparison of two silica-based extraction methods for DNA isolation from bones. <i>Legal Medicine</i> , 2016, 22, 36-41.	1.3	22
14	Evaluation of whole-genome amplification of low-copy-number DNA in chimerism analysis after allogeneic stem cell transplantation using STR marker typing. <i>Electrophoresis</i> , 2006, 27, 3028-3037.	2.4	12
15	Results of a collaborative study on DNA identification of aged bone samples. <i>Croatian Medical Journal</i> , 2017, 58, 203-213.	0.7	12
16	Individual specific extraction of DNA from male mixtures – First evaluation studies. <i>Forensic Science International: Genetics</i> , 2011, 5, 117-121.	3.1	10
17	Investigation of HLA-DPA1 genotypes as predictors of inflammatory bowel disease in the German, South African, and South Korean populations. <i>International Journal of Colorectal Disease</i> , 2002, 17, 238-244.	2.2	9
18	Genetic research at a fivefold children's burial from medieval Berlin. <i>Forensic Science International: Genetics</i> , 2015, 15, 90-97.	3.1	9

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19	Separation of Y-chromosomal haplotypes from male DNA mixtures via multiplex haplotype-specific extraction. <i>Forensic Science International: Genetics</i> , 2015, 19, 223-231.	3.1	7
20	Nonmyeloablative stem cell transplantation in patients with ALL and AML results in low nonrelapse mortality despite high rate of infections and GVHD. <i>The Hematology Journal</i> , 2004, 5, 395-402.	1.4	6
21	New Prediction Model for Probe Specificity in an Allele-Specific Extension Reaction for Haplotype-Specific Extraction (HSE) of Y Chromosome Mixtures. <i>PLoS ONE</i> , 2012, 7, e45955.	2.5	6
22	Nonelectrophoretic method for high-throughput HLA-DRB1 group genotyping. <i>BioTechniques</i> , 2004, 36, 148-151.	1.8	4
23	Validation of haplotype-specific extraction for separating a mitochondrial DNA model mixture and application to simulated casework. <i>Forensic Science International: Genetics</i> , 2018, 35, 57-64.	3.1	4
24	Genetic and isotope analysis of a triple burial from medieval St. Peter's cemetery in Cölln/Berlin. <i>Forensic Science International: Genetics</i> , 2022, 59, 102718.	3.1	3
25	Strategies for excluding false Y-chromosomal SNP entries from human genome databases. <i>Electrophoresis</i> , 2012, 33, 1488-1491.	2.4	1
26	SNP-Array Karyotyping Reveals the Presence of Previously Cryptic Clonal Chromosomal Aberrations Including Segmental UPD in Patients with Fanconi Anemia. <i>Blood</i> , 2007, 110, 1678-1678.	1.4	0