

Lareu Mv

List of Publications by Year in descending order

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135
papers

6,490
citations

50170

46
h-index

76769

74
g-index

136
all docs

136
docs citations

136
times ranked

4136
citing authors

#	ARTICLE	IF	CITATIONS
1	The Making of the African mtDNA Landscape. <i>American Journal of Human Genetics</i> , 2002, 71, 1082-1111.	2.6	451
2	An overview of STRUCTURE: applications, parameter settings, and supporting software. <i>Frontiers in Genetics</i> , 2013, 4, 98.	1.1	432
3	Inferring ancestral origin using a single multiplex assay of ancestry-informative marker SNPs. <i>Forensic Science International: Genetics</i> , 2007, 1, 273-280.	1.6	332
4	The African Diaspora: Mitochondrial DNA and the Atlantic Slave Trade. <i>American Journal of Human Genetics</i> , 2004, 74, 454-465.	2.6	213
5	Typing of mitochondrial DNA coding region SNPs of forensic and anthropological interest using SNaPshot minisequencing. <i>Forensic Science International</i> , 2004, 140, 251-257.	1.3	161
6	mtDNA analysis of the Galician population: a genetic edge of European variation. <i>European Journal of Human Genetics</i> , 1998, 6, 365-375.	1.4	141
7	Development of a methylation marker set for forensic age estimation using analysis of public methylation data and the Agena Bioscience EpiTYPER system. <i>Forensic Science International: Genetics</i> , 2016, 24, 65-74.	1.6	127
8	Further development of forensic eye color predictive tests. <i>Forensic Science International: Genetics</i> , 2013, 7, 28-40.	1.6	119
9	Building a forensic ancestry panel from the ground up: The EUROFORGEN Global AIM-SNP set. <i>Forensic Science International: Genetics</i> , 2014, 11, 13-25.	1.6	116
10	Ancestry Analysis in the 11-M Madrid Bomb Attack Investigation. <i>PLoS ONE</i> , 2009, 4, e6583.	1.1	110
11	Report of the European DNA profiling group (EDNAP) "towards standardisation of short tandem repeat (STR) loci. <i>Forensic Science International</i> , 1994, 65, 51-59.	1.3	109
12	Inter-laboratory evaluation of SNP-based forensic identification by massively parallel sequencing using the Ion PGM. <i>Forensic Science International: Genetics</i> , 2015, 17, 110-121.	1.6	105
13	Analysis of global variability in 15 established and 5 new European Standard Set (ESS) STRs using the CEPH human genome diversity panel. <i>Forensic Science International: Genetics</i> , 2011, 5, 155-169.	1.6	103
14	Revision of the SNPforID 34-plex forensic ancestry test: Assay enhancements, standard reference sample genotypes and extended population studies. <i>Forensic Science International: Genetics</i> , 2013, 7, 63-74.	1.6	102
15	Eurasiaplex: A forensic SNP assay for differentiating European and South Asian ancestries. <i>Forensic Science International: Genetics</i> , 2013, 7, 359-366.	1.6	102
16	Resolving relationship tests that show ambiguous STR results using autosomal SNPs as supplementary markers. <i>Forensic Science International: Genetics</i> , 2008, 2, 198-204.	1.6	100
17	Effect of environmental factors on PCR-DNA analysis from dental pulp. <i>International Journal of Legal Medicine</i> , 1996, 109, 125-129.	1.2	92
18	Hierarchical analysis of 30 Y-chromosome SNPs in European populations. <i>International Journal of Legal Medicine</i> , 2005, 119, 10-15.	1.2	92

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19	A SNaPshot of next generation sequencing for forensic SNP analysis. <i>Forensic Science International: Genetics</i> , 2015, 14, 50-60.	1.6	85
20	A new SNP assay for identification of highly degraded human DNA. <i>Forensic Science International: Genetics</i> , 2012, 6, 341-349.	1.6	82
21	Heteroplasmy in mtDNA and the weight of evidence in forensic mtDNA analysis: a case report. <i>International Journal of Legal Medicine</i> , 2001, 114, 186-190.	1.2	75
22	New Population and Phylogenetic Features of the Internal Variation within Mitochondrial DNA Macro-Haplogroup R0. <i>PLoS ONE</i> , 2009, 4, e5112.	1.1	75
23	A highly variable STR at the D12S391 locus. <i>International Journal of Legal Medicine</i> , 1996, 109, 134-138.	1.2	74
24	The recombination landscape around forensic STRs: Accurate measurement of genetic distances between syntenic STR pairs using HapMap high density SNP data. <i>Forensic Science International: Genetics</i> , 2012, 6, 354-365.	1.6	73
25	Investigation of the STR locus HUMTH01 using PCR and two electrophoresis formats: UK and Galician Caucasian population surveys and usefulness in paternity investigations. <i>Forensic Science International</i> , 1994, 66, 41-52.	1.3	70
26	Robustness of the Y STRs DYS19, DYS389 I and II, DYS390 and DYS393: optimization of a PCR pentaplex. <i>Forensic Science International</i> , 1999, 106, 163-172.	1.3	70
27	Forensic performance of two insertion-deletion marker assays. <i>International Journal of Legal Medicine</i> , 2012, 126, 725-737.	1.2	70
28	Building a custom large-scale panel of novel microhaplotypes for forensic identification using MiSeq and Ion S5 massively parallel sequencing systems. <i>Forensic Science International: Genetics</i> , 2020, 45, 102213.	1.6	70
29	Development of a forensic skin colour predictive test. <i>Forensic Science International: Genetics</i> , 2014, 13, 34-44.	1.6	69
30	Case report: Identification of skeletal remains using short-amplicon marker analysis of severely degraded DNA extracted from a decomposed and charred femur. <i>Forensic Science International: Genetics</i> , 2008, 2, 212-218.	1.6	66
31	Inter-laboratory evaluation of the EUROFORGEN Global ancestry-informative SNP panel by massively parallel sequencing using the Ion PGM. <i>Forensic Science International: Genetics</i> , 2016, 23, 178-189.	1.6	65
32	Report on the second EDNAP collaborative STR exercise. <i>Forensic Science International</i> , 1995, 71, 137-152.	1.3	64
33	Sequence variation of a hypervariable short tandem repeat at the D1S1656 locus. <i>International Journal of Legal Medicine</i> , 1998, 111, 244-247.	1.2	63
34	MAPlex - A massively parallel sequencing ancestry analysis multiplex for Asia-Pacific populations. <i>Forensic Science International: Genetics</i> , 2019, 42, 213-226.	1.6	63
35	Typing short amplicon binary polymorphisms: Supplementary SNP and Indel genetic information in the analysis of highly degraded skeletal remains. <i>Forensic Science International: Genetics</i> , 2012, 6, 469-476.	1.6	60
36	Pacifplex : an ancestry-informative SNP panel centred on Australia and the Pacific region. <i>Forensic Science International: Genetics</i> , 2016, 20, 71-80.	1.6	60

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37	Results of a collaborative study of the EDNAP group regarding mitochondrial DNA heteroplasmy and segregation in hair shafts. <i>Forensic Science International</i> , 2004, 140, 1-11.	1.3	59
38	Ethical-legal problems of DNA databases in criminal investigation. <i>Journal of Medical Ethics</i> , 2000, 26, 266-271.	1.0	57
39	The Global AIMs Nano set: A 31-plex SNaPshot assay of ancestry-informative SNPs. <i>Forensic Science International: Genetics</i> , 2016, 22, 81-88.	1.6	57
40	Performance of ancestry-informative SNP and microhaplotype markers. <i>Forensic Science International: Genetics</i> , 2019, 43, 102141.	1.6	55
41	Shipwrecks and founder effects: Divergent demographic histories reflected in Caribbean mtDNA. <i>American Journal of Physical Anthropology</i> , 2005, 128, 855-860.	2.1	52
42	SNPs as Supplements in Simple Kinship Analysis or as Core Markers in Distant Pairwise Relationship Tests: When Do SNPs Add Value or Replace Well-Established and Powerful STR Tests?. <i>Transfusion Medicine and Hemotherapy</i> , 2012, 39, 202-210.	0.7	52
43	The use of the STRs HUMTH01, HUMVWA31/A, HUMF13A1, HUMFES/FPS, HUMLPL in forensic application: Validation studies and population data for Galicia (NW Spain). <i>International Journal of Legal Medicine</i> , 1995, 107, 283-290.	1.2	51
44	Global patterns of STR sequence variation: Sequencing the CEPH human genome diversity panel for 58 forensic STRs using the Illumina ForenSeq DNA Signature Prep Kit. <i>Electrophoresis</i> , 2018, 39, 2708-2724.	1.3	51
45	Chimpanzee homologous of human Y specific STRs. <i>Forensic Science International</i> , 2002, 126, 129-136.	1.3	50
46	EuroforGen-NoE collaborative exercise on LRmix to demonstrate standardization of the interpretation of complex DNA profiles. <i>Forensic Science International: Genetics</i> , 2014, 9, 47-54.	1.6	50
47	Pharmacogenetics of OATP Transporters Reveals That SLCO1B1 c.388A>G Variant Is Determinant of Increased Atorvastatin Response. <i>International Journal of Molecular Sciences</i> , 2011, 12, 5815-5827.	1.8	49
48	The SNPforID browser: an online tool for query and display of frequency data from the SNPforID project. <i>International Journal of Legal Medicine</i> , 2008, 122, 435-440.	1.2	47
49	Results of the 1999-2000 collaborative exercise and proficiency testing program on mitochondrial DNA of the GEP-ISFG: an inter-laboratory study of the observed variability in the heteroplasmy level of hair from the same donor. <i>Forensic Science International</i> , 2002, 125, 1-7.	1.3	45
50	The mtDNA ancestry of admixed Colombian populations. <i>American Journal of Human Biology</i> , 2008, 20, 584-591.	0.8	44
51	Tetra-allelic SNPs: Informative forensic markers compiled from public whole-genome sequence data. <i>Forensic Science International: Genetics</i> , 2015, 19, 100-106.	1.6	44
52	Distribution of Y-chromosome STR defined haplotypes in Iberia. <i>Forensic Science International</i> , 2000, 110, 117-126.	1.3	43
53	Tracking age-correlated DNA methylation markers in the young. <i>Forensic Science International: Genetics</i> , 2018, 36, 50-59.	1.6	41
54	Y chromosome STR haplotypes: genetic and sequencing data of the Galician population (NW Spain). <i>International Journal of Legal Medicine</i> , 1998, 112, 15-21.	1.2	39

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55	Micro-geographical differentiation in Northern Iberia revealed by Y-chromosomal DNA analysis. <i>Gene</i> , 2004, 329, 17-25.	1.0	38
56	Challenging DNA: Assessment of a range of genotyping approaches for highly degraded forensic samples. <i>Forensic Science International: Genetics Supplement Series</i> , 2008, 1, 26-28.	0.1	38
57	pop.STR® An online population frequency browser for established and new forensic STRs. <i>Forensic Science International: Genetics Supplement Series</i> , 2009, 2, 361-362.	0.1	38
58	Development of a novel forensic STR multiplex for ancestry analysis and extended identity testing. <i>Electrophoresis</i> , 2013, 34, 1151-1162.	1.3	34
59	A compilation of tri-allelic SNPs from 1000 Genomes and use of the most polymorphic loci for a large-scale human identification panel. <i>Forensic Science International: Genetics</i> , 2020, 46, 102232.	1.6	34
60	Sequence structure of 12 novel Y chromosome microsatellites and PCR amplification strategies. <i>Forensic Science International</i> , 2001, 122, 19-26.	1.3	33
61	Evaluation of the Qiagen 140-SNP forensic identification multiplex for massively parallel sequencing. <i>Forensic Science International: Genetics</i> , 2017, 28, 35-43.	1.6	33
62	Exploring iris colour prediction and ancestry inference in admixed populations of South America. <i>Forensic Science International: Genetics</i> , 2014, 13, 3-9.	1.6	32
63	“New turns from old STAs” Enhancing the capabilities of forensic short tandem repeat analysis. <i>Electrophoresis</i> , 2014, 35, 3173-3187.	1.3	31
64	Exploration of SNP variants affecting hair colour prediction in Europeans. <i>International Journal of Legal Medicine</i> , 2015, 129, 963-975.	1.2	31
65	Alternative primers for DYS391 typing: advantages of their application to forensic genetics. <i>Forensic Science International</i> , 2000, 112, 49-57.	1.3	30
66	Insights into Iberian population origins through the construction of highly informative Y-chromosome haplotypes using biallelic markers, STRs, and the MSY1 minisatellite. <i>American Journal of Physical Anthropology</i> , 2003, 122, 147-161.	2.1	30
67	Completion of a worldwide reference panel of samples for an ancestry informative Indel assay. <i>Forensic Science International: Genetics</i> , 2015, 17, 75-80.	1.6	30
68	Rapid and enhanced detection of mitochondrial DNA variation using single-strand conformation analysis of superposed restriction enzyme fragments from polymerase chain reaction-amplified products. <i>Electrophoresis</i> , 1997, 18, 52-54.	1.3	29
69	The use of the LightCycler for the detection of Y chromosome SNPs. <i>Forensic Science International</i> , 2001, 118, 163-168.	1.3	29
70	mtDNA hypervariable region II (HVII) sequences in human evolution studies. <i>European Journal of Human Genetics</i> , 2000, 8, 964-974.	1.4	27
71	Allele frequencies of fifteen STRs in a representative sample of the Italian population. <i>Forensic Science International: Genetics</i> , 2009, 3, e29-e30.	1.6	27
72	Sequence variation of a hypervariable short tandem repeat at the D12S391 locus. <i>Gene</i> , 1996, 182, 151-153.	1.0	26

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73	Failed PCR amplifications of MBP-STR alleles due to polymorphism in the primer annealing region. <i>International Journal of Legal Medicine</i> , 1996, 108, 313-315.	1.2	26
74	Report of the European DNA profiling group (EDNAP): an investigation of the complex STR loci D21S11 and HUMFIBRA (FGA). <i>Forensic Science International</i> , 1997, 86, 25-33.	1.3	25
75	Results of a collaborative study regarding the standardization of the Y-linked STR system DYS385 by the European DNA Profiling (EDNAP) group. <i>Forensic Science International</i> , 1999, 102, 159-165.	1.3	25
76	D9S1120, a simple STR with a common Native American-specific allele: Forensic optimization, locus characterization and allele frequency studies. <i>Forensic Science International: Genetics</i> , 2008, 3, 7-13.	1.6	25
77	Duplications of the Y-chromosome specific loci P25 and 92R7 and forensic implications. <i>Forensic Science International</i> , 2004, 140, 241-250.	1.3	24
78	SNP Markers as Additional Information to Resolve Complex Kinship Cases. <i>Transfusion Medicine and Hemotherapy</i> , 2015, 42, 385-388.	0.7	24
79	D18S535, D1S1656 and D10S2325: three efficient short tandem repeats for forensic genetics. <i>International Journal of Legal Medicine</i> , 1999, 112, 360-363.	1.2	23
80	Mitochondrial Echoes of First Settlement and Genetic Continuity in El Salvador. <i>PLoS ONE</i> , 2009, 4, e6882.	1.1	23
81	Testing the performance of mtSNP minisequencing in forensic samples. <i>Forensic Science International: Genetics</i> , 2009, 3, 261-264.	1.6	22
82	Analysis of a claimed distant relationship in a deficient pedigree using high density SNP data. <i>Forensic Science International: Genetics</i> , 2012, 6, 350-353.	1.6	22
83	Nonbinary single-nucleotide polymorphism markers. <i>International Congress Series</i> , 2004, 1261, 27-29.	0.2	21
84	Humanized Medium (h7H) Allows Long-Term Primary Follicular Thyroid Cultures From Human Normal Thyroid, Benign Neoplasm, and Cancer. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2013, 98, 2431-2441.	1.8	20
85	Electrophoretic human leukocyte antigenHLA-DQA1 DNA typing after polymerase chain reaction amplification. <i>Electrophoresis</i> , 1991, 12, 1041-1045.	1.3	19
86	Finding genes that underlie physical traits of forensic interest using genetic tools. <i>Forensic Science International: Genetics</i> , 2007, 1, 100-104.	1.6	19
87	Autosomal STR genetic variation in negroid ChocÃ³ and BogotÃ¡ populations. <i>International Journal of Legal Medicine</i> , 2001, 115, 102-104.	1.2	18
88	Y chromosome specific polymorphisms in forensic analysis. <i>Legal Medicine</i> , 1999, 1, 55-60.	0.6	17
89	Genetic diversity of nine STRs in two northwest Iberian populations: Galicia and northern Portugal. <i>International Journal of Legal Medicine</i> , 2000, 114, 109-113.	1.2	17
90	Whole genome amplificationâ€”the solution for a common problem in forensic casework?. <i>International Congress Series</i> , 2004, 1261, 24-26.	0.2	17

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91	Inference of biogeographical ancestry across central regions of Eurasia. <i>International Journal of Legal Medicine</i> , 2016, 130, 73-79.	1.2	17
92	Y-chromosome STR haplotypes in Córdoba (Argentina). <i>Forensic Science International</i> , 2003, 137, 217-220.	1.3	16
93	Band shift analysis of three base-pair repeat alleles in the short tandem repeat locus D12S391. <i>Forensic Science International</i> , 1998, 93, 79-88.	1.3	15
94	Fluorescent SSCP of overlapping fragments (FSSCP-OF): a highly sensitive method for the screening of mitochondrial DNA variation. <i>Forensic Science International</i> , 2001, 124, 97-103.	1.3	15
95	The genetic male legacy from El Salvador. <i>Forensic Science International</i> , 2007, 171, 198-203.	1.3	15
96	Genetic variability of the SNPforID 52-plex identification-SNP panel in Central West Colombia. <i>Forensic Science International: Genetics</i> , 2009, 4, e9-e10.	1.6	15
97	Population data on the D1S1656 and D12S391 STR loci in Andalusia (South Spain) and the Maghreb (North Africa). <i>Forensic Science International</i> , 1999, 104, 33-36.	1.3	14
98	Detection of polymorphisms of human DNA after polymerase chain reaction by miniaturized SDS-PAGE. <i>Forensic Science International</i> , 1992, 55, 27-36.	1.3	13
99	A study of East Timor variability using the SNPforID 52-plex SNP panel. <i>Forensic Science International: Genetics</i> , 2011, 5, e25-e26.	1.6	13
100	Report of the European DNA profiling group (EDNAP)-an investigation of the hypervariable STR loci ACTBP2, APOA11 and D11S554 and the compound loci D12S391 and D1S1656. <i>Forensic Science International</i> , 1998, 98, 193-200.	1.3	12
101	Double- and single-strand conformation polymorphism analysis of point mutations and short tandem repeats. <i>Electrophoresis</i> , 1994, 15, 566-571.	1.3	11
102	Genetic variability at nine STR loci in the Chueta (Majorcan Jews) and the Balearic populations investigated by a single multiplex reaction. <i>International Journal of Legal Medicine</i> , 2000, 113, 263-267.	1.2	11
103	Spanish population data and forensic usefulness of a novel Y-STR set (DYS437, DYS438, DYS439, DYS460,) Tj ETQq _{1,1} 0.784314 rgB _{1,2} 11	1.2	11
104	STR-CODIS typing in Greece. <i>Forensic Science International</i> , 2003, 137, 104-106.	1.3	11
105	Y-chromosome STR-haplotype typing in El Salvador. <i>Forensic Science International</i> , 2004, 142, 45-49.	1.3	11
106	Selecting single nucleotide polymorphisms for forensic applications. <i>International Congress Series</i> , 2004, 1261, 18-20.	0.2	11
107	The genetic male component of two South-Western Colombian populations. <i>Forensic Science International: Genetics</i> , 2009, 3, e59-e61.	1.6	11
108	Population data on 15 autosomal STRs in a sample from Colombia. <i>Forensic Science International: Genetics</i> , 2009, 3, e81-e82.	1.6	11

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109	Sequence variation of a variable short tandem repeat at the D18S535 locus. <i>International Journal of Legal Medicine</i> , 1998, 111, 337-339.	1.2	10
110	STR data for the AmpFISTR profiler plus loci from Macau (China). <i>Forensic Science International</i> , 2001, 123, 74-75.	1.3	10
111	Genetic data on eight STRs (D5S818, D7S820, F13B, LPL, TH01, TPOX, VWA31, CSF1PO) from a Colombian population. <i>Forensic Science International</i> , 2002, 129, 216-218.	1.3	10
112	New method to measure minisatellite variant repeat variation in population genetic studies. <i>American Journal of Human Biology</i> , 2002, 14, 421-428.	0.8	9
113	Significance of micro-geographical population structure in forensic cases: a bayesian exploration. <i>International Journal of Legal Medicine</i> , 2003, 117, 302-305.	1.2	9
114	Y-chromosome haplotype analysis in Antioquia (Colombia). <i>Forensic Science International</i> , 2005, 151, 85-91.	1.3	9
115	Allele frequencies of 20 STRs from Northwest Spain (Galicia). <i>Forensic Science International: Genetics</i> , 2012, 6, e149-e150.	1.6	9
116	Forensic DNA analysis in Europe: current situation and standardization efforts. <i>Forensic Science International</i> , 1997, 86, 87-102.	1.3	8
117	Nine autosomal STRs genotype profiles in a sample from Córdoba (Argentina). <i>Forensic Science International</i> , 2004, 139, 81-83.	1.3	8
118	Genetic Markers in Alcoholic Liver Cirrhosis. <i>Human Heredity</i> , 1992, 42, 235-241.	0.4	7
119	Sequence variation of two hypervariable short tandem repeats at the D22S683 and D6S477 loci. <i>International Journal of Legal Medicine</i> , 2000, 113, 146-149.	1.2	7
120	Population data of Galicia (NW Spain) on the new Y-STRs DYS437, DYS438, DYS439, GATA A10, GATA A7.1, GATA A7.2, GATA C4 and GATA H4. <i>Forensic Science International</i> , 2003, 131, 220-224.	1.3	7
121	Analysis of the SNPforID 52-plex markers in four Native American populations from Venezuela. <i>Forensic Science International: Genetics</i> , 2012, 6, e142-e145.	1.6	7
122	Fluorescence-based amplification of the STR loci D18S535, D1S1656 and D12S391 in a population sample from Aragon (North Spain). <i>International Journal of Legal Medicine</i> , 1999, 113, 58-59.	1.2	6
123	Typing Y-chromosome single nucleotide polymorphisms with DNA microarray technology. <i>International Congress Series</i> , 2003, 1239, 21-25.	0.2	5
124	Differentiation of African Components of Ancestry to Stratify Groups in a Caseâ€“Control Study of a Brazilian Urban Population. <i>Genetic Testing and Molecular Biomarkers</i> , 2012, 16, 524-530.	0.3	5
125	Distribution of allele frequencies of 20 STRs loci in a population sample from Calabria, Southern Italy. <i>Forensic Science International: Genetics</i> , 2012, 6, e137-e138.	1.6	5
126	Population specific single nucleotide polymorphisms. <i>International Congress Series</i> , 2004, 1261, 233-235.	0.2	4

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127	Development and validation of a next generation STR ESS-pentaplex. Forensic Science International: Genetics Supplement Series, 2009, 2, 25-26.	0.1	4
128	Genetic data on three complex STRs (ACTBP2, D21S11 and HUMFIBRA/FGA) in the Galician population (NW) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.2	3
129	Y-chromosome STRs in populations of Bantu origin from Mozambique: male contribution to the Africa genetic pool and forensic implications. International Congress Series, 2003, 1239, 419-424.	0.2	3
130	Genetic variability of the SNPforID 52-plex identification SNP panel in Italian population samples. Forensic Science International: Genetics, 2012, 6, e185-e186.	1.6	3
131	¹⁶⁰Thr Mutation in the Rhodopsin Gene Associated withRet initis pigmentosa. Human Heredity, 1998, 48, 237-240.	0.4	2
132	Behavior of loci D1S1656 and D12S391 in a sample from Maracaibo, Venezuela. American Journal of Human Biology, 2003, 15, 68-71.	0.8	2
133	Typing mtDNA SNPs of forensic and population interest with snapshot. International Congress Series, 2004, 1261, 419-421.	0.2	1
134	Microgeographic patterns of highly informative Y-chromosome haplotypes (using biallelic markers) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 Series, 2003, 1239, 61-66.	0.2	0
135	Mitochondrial DNA variability patterns in Southeast Africa and forensic implications. International Congress Series, 2003, 1239, 541-545.	0.2	0