Weibo Liang

List of Publications by Year
in descending order

148 papers	2,673 citations	172457 29 h-index	254184 43 g-index
150 all docs	150 docs citations	150 times ranked	2724 citing authors

#	Article	IF	Citations
1	miR-212-5p attenuates ferroptotic neuronal death after traumatic brain injury by targeting Ptgs2. Molecular Brain, 2019, 12, 78.	2.6	123
2	The association of interleukin-16 polymorphisms with IL-16 serum levels and risk of colorectal and gastric cancer. Carcinogenesis, 2008, 30, 295-299.	2.8	95
3	The association between two polymorphisms in pre-miRNAs and breast cancer risk: a meta-analysis. Breast Cancer Research and Treatment, 2011, 125, 571-574.	2.5	91
4	Fermented dairy foods intake and risk of cancer. International Journal of Cancer, 2019, 144, 2099-2108.	5.1	79
5	Evaluation of the Microhaplotypes panel for DNA mixture analyses. Forensic Science International: Genetics, 2018, 35, 149-155.	3.1	64
6	RAD51 135G/C polymorphism and breast cancer risk: a meta-analysis from 21 studies. Breast Cancer Research and Treatment, 2011, 125, 827-835.	2.5	60
7	Effect of aging on the microstructure, hardness and chemical composition of dentin. Archives of Oral Biology, 2015, 60, 1811-1820.	1.8	59
8	A microhaplotypes panel for massively parallel sequencing analysis of DNA mixtures. Forensic Science International: Genetics, 2019, 40, 140-149.	3.1	58
9	A genetic variant in the promoter region of miR-34b/c is associated with a reduced risk of colorectal cancer. Biological Chemistry, 2013, 394, 415-420.	2.5	52
10	Forensic age estimation for pelvic X-ray images using deep learning. European Radiology, 2019, 29, 2322-2329.	4.5	51
11	Null Genotypes of GSTM1 and GSTT1 Contribute to Risk of Cervical Neoplasia: An Evidence-Based Meta-Analysis. PLoS ONE, 2011, 6, e20157.	2.5	49
12	Semen-specific miRNAs: Suitable for the distinction of infertile semen in the body fluid identification?. Forensic Science International: Genetics, 2018, 33, 161-167.	3.1	49
13	Development and application of a nonbinary SNP-based microhaplotype panel for paternity testing involving close relatives. Forensic Science International: Genetics, 2020, 46, 102255.	3.1	48
14	Identifying novel microhaplotypes for ancestry inference. International Journal of Legal Medicine, 2019, 133, 983-988.	2.2	47
15	Association of transforming growth factor-β1 gene polymorphisms with genetic susceptibility to nasopharyngeal carcinoma. Clinica Chimica Acta, 2007, 380, 165-169.	1.1	46
16	A comparative study of insertion/deletion polymorphisms applied among Southwest, South and Northwest Chinese populations using InvestigatorÂ® DIPplex. Forensic Science International: Genetics, 2016, 21, 10-14.	3.1	45
17	Genetic diversity of 21 autosomal STR loci in the Han population from Sichuan province, Southwest China. Forensic Science International: Genetics, 2017, 31, e33-e35.	3.1	41
18	Interleukin-10 gene promoter polymorphisms and the risk of nasopharyngeal carcinoma. Tissue Antigens, 2007, 70, 12-17.	1.0	40

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19	IL-8 â€"251A/T polymorphism is associated with decreased cancer risk among population-based studies: Evidence from a meta-analysis. European Journal of Cancer, 2010, 46, 1333-1343.	2.8	37
20	Hippocampus-dependent spatial memory impairment due to molar tooth loss is ameliorated by an enriched environment. Archives of Oral Biology, 2016, 61, 1-7.	1.8	37
21	Genetic Polymorphisms of the DNA Repair Gene and Risk of Nasopharyngeal Carcinoma. DNA and Cell Biology, 2007, 26, 491-496.	1.9	36
22	The IL-16 gene polymorphisms and the risk of the systemic lupus erythematosus. Clinica Chimica Acta, 2009, 403, 223-225.	1.1	36
23	Genetic polymorphism of Interleukin-16 and risk of nasopharyngeal carcinoma. Clinica Chimica Acta, 2009, 409, 132-135.	1.1	36
24	A Functional Polymorphism in the Promoter of MiR-143/145 Is Associated With the Risk of Cervical Squamous Cell Carcinoma in Chinese Women. Medicine (United States), 2015, 94, e1289.	1.0	36
25	Association Between Single-Nucleotide Polymorphisms in Pre-miRNAs and the Risk of Asthma in a Chinese Population. DNA and Cell Biology, 2011, 30, 919-923.	1.9	35
26	Characteristics of eight X-STR loci for forensic purposes in the Chinese population. International Journal of Legal Medicine, 2011, 125, 127-131.	2.2	35
27	Genotyping polymorphic microhaplotype markers through the IlluminaÂ® MiSeq platform for forensics. Forensic Science International: Genetics, 2019, 39, 1-7.	3.1	35
28	Identification of serum biomarkers for nasopharyngeal carcinoma by proteomic analysis. Cancer, 2008, 112, 544-551.	4.1	31
29	Screening and confirmation of microRNA markers for distinguishing between menstrual and peripheral blood. Forensic Science International: Genetics, 2017, 30, 24-33.	3.1	31
30	Moringa oleifera Lam and its Therapeutic Effects in Immune Disorders. Frontiers in Pharmacology, 2020, 11, 566783.	3.5	31
31	CTLA4 and CD86 gene polymorphisms and susceptibility to chronic obstructive pulmonary disease. Human Immunology, 2010, 71, 1141-1146.	2.4	28
32	The association between ATM D1853N polymorphism and breast cancer susceptibility: a meta-analysis. Journal of Experimental and Clinical Cancer Research, 2010, 29, 117.	8.6	28
33	Association between SNPs in pre-miRNA and risk of chronic obstructive pulmonary disease. Clinical Biochemistry, 2011, 44, 813-816.	1.9	28
34	Circular Ribonucleic Acid Expression Profile in Mouse Cortex after Traumatic Brain Injury. Journal of Neurotrauma, 2019, 36, 1018-1028.	3.4	28
35	Association of IL-1B Gene Polymorphisms with Nasopharyngeal Carcinoma in a Chinese Population. Clinical Oncology, 2008, 20, 207-211.	1.4	27
36	Association of single nucleotide polymorphisms in interleukin 12 (IL-12A and -B) with asthma in a Chinese population. Human Immunology, 2011, 72, 603-606.	2.4	27

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37	Detection of promoter methylation status of suppressor of cytokine signaling 3 (SOCS3) in tissue and plasma from Chinese patients with different hepatic diseases. Clinical and Experimental Medicine, 2018, 18, 79-87.	3.6	27
38	The xeroderma pigmentosum group C gene polymorphisms and genetic susceptibility of nasopharyngeal carcinoma. Acta OncolÃ³gica, 2008, 47, 379-384.	1.8	26
39	Association Between<i>pri-miR-218</i>Polymorphism and Risk of Hepatocellular Carcinoma in a Han Chinese Population. DNA and Cell Biology, 2012, 31, 761-765.	1.9	26
40	Mutational analysis of 33 autosomal short tandem repeat (STR) loci in southwest Chinese Han population based on trio parentage testing. Forensic Science International: Genetics, 2016, 23, 86-90.	3.1	25
41	Genetic portrait of 27 Y-STR loci in the Tibetan ethnic population of the Qinghai province of China. Forensic Science International: Genetics, 2018, 34, e18-e19.	3.1	25
42	Two-person DNA mixture interpretation based on a novel set of SNP-STR markers. Forensic Science International: Genetics, 2018, 37, 37-45.	3.1	25
43	Expression difference of miR-10b and miR-135b between the fertile and infertile semen samples (p). Forensic Science International: Genetics Supplement Series, 2017, 6, e257-e259.	0.3	24
44	Population genetics for 17 Y-STR loci(AmpFISTRÂ®Y-filerTM) in Luzhou Han ethnic group. Forensic Science International: Genetics, 2013, 7, e23-e26.	3.1	23
45	Interactions ofmiR-34b/candTP53Polymorphisms on the Risk of Intracranial Aneurysm. Clinical and Developmental Immunology, 2012, 2012, 1-7.	3.3	22
46	Evaluation of the microhaplotype markers in kinship analysis. Electrophoresis, 2019, 40, 1091-1095.	2.4	22
47	Single nucleotide polymorphisms of VEGF gene and Psoriasis risk. Journal of Dermatological Science, 2008, 49, 263-265.	1.9	21
48	An investigation of a set of DIP-STR markers to detect unbalanced DNA mixtures among the southwest Chinese Han population. Forensic Science International: Genetics, 2017, 31, 34-39.	3.1	21
49	Association of Matrix Metalloproteinases 1, 7, and 9 Gene Polymorphisms with Genetic Susceptibility to Colorectal Carcinoma in a Han Chinese Population. DNA and Cell Biology, 2010, 29, 657-661.	1.9	20
50	Multi-Indel: A Microhaplotype Marker Can Be Typed Using Capillary Electrophoresis Platforms. Frontiers in Genetics, 2020, 11, 567082.	2.3	19
51	No association between epidermal growth factor and epidermal growth factor receptor polymorphisms and nasopharyngeal carcinoma. Cancer Genetics and Cytogenetics, 2008, 185, 69-73.	1.0	18
52	5 miRNA expression analyze in post-mortem interval (PMI) within 48h. Forensic Science International: Genetics Supplement Series, 2013, 4, e190-e191.	0.3	18
53	Development of a SNP-STRs multiplex for forensic identification. Forensic Science International: Genetics Supplement Series, 2015, 5, e598-e600.	0.3	18
54	Postmortem interval determination using mRNA markers and DNA normalization. International Journal of Legal Medicine, 2020, 134, 149-157.	2.2	18

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55	A new method to detect methylation profiles for forensic body fluid identification combining ARMS-PCR technique and random forest model. Forensic Science International: Genetics, 2020, 49, 102371.	3.1	16
56	Association of Tumor Necrosis Factor Gene Polymorphisms with Susceptibility to Dilated Cardiomyopathy in a Han Chinese Population. DNA and Cell Biology, 2010, 29, 625-628.	1.9	15
57	NGS technology makes microhaplotype a potential forensic marker. Forensic Science International: Genetics Supplement Series, 2015, 5, e233-e234.	0.3	15
58	Microhaplotype identified and performed in genetic investigation using PCR-SSCP. Forensic Science International: Genetics, 2017, 28, e1-e7.	3.1	15
59	The association between dilated cardiomyopathy and RTN4 3â€²UTR insertion/deletion polymorphisms. Clinica Chimica Acta, 2009, 400, 21-24.	1.1	14
60	CD86 +1057 G/A Polymorphism and the Risk of Colorectal Cancer. DNA and Cell Biology, 2010, 29, 381-386.	1.9	14
61	Interactions of interleukin-12A and interleukin-12B polymorphisms on the risk of intracranial aneurysm. Molecular Biology Reports, 2012, 39, 11217-11223.	2.3	14
62	Forensic parameters of 19 X-STR polymorphisms in two Chinese populations. International Journal of Legal Medicine, 2017, 131, 975-977.	2.2	14
63	Association of CD40 â^'1C/T polymorphism in the 5â€²-untranslated region and chronic obstructive pulmonary disease. Clinica Chimica Acta, 2009, 408, 56-59.	1.1	13
64	The Association Between Interleukin-23 Receptor Gene Polymorphisms and Systemic Lupus Erythematosus. DNA and Cell Biology, 2010, 29, 79-82.	1.9	13
65	FLfinder: A novel software for the microhaplotype marker. Forensic Science International: Genetics Supplement Series, 2015, 5, e622-e624.	0.3	13
66	Influences of different RT-qPCR methods on forensic body fluid identification by microRNA. Forensic Science International: Genetics Supplement Series, 2015, 5, e295-e297.	0.3	13
67	Establishing a second-tier panel of 18 ancestry informative markers to improve ancestry distinctions among Asian populations. Forensic Science International: Genetics, 2019, 41, 159-167.	3.1	13
68	mRNA and microRNA stability validation of blood samples under different environmental conditions. Forensic Science International: Genetics, 2021, 55, 102567.	3.1	13
69	Population genetic analysis of 30 insertionâ€"deletion (INDEL) loci in a Qinghai Tibetan group using the Investigator DIPplex Kit. International Journal of Legal Medicine, 2019, 133, 1039-1041.	2.2	12
70	Intermittent fasting enhances hippocampal NPY expression to promote neurogenesis after traumatic brain injury. Nutrition, 2022, 97, 111621.	2.4	12
71	SNPâ€"STR polymorphism: A sensitive compound marker for forensic genetic applications. Forensic Science International: Genetics Supplement Series, 2013, 4, e206-e207.	0.3	11
72	A Novel SNP-STR System Based on a Capillary Electrophoresis Platform. Frontiers in Genetics, 2021, 12, 636821.	2.3	11

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73	Association of ADAM33 Polymorphisms and Susceptibility to Psoriasis. DNA and Cell Biology, 2010, 29, 435-439.	1.9	10
74	Association Between IRF-5 Polymorphisms and Risk of Acute Coronary Syndrome. DNA and Cell Biology, 2010, 29, 19-23.	1.9	10
75	Association of TNF-α Gene Promoter Polymorphisms With Susceptibility of Cervical Cancer in Southwest China. Laboratory Medicine, 2011, 42, 287-290.	1.2	10
76	A new approach to detect a set of SNP–SNP markers: Combining ARMS–PCR with SNaPshot technology. Electrophoresis, 2020, 41, 1189-1197.	2.4	10
77	Computer-aided superimposition of the frontal sinus via 3D reconstruction for comparative forensic identification. International Journal of Legal Medicine, 2021, 135, 1993-2001.	2.2	10
78	Assess the diversity of gut microbiota among healthy adults for forensic application. Microbial Cell Factories, 2022, 21, 46.	4.0	10
79	Association Between Single-Nucleotide Polymorphisms in Interleukin-12A and Risk of Chronic Obstructive Pulmonary Disease. DNA and Cell Biology, 2012, 31, 1475-1479.	1.9	9
80	Population study and mutation analysis for 28 short tandem repeat loci in southwest Chinese Han population. Journal of Clinical Forensic and Legal Medicine, 2016, 44, 10-13.	1.0	9
81	Integrated Bioinformatics Analysis for the Identification of Key Molecules and Pathways in the Hippocampus of Rats After Traumatic Brain Injury. Neurochemical Research, 2020, 45, 928-939.	3.3	9
82	Rapidly mutating Y-STRs study in Chinese Yi population. International Journal of Legal Medicine, 2019, 133, 45-50.	2.2	8
83	Morphological analysis of three-dimensionally reconstructed frontal sinuses from Chinese Han population using computed tomography. International Journal of Legal Medicine, 2021, 135, 1015-1023.	2.2	8
84	Expression of basigin in the early phase of acute myocardial ischemia in rats. Molecular Medicine Reports, 2013, 7, 1494-1500.	2.4	7
85	Validation of the Microreader 40Y ID System: a Y-STR multiplex for casework and database samples. International Journal of Legal Medicine, 2021, 135, 23-41.	2.2	7
86	Two X-Chromosome STR Loci DXS6803 and XS6793 Frequency Data in Chinese Population. Journal of Forensic Sciences, 2004, 49, 1-2.	1.6	7
87	Optogenetics for Understanding and Treating Brain Injury: Advances in the Field and Future Prospects. International Journal of Molecular Sciences, 2022, 23, 1800.	4.1	7
88	Micro RNA profiling for the detection and differentiation of body fluids in forensic stain analysis. Forensic Science International: Genetics Supplement Series, 2013, 4, e216-e217.	0.3	6
89	mRNA degradation pattern analysis in post-mortem normalized using the DNA. Forensic Science International: Genetics Supplement Series, 2013, 4, e266-e267.	0.3	6
90	Genotyping microhaplotype markers through massively parallel sequencing. Forensic Science International: Genetics Supplement Series, 2017, 6, e314-e316.	0.3	6

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91	Estimate the heterozygote balance of microhaplotype marker with massively parallel sequencing. Forensic Science International: Genetics Supplement Series, 2017, 6, e375-e376.	0.3	6
92	Genetic polymorphism of 21 non-CODIS STR loci in Chengdu Han population and its interpopulation analysis between 25 populations in China. Legal Medicine, 2018, 31, 14-16.	1.3	6
93	A functional variant in the flanking region of priâ€letâ€7f contributes to colorectal cancer risk in a Chinese population. Journal of Cellular Physiology, 2019, 234, 15717-15725.	4.1	6
94	CR-GAN: Automatic craniofacial reconstruction for personal identification. Pattern Recognition, 2022, 124, 108400.	8.1	6
95	Scientific Evidences of Calorie Restriction and Intermittent Fasting for Neuroprotection in Traumatic Brain Injury Animal Models: A Review of the Literature. Nutrients, 2022, 14, 1431.	4.1	6
96	An overview of SNP-SNP microhaplotypes in the 26 populations of the 1000 Genomes Project. International Journal of Legal Medicine, 2022, 136, 1211-1226.	2.2	6
97	Genetic data for 30 insertion/deletion polymorphisms in six Chinese populations with Qiagen Investigator DIPplex Kit. Forensic Science International: Genetics Supplement Series, 2013, 4, e268-e269.	0.3	5
98	Microhaplotype: Ability of personal identification and being ancestry informative marker. Forensic Science International: Genetics Supplement Series, 2017, 6, e442-e444.	0.3	5
99	SNP-STR analysis for non-invasive paternity test for fetus. Forensic Science International: Genetics Supplement Series, 2017, 6, e413-e414.	0.3	5
100	Systemic Lupus Erythematosus (SLE) Risk Factors: Novel Proteins Detected From Familial SLE Using Proteomics. Laboratory Medicine, 2009, 40, 408-411.	1.2	4
101	A comparison of malpractice lawsuits mediated and judged in court in China. Journal of Clinical Forensic and Legal Medicine, 2018, 54, 109-113.	1.0	4
102	Population genetic analysis of a 21-plex DIP panel in seven Chinese ethnic populations. International Journal of Legal Medicine, 2018, 132, 145-147.	2.2	4
103	Developmental validation of the Microreaderâ„¢ 20A ID system. Electrophoresis, 2019, 40, 3099-3107.	2.4	4
104	Detection of cellâ€free fetal DNA in maternal plasma using two types of compound markers. Electrophoresis, 2021, 42, 1158-1167.	2.4	4
105	A fully automated sex estimation for proximal femur X-ray images through deep learning detection and classification. Legal Medicine, 2022, 57, 102056.	1.3	4
106	A primary investigation on SNPs associated with eyelid traits of Chinese Han Adults. Forensic Science International: Genetics Supplement Series, 2015, 5, e669-e670.	0.3	3
107	Genetic polymorphisms of 17 Y-chromosomal STRs in the Chengdu Han population of China. International Journal of Legal Medicine, 2017, 131, 967-968.	2.2	3
108	Developing eight SNP-STR markers for DNA mixture detection. Forensic Science International: Genetics Supplement Series, 2017, 6, e351-e352.	0.3	3

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109	The expression of 10 candidate specific microRNA markers for human body fluid identification in animal buccal swabs. Forensic Science International, 2019, 300, e44-e49.	2.2	3
110	Validation of the Microreader 28A ID System: A 6â€dye multiplex amplification assay for forensic application. Electrophoresis, 2021, 42, 1928-1935.	2.4	3
111	Sequence Polymorphisms of the Mitochondrial DNA Control Region in 105 Chinese Han Population. Journal of Forensic Sciences, 2003, 48, 1-5.	1.6	3
112	Allele Frequency Distribution of Two X-Chromosomal STR Loci in Han Population in China. Journal of Forensic Sciences, 2004, 49, 1-2.	1.6	3
113	Allele Frequency Distribution of STR Loci D5S2848 in Four Populations. Journal of Forensic Sciences, 2004, 49, 1-2.	1.6	3
114	Set of 15 SNP-SNP Markers for Detection of Unbalanced Degraded DNA Mixtures and Noninvasive Prenatal Paternity Testing. Frontiers in Genetics, 2021, 12, 800598.	2.3	3
115	Association between BMP4 gene polymorphisms and eyelid traits in Chinese Han population. Forensic Science International: Genetics Supplement Series, 2017, 6, e355-e356.	0.3	2
116	Multiplex DNA methylation profiling by ARMS-PCR for body fluid identification. Forensic Science International: Genetics Supplement Series, 2019, 7, 820-822.	0.3	2
117	DNA-based eyelid trait prediction in Chinese Han population. International Journal of Legal Medicine, 2021, 135, 1743-1752.	2.2	2
118	The effect of infertile semen on the mRNAâ€based body fluid identification. Electrophoresis, 2021, 42, 1614-1622.	2.4	2
119	Feasibility of using probabilistic methods to analyse microRNA quantitative data in forensically relevant body fluids: a proof-of-principle study. International Journal of Legal Medicine, 2021, 135, 2247-2261.	2.2	2
120	Effect of infertile semen samples on mRNA-based body fluid identification by KLK3 and PRM1. Forensic Science International: Genetics Supplement Series, 2019, 7, 507-508.	0.3	2
121	Allele Frequency Distributions for 9 STR Loci of Tibetan Population in Chinese Tibet. Journal of Forensic Sciences, 2005, 50, 1-2.	1.6	2
122	Allele frequency distribution of two X-chromosomal STR loci in the Han population in China. International Congress Series, 2004, 1261, 145-147.	0.2	1
123	Construction and characterization of monoclonal antibodies specific for the R transactivator 185 of Epstein-Barr virus. Journal of Virological Methods, 2007, 144, 12-16.	2.1	1
124	The species specific of 3 microRNA markers in saliva. Forensic Science International: Genetics Supplement Series, 2015, 5, e674-e676.	0.3	1
125	Screening and confirmation of microRNA markers for distinguishing between menstrual and peripheral blood. Forensic Science International: Genetics Supplement Series, 2015, 5, e353-e355.	0.3	1
126	Mutation Study of 28 Autosomal STR Loci in Southwest Chinese Han Population. Forensic Science International: Genetics Supplement Series, 2015, 5, e298-e299.	0.3	1

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127	Postmortem interval (PMI) determination by profiling of HAF mRNA degradation using RT-qPCR. Forensic Science International: Genetics Supplement Series, 2017, 6, e182-e183.	0.3	1
128	A new approach to detect a set of SNP-SNP markers: Combining ARMS-PCR with SNaPshot technology. Forensic Science International: Genetics Supplement Series, 2019, 7, 150-151.	0.3	1
129	Population genetics of 27 Y-STRs for the Yi population from Liangshan Yi Autonomous Prefecture, China. International Journal of Legal Medicine, 2021, 135, 441-442.	2.2	1
130	STR Loci D19S400's Allele Frequency Distribution in Ten Populations. Journal of Forensic Sciences, 2005, 50, 1-1.	1.6	1
131	Application of MHanalyser software in the study of microhaplotypes in forensics. Forensic Science International: Genetics Supplement Series, 2019, 7, 271-273.	0.3	1
132	A population study of three Y-STR loci by multiplexing in Han population in Chengdu, China. International Congress Series, 2004, 1261, 254-256.	0.2	0
133	Allele Frequencies of D2S2960 and GATA149B10 in Two Populations. Journal of Forensic Sciences, 2006, 51, 1204-1204.	1.6	0
134	A 21-plex DIP panel's application in multinational Chinese population. Forensic Science International: Genetics Supplement Series, 2015, 5, e537-e538.	0.3	0
135	A novel system for forensic SNP analysis through PCR–ligase detection reaction. Forensic Science International: Genetics Supplement Series, 2015, 5, e231-e232.	0.3	0
136	Comparative study on methods of DNA genotyping between single piece of dandruff and EZ-tape. Forensic Science International: Genetics Supplement Series, 2017, 6, e244-e245.	0.3	0
137	Degradation of AIF in mouse heart tissue for estimating postmortem interval (PMI). Forensic Science International: Genetics Supplement Series, 2017, 6, e575-e576.	0.3	0
138	Allele Frequency Distribution of STR Loci D5S1486 in Three Populations. Journal of Forensic Sciences, 2005, 50, 1-2.	1.6	0
139	Allele Frequency Distributions for 15 STR Loci in Chinese Chengdu Han Population. Journal of Forensic Sciences, 2005, 50, 1-2.	1.6	0
140	Allele Frequency Distribution of STR Loci D5S814 in Four Populations. Journal of Forensic Sciences, 2005, 50, 1-2.	1.6	0
141	Allele Frequency Distribution of STR D5S819 in Four Populations. Journal of Forensic Sciences, 2005, 50, 1-2.	1.6	0
142	Allele Frequency Distribution of STR Loci D11S1390 and D11S2008 in Two Populations. Journal of Forensic Sciences, 2005, 50, 1-1.	1.6	0
143	Allele Frequency Distribution of STR Loci D5S2845 in Four Populations. Journal of Forensic Sciences, 2005, 50, 1-2.	1.6	0
144	A new proposed nomenclature for microhaplotypes. Forensic Science International: Genetics Supplement Series, 2019, 7, 813-815.	0.3	0

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145	What makes your â€œeyesâ€•look different?. Forensic Science International: Genetics Supplement Series, 2019, 7, 105-106.	0.3	0
146	Allele distributions for D21 S1435 and D21S2055 loci in two Chinese populations. Journal of Forensic Sciences, 2002, 47, 667-8.	1.6	0
147	Allele frequency distribution of STR loci D5S814 in four populations. Journal of Forensic Sciences, 2005, 50, 226-7.	1.6	0
148	STR loci D19S400's allele frequency distribution in ten populations. Journal of Forensic Sciences, 2005, 50, 725.	1.6	0