Niels Morling

List of Publications by Year
in descending order

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139	4,070	117453	143772
papers	citations	34 h-index	57 g-index

141	141	141	3378
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Next generation sequencing and its applications in forensic genetics. Forensic Science International: Genetics, 2015, 18, 78-89.	1.6	338
2	A global analysis of Y-chromosomal haplotype diversity for 23 STR loci. Forensic Science International: Genetics, 2014, 12, 12-23.	1.6	214
3	Massively parallel sequencing of forensic STRs: Considerations of the DNA commission of the International Society for Forensic Genetics (ISFG) on minimal nomenclature requirements. Forensic Science International: Genetics, 2016, 22, 54-63.	1.6	190
4	Toward Male Individualization with Rapidly Mutating Y-Chromosomal Short Tandem Repeats. Human Mutation, 2014, 35, 1021-1032.	1.1	151
5	Recommendations of the DNA Commission of the International Society for Forensic Genetics (ISFG) on quality control of autosomal Short Tandem Repeat allele frequency databasing (STRidER). Forensic Science International: Genetics, 2016, 24, 97-102.	1.6	130
6	Second-generation sequencing of forensic STRs using the Ion Torrentâ„¢ HID STR 10-plex and the Ion PGMâ„¢. Forensic Science International: Genetics, 2015, 14, 132-140.	1.6	112
7	Evaluation of the Ion Torrentâ„¢ HID SNP 169-plex: A SNP typing assay developed for human identification by second generation sequencing. Forensic Science International: Genetics, 2014, 12, 144-154.	1.6	95
8	Paternity Testing Commission of the International Society of Forensic Genetics: recommendations on genetic investigations in paternity cases. Forensic Science International, 2002, 129, 148-157.	1.3	85
9	DNA commission of the International society for forensic genetics: Assessing the value of forensic biological evidence - Guidelines highlighting the importance of propositions. Forensic Science International: Genetics, 2018, 36, 189-202.	1.6	83
10	Evaluation of the Precision ID Ancestry Panel for crime case work: A SNP typing assay developed for typing of 165 ancestral informative markers. Forensic Science International: Genetics, 2017, 28, 138-145.	1.6	82
11	Typing of 30 insertion/deletions in Danes using the first commercial indel kitâ€"MentypeÂ® DIPplex. Forensic Science International: Genetics, 2012, 6, e72-e74.	1.6	77
12	Analysis of 12 X-STRs in Greenlanders, Danes and Somalis using Argus X-12. International Journal of Legal Medicine, 2012, 126, 121-128.	1.2	70
13	Characterization of mutations and sequence variants in the D21S11 locus by next generation sequencing. Forensic Science International: Genetics, 2014, 8, 68-72.	1.6	70
14	Second generation sequencing of three STRs D3S1358, D12S391 and D21S11 in Danes and a new nomenclature for sequenced STR alleles. Forensic Science International: Genetics, 2014, 12, 38-41.	1.6	70
15	Genetic investigation of 100 heart genes in sudden unexplained death victims in a forensic setting. European Journal of Human Genetics, 2016, 24, 1797-1802.	1.4	65
16	DNA commission of the International society for forensic genetics: Assessing the value of forensic biological evidence - Guidelines highlighting the importance of propositions. Part II: Evaluation of biological traces considering activity level propositions. Forensic Science International: Genetics, 2020, 44, 102186.	1.6	59
17	Genetic determinants of hair and eye colours in the Scottish and Danish populations. BMC Genetics, 2009, 10, 88.	2.7	57
18	The discrete Laplace exponential family and estimation of Y-STR haplotype frequencies. Journal of Theoretical Biology, 2013, 329, 39-51.	0.8	55

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19	Genetic investigations of sudden unexpected deaths in infancy using next-generation sequencing of 100 genes associated with cardiac diseases. European Journal of Human Genetics, 2016, 24, 817-822.	1.4	55
20	Forensic genetic SNP typing of low-template DNA and highly degraded DNA from crime case samples. Forensic Science International: Genetics, 2013, 7, 345-352.	1.6	53
21	ISO 17025 validation of a nextâ€generation sequencing assay for relationship testing. Electrophoresis, 2016, 37, 2822-2831.	1.3	52
22	Evaluation of DNA Variants Associated with Androgenetic Alopecia and Their Potential to Predict Male Pattern Baldness. PLoS ONE, 2015, 10, e0127852.	1.1	51
23	Statistical model for degraded DNA samples and adjusted probabilities for allelic drop-out. Forensic Science International: Genetics, 2012, 6, 97-101.	1.6	50
24	Forensic and population genetic analyses of Danes, Greenlanders and Somalis typed with the Yfiler Â® Plus PCR amplification kit. Forensic Science International: Genetics, 2015, 16, 232-236.	1.6	48
25	A method for the analysis of 32 X chromosome insertion deletion polymorphisms in a single PCR. International Journal of Legal Medicine, 2012, 126, 97-105.	1.2	45
26	miR-125b induces cellular senescence in malignant melanoma. BMC Dermatology, 2014, 14, 8.	2.1	45
27	Massively parallel pyrosequencing of the mitochondrial genome with the 454 methodology in forensic genetics. Forensic Science International: Genetics, 2014, 12, 30-37.	1.6	41
28	Allelic drop-out probabilities estimated by logistic regressionâ€"Further considerations and practical implementation. Forensic Science International: Genetics, 2012, 6, 263-267.	1.6	40
29	Analysis of ancestry informative markers in three main ethnic groups from Ecuador supports a trihybrid origin of Ecuadorians. Forensic Science International: Genetics, 2017, 31, 29-33.	1.6	40
30	Increasing the reference populations for the 55 AISNP panel: the need and benefits. International Journal of Legal Medicine, 2017, 131, 913-917.	1.2	38
31	Towards broadening Forensic DNA Phenotyping beyond pigmentation: Improving the prediction of head hair shape from DNA. Forensic Science International: Genetics, 2018, 37, 241-251.	1.6	38
32	MDM2 Inhibitor Nutlin-3a Induces Apoptosis and Senescence in Cutaneous T-Cell Lymphoma: Role of p53. Journal of Investigative Dermatology, 2012, 132, 1487-1496.	0.3	37
33	Sequencing of 231 forensic genetic markers using the MiSeq FGxâ„¢ forensic genomics systemÂâ€"Âan evaluation of the assay and software. Forensic Sciences Research, 2018, 3, 111-123.	0.9	37
34	Evaluation of the predictive capacity of DNA variants associated with straight hair in Europeans. Forensic Science International: Genetics, 2015, 19, 280-288.	1.6	36
35	Whose DNA is this? How relevant a question? (a note for forensic scientists). Forensic Science International: Genetics, 2013, 7, 467-470.	1.6	35
36	Evaluation of Forensic DNA Traces When Propositions of Interest Relate to Activities: Analysis and Discussion of Recurrent Concerns. Frontiers in Genetics, 2016, 7, 215.	1.1	35

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37	Introduction of the Python script STRinNGS for analysis of STR regions in FASTQ or BAM files and expansion of the Danish STR sequence database to 11 STRs. Forensic Science International: Genetics, 2016, 21, 68-75.	1.6	34
38	The half-life of 25(OH)D after UVB exposure depends on gender and vitamin D receptor polymorphism but mainly on the start level. Photochemical and Photobiological Sciences, 2017, 16, 985-995.	1.6	33
39	Multiplex PCR with minisequencing as an effective high-throughput SNP typing method for formalin-fixed tissue. Electrophoresis, 2007, 28, 2361-2367.	1.3	31
40	PCR in forensic genetics. Biochemical Society Transactions, 2009, 37, 438-440.	1.6	31
41	Genetic analyses of the human eye colours using a novel objective method for eye colour classification. Forensic Science International: Genetics, 2013, 7, 508-515.	1.6	31
42	Importance of nonsynonymous <scp>OCA</scp>2 variants in human eye color prediction. Molecular Genetics & Genomic Medicine, 2016, 4, 420-430.	0.6	31
43	Quantification of massively parallel sequencing libraries â€“ a comparative study of eight methods. Scientific Reports, 2018, 8, 1110.	1.6	30
44	The effect of gender on eye colour variation in European populations and an evaluation of the IrisPlex prediction model. Forensic Science International: Genetics, 2014, 11, 1-6.	1.6	29
45	Characterization of a new HLA-G allele encoding a nonconservative amino acid substitution in the Î±3 domain (exon 4) and its relevance to certain complications in pregnancy. Immunogenetics, 2001, 53, 48-53.	1.2	28
46	Performance of two 17 locus forensic identification STR kitsâ€”Applied Biosystems's AmpFâ„“STRÂ® NGMSElectâ„¢ and Promega's PowerPlexÂ® ESI17 kits. Forensic Science International: Genetics, 2012, 6, 523-531.	1.6	26
47	Frequencies of HID-ion ampliseq ancestry panel markers among greenlanders. Forensic Science International: Genetics, 2016, 24, 60-64.	1.6	26
48	New clues to the evolutionary history of the main European paternal lineage M269: dissection of the Y-SNP S116 in Atlantic Europe and Iberia. European Journal of Human Genetics, 2016, 24, 437-441.	1.4	26
49	Evaluation of Four Automated Protocols for Extraction of DNA from FTA Cards. Journal of the Association for Laboratory Automation, 2013, 18, 404-410.	2.8	25
50	Technological Innovations in Forensic Genetics: Social, Legal and Ethical Aspects. Recent Advances in DNA & Gene Sequences, 2015, 8, 98-103.	0.7	25
51	Graft-derived anti-HPA-2b production after allogeneic bone-marrow transplantation. British Journal of Haematology, 1994, 86, 651-653.	1.2	23
52	Kinship Analysis with Diallelic SNPs Â– Experiences with the SNPforID Multiplex in an ISO17025 Accreditated Laboratory. Transfusion Medicine and Hemotherapy, 2012, 39, 195-201.	0.7	23
53	Collaborative EDNAP exercise on the IrisPlex system for DNA-based prediction of human eye colour. Forensic Science International: Genetics, 2014, 11, 241-251.	1.6	23
54	High-throughput sequencing of forensic genetic samples using punches of FTA cards with buccal swabs. BioTechniques, 2016, 61, 149-151.	0.8	23

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55	Major inter-personal variation in the increase and maximal level of 25-hydroxy vitamin D induced by UVB. Photochemical and Photobiological Sciences, 2016, 15, 536-545.	1.6	23
56	Whole genome and transcriptome sequencing of post-mortem cardiac tissues from sudden cardiac death victims identifies a gene regulatory variant in NEXN. International Journal of Legal Medicine, 2019, 133, 1699-1709.	1.2	23
57	Comparison of manual and automated AmpliSeqâ„¢ workflows in the typing of a Somali population with the Precision ID Identity Panel. Forensic Science International: Genetics, 2017, 31, 118-125.	1.6	22
58	Platelet Alloimmunization after Transfusion. A Prospective Study in 117 Heart Surgery Patients. Vox Sanguinis, 1997, 72, 238-241.	0.7	21
59	Decrease DNA contamination in the laboratories. Forensic Science International: Genetics Supplement Series, 2017, 6, e577-e578.	0.1	21
60	Forensic genetics. Lancet, The, 2004, 364, 10-11.	6.3	20
61	Identifying Contributors of DNA Mixtures by Means of Quantitative Information of STR Typing. Journal of Computational Biology, 2012, 19, 887-902.	0.8	20
62	Stutter analysis of complex STR MPS data. Forensic Science International: Genetics, 2018, 35, 107-112.	1.6	20
63	Weight of the evidence of genetic investigations of ancestry informative markers. Theoretical Population Biology, 2018, 120, 1-10.	0.5	19
64	Ancestry prediction efficiency of the software GenoGeographer using a z-score method and the ancestry informative markers in the Precision ID Ancestry Panel. Forensic Science International: Genetics, 2020, 44, 102154.	1.6	19
65	SPERM HY-LITERâ„¢ for the identification of spermatozoa from sexual assault evidence. Forensic Science International: Genetics, 2014, 12, 161-167.	1.6	18
66	Identifying the most likely contributors to a Y-STR mixture using the discrete Laplace method. Forensic Science International: Genetics, 2015, 15, 76-83.	1.6	18
67	Forensic and phylogeographic characterisation of mtDNA lineages from Somalia. International Journal of Legal Medicine, 2012, 126, 573-579.	1.2	17
68	A comparison of genetic findings in sudden cardiac death victims and cardiac patients: the importance of phenotypic classification. Europace, 2015, 17, 350-357.	0.7	17
69	Characterization of mutations and sequence variations in complex STR loci by second generation sequencing. Forensic Science International: Genetics Supplement Series, 2013, 4, e218-e219.	0.1	16
70	Peopling of the North Circumpolar Region â€“ Insights from Y Chromosome STR and SNP Typing of Greenlanders. PLoS ONE, 2015, 10, e0116573.	1.1	16
71	Analysis of mainland Japanese and Okinawan Japanese populations using the precision ID Ancestry Panel. Forensic Science International: Genetics, 2018, 33, 106-109.	1.6	16
72	DNA polymorphism of HLA class II genes in systemic lupus erythematosus. Tissue Antigens, 1994, 43, 34-37.	1.0	15

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73	The peopling of Greenland: further insights from the analysis of genetic diversity using autosomal and X-chromosomal markers. European Journal of Human Genetics, 2015, 23, 245-251.	1.4	15
74	A study of the peopling of Greenland using next generation sequencing of complete mitochondrial genomes. American Journal of Physical Anthropology, 2016, 161, 698-704.	2.1	15
75	Genomic and immunohistochemical characterisation of a lacrimal gland oncocytoma and review of literature. Oncology Letters, 2017, 14, 4176-4182.	0.8	15
76	Pigment genes not skin pigmentation affect UVB-induced vitamin D. Photochemical and Photobiological Sciences, 2019, 18, 448-458.	1.6	15
77	HLAâ€DPB1 typing with polymerase chain reaction and restriction fragment length polymorphism technique in Danes. Tissue Antigens, 1992, 40, 140-144.	1.0	14
78	Statistical modelling of Ion PGM HID STR 10-plex MPS data. Forensic Science International: Genetics, 2017, 28, 82-89.	1.6	14
79	Association between brown eye colour in rs12913832:GG individuals and SNPs in TYR, TYRP1, and SLC24A4. PLoS ONE, 2020, 15, e0239131.	1.1	14
80	The role of the glutathione S-transferase genes GSTT1, GSTM1, and GSTP1 in acetaminophen-poisoned patients. Clinical Toxicology, 2012, 50, 27-33.	0.8	13
81	A report of the 2009â€“2011 paternity and relationship testing workshops of the English Speaking Working Group of the International Society For Forensic Genetics. Forensic Science International: Genetics, 2014, 9, e1-e2.	1.6	13
82	Targeted molecular genetic testing in young sudden cardiac death victims from Western Denmark. International Journal of Legal Medicine, 2020, 134, 111-121.	1.2	13
83	Reinvestigations of six unusual paternity cases by typing of autosomal singleâ€nucleotide polymorphisms. Transfusion, 2012, 52, 425-430.	0.8	12
84	Prediction of Eye Colour in Scandinavians Using the EyeColour 11 (EC11) SNP Set. Genes, 2021, 12, 821.	1.0	12
85	The transcriptome of hand eczema assessed by tape stripping. Contact Dermatitis, 2022, 86, 71-79.	0.8	12
86	Cluster analysis of European Y-chromosomal STR haplotypes using the discrete Laplace method. Forensic Science International: Genetics, 2014, 11, 182-194.	1.6	11
87	The stratum corneum transcriptome in atopic dermatitis can be assessed by tape stripping. Journal of Dermatological Science, 2021, 101, 14-21.	1.0	11
88	Detection of fetal-specific DNA after enrichment for trophoblasts using the monoclonal antibody LK26 in model systems but failure to demonstrate fetal DNA in maternal peripheral blood. , 1999, 19, 271-278.		10
89	Evaluation of the iPLEXÂ® ADME PGx Pro Panel and allele frequencies of pharmacogenetic markers in Danes. Clinical Biochemistry, 2016, 49, 1299-1301.	0.8	10
90	Non-invasive prenatal paternity testing using a standard forensic genetic massively parallel sequencing assay for amplification of human identification SNPs. International Journal of Legal Medicine, 2019, 133, 1361-1368.	1.2	10

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91	Evaluation of the Precision of Ancestry Inferences in South American Admixed Populations. Frontiers in Genetics, 2020, 11, 966.	1.1	10
92	Refining the genetic portrait of Portuguese Roma through Xâ€chromosomal markers. American Journal of Physical Anthropology, 2012, 148, 389-394.	2.1	9
93	Estimating Y-STR allelic drop-out rates and adjusting for interlocus balances. Forensic Science International: Genetics, 2013, 7, 327-336.	1.6	9
94	eDNAâ€"An expert software system for comparison and evaluation of DNA profiles in forensic casework. Forensic Science International: Genetics Supplement Series, 2015, 5, e400-e402.	0.1	9
95	On the Bayesian approach to forensic age estimation of living individuals. Forensic Science International, 2017, 281, e24-e29.	1.3	9
96	Analysis of 49 autosomal SNPs in an Iraqi population. Forensic Science International: Genetics, 2013, 7, 198-199.	1.6	8
97	GenoGeographer â€“ A tool for genogeographic inference. Forensic Science International: Genetics Supplement Series, 2017, 6, e463-e465.	0.1	8
98	A comparative study of single nucleotide variant detection performance using three massively parallel sequencing methods. PLoS ONE, 2020, 15, e0239850.	1.1	8
99	Monozygotic twins discordant for narcolepsy type 1 and multiple sclerosis. Neurology: Neuroimmunology and NeuroInflammation, 2016, 3, e249.	3.1	7
100	Modelling allelic drop-outs in STR sequencing data generated by MPS. Forensic Science International: Genetics, 2018, 37, 6-12.	1.6	7
101	Sequencing of human identification markers in an Uyghur population using the MiSeq FGx^{TM} Forensic Genomics System. Forensic Sciences Research, 2022, 7, 154-162.	0.9	7
102	Evaluation of a custom GeneReadâ„¢ massively parallel sequencing assay with 210 ancestry informative SNPs using the Ion S5â„¢ and MiSeq platforms. Forensic Science International: Genetics, 2021, 50, 102411.	1.6	7
103	Reproducibility of the Infinium methylationEPIC BeadChip assay using low DNA amounts. Epigenetics, 2022, 17, 1636-1645.	1.3	7
104	Repeated extraction of DNA from FTA cards. Forensic Science International: Genetics Supplement Series, 2011, 3, e345-e346.	0.1	6
105	Sequence variants of allele 22 and 23 of DYS635 causing different stutter rates. Forensic Science International: Genetics, 2012, 6, e161-e162.	1.6	6
106	Associations between second to fourth digit ratio, cortisol, vitamin D, and body composition among Polish children. Scientific Reports, 2021, 11, 7029.	1.6	6
107	Typing of 49 Autosomal SNPs by Single Base Extension and Capillary Electrophoresis for Forensic Genetic Testing. Methods in Molecular Biology, 2012, 830, 87-107.	0.4	5
108	Body fluid identification of blood, saliva and semen using second generation sequencing of micro-RNA. Forensic Science International: Genetics Supplement Series, 2013, 4, e204-e205.	0.1	5

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109	Characterization of sequence variations in the D21S11 locus in Danes, Somalis and Greenlanders by second generation sequencing. Forensic Science International: Genetics Supplement Series, 2013, 4, e302-e303.	0.1	5
110	Estimating drop-out probabilities of STR alleles accounting for stutters, detection threshold truncation and degradation. Forensic Science International: Genetics Supplement Series, 2013, 4, e51-e52.	0.1	5
111	Next-generation sequencing of multiple individuals per barcoded library by deconvolution of sequenced amplicons using endonuclease fragment analysis. BioTechniques, 2014, 57, 91-4.	0.8	5
112	Modelling noise in second generation sequencing forensic genetics STR data using a one-inflated (zero-truncated) negative binomial model. Forensic Science International: Genetics Supplement Series, 2015, 5, e416-e417.	0.1	5
113	Cortisol concentration affects fat and muscle mass among Polish children aged 6â€“13 years. BMC Pediatrics, 2021, 21, 365.	0.7	5
114	Pigmentary Markers in Danes â€“ Associations with Quantitative Skin Colour, Nevi Count, Familial Atypical Multiple-Mole, and Melanoma Syndrome. PLoS ONE, 2016, 11, e0150381.	1.1	5
115	Sequences of microvariant/â€œoff-ladderâ€•STR alleles. Forensic Science International: Genetics Supplement Series, 2011, 3, e204-e205.	0.1	4
116	Biomek 3000. Journal of the Association for Laboratory Automation, 2012, 17, 378-386.	2.8	4
117	Yfilerâ“‡ Plus population samples and dilution series: stutters, analytic thresholds, and drop-out probabilities. International Journal of Legal Medicine, 2017, 131, 1503-1511.	1.2	4
118	Sequence variants in muscle tissueâ€related genes may determine the severity of muscle contractures in cerebral palsy. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 12-24.	1.1	4
119	Skin pigmentation and genetic variants in an admixed Brazilian population of primarily European ancestry. International Journal of Legal Medicine, 2020, 134, 1569-1579.	1.2	4
120	Forensic application and genetic diversity of 21 autosomal STR loci in five major population groups of Pakistan. International Journal of Legal Medicine, 2021, 135, 775-777.	1.2	4
121	Differential Methylation in the GSTT1 Regulatory Region in Sudden Unexplained Death and Sudden Unexpected Death in Epilepsy. International Journal of Molecular Sciences, 2021, 22, 2790.	1.8	4
122	Genetic investigations of 100 inherited cardiac disease-related genes in deceased individuals with schizophrenia. International Journal of Legal Medicine, 2021, 135, 1395-1405.	1.2	4
123	Association of saliva 25(OH)D concentration with body composition and proportion among preâ€pubertal and pubertal Polish children. American Journal of Human Biology, 2020, 32, e23397.	0.8	4
124	Evaluation of Y-STR analyses of sperm cell negative vaginal samples. Forensic Science International: Genetics Supplement Series, 2011, 3, e141-e142.	0.1	3
125	Identical twins in forensic genetics â€” Epidemiology and risk based estimation of weight of evidence. Science and Justice - Journal of the Forensic Science Society, 2015, 55, 408-414.	1.3	3
126	The multivariate Dirichlet-multinomial distribution and its application in forensic genetics to adjust for subpopulation effects using the <mml:math altimg="si12.gif" display="inline" overflow="scroll" xmlns:xocs="http://www.elsevier.com/xml/xocs/dtd" xmlns:xs="http://www.w3.org/2001/XMLSchema" xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance" xmlns="http://www.elsevier.com/xml/ja/dtd" xmlns:ja="http://www.elsevier.com/xml/ja/dtd" xmlns:mml="http://www.w3.org/1998/Math/MathML" xmlns:tb="http://www.elsev. Theoretical Population Biology, 2015, 105, 24-32.	0.5	3

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127	Serum 25(OH)D levels after oral vitamin D 3 supplementation and UVB exposure correlate. Photodermatology Photoimmunology and Photomedicine, 2019, 35, 344-353.	0.7	3
128	Test of Investigator ESSPLEX SE QS with quality sensors. Forensic Science International: Genetics Supplement Series, 2015, 5, e490-e491.	0.1	2
129	Targeted exon sequencing in deceased schizophrenia patients in Denmark. International Journal of Legal Medicine, 2020, 134, 135-147.	1.2	2
130	Analysis of Skin Pigmentation and Genetic Ancestry in Three Subpopulations from Pakistan: Punjabi, Pashtun, and Baloch. Genes, 2021, 12, 733.	1.0	2
131	Survival of Fetuses and Viruses: Universal Mechanisms of Coâ€Existence with an Immunological Potent Host. American Journal of Reproductive Immunology, 1999, 41, 353-355.	1.2	1
132	Study of 25 X-chromosome Single Nucleotide Polymorphisms in African and Asian populations. Forensic Science International: Genetics Supplement Series, 2011, 3, e139-e140.	0.1	1
133	Correlation of iris biometrics and DNA. , 2013, , .		1
134	Results of the 2015 Relationship Testing Workshop of the English Speaking Working Group. Forensic Science International: Genetics Supplement Series, 2015, 5, e320-e321.	0.1	1
135	Anthropological analyses of 30 insertion/deletion autosomal markers in five major ethnic groups of Pakistan. Forensic Sciences Research, 0, , 1-5.	0.9	1
136	GENETIC PORTRAIT OF THE PUNJABI POPULATION FROM PAKISTAN USING THE PRECISION ID ANCESTRY PANEL. Forensic Science International: Genetics Supplement Series, 2019, 7, 87-89.	0.1	1
137	Group Specific Component in Serum and Otosclerosis: No Association. Acta Oto-Laryngologica, 1994, 114, 303-304.	0.3	0
138	The effect of wild card designations and rare alleles in forensic DNA database searches. Forensic Science International: Genetics, 2015, 16, 98-104.	1.6	0
139	A 48-plex Autosomal SNP GenPlexâ„¢ Assay for Human Individualization and Relationship Testing. Methods in Molecular Biology, 2012, 830, 73-85.	0.4	0