

Weibo Liang

List of Publications by Year in descending order

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148
papers

2,673
citations

196777

29
h-index

286692

43
g-index

150
all docs

150
docs citations

150
times ranked

2918
citing authors

#	ARTICLE	IF	CITATIONS
1	miR-212-5p attenuates ferroptotic neuronal death after traumatic brain injury by targeting Ptgs2. <i>Molecular Brain</i> , 2019, 12, 78.	1.3	123
2	The association of interleukin-16 polymorphisms with IL-16 serum levels and risk of colorectal and gastric cancer. <i>Carcinogenesis</i> , 2008, 30, 295-299.	1.3	95
3	The association between two polymorphisms in pre-miRNAs and breast cancer risk: a meta-analysis. <i>Breast Cancer Research and Treatment</i> , 2011, 125, 571-574.	1.1	91
4	Fermented dairy foods intake and risk of cancer. <i>International Journal of Cancer</i> , 2019, 144, 2099-2108.	2.3	79
5	Evaluation of the Microhaplotypes panel for DNA mixture analyses. <i>Forensic Science International: Genetics</i> , 2018, 35, 149-155.	1.6	64
6	RAD51 135G/C polymorphism and breast cancer risk: a meta-analysis from 21 studies. <i>Breast Cancer Research and Treatment</i> , 2011, 125, 827-835.	1.1	60
7	Effect of aging on the microstructure, hardness and chemical composition of dentin. <i>Archives of Oral Biology</i> , 2015, 60, 1811-1820.	0.8	59
8	A microhaplotypes panel for massively parallel sequencing analysis of DNA mixtures. <i>Forensic Science International: Genetics</i> , 2019, 40, 140-149.	1.6	58
9	A genetic variant in the promoter region of miR-34b/c is associated with a reduced risk of colorectal cancer. <i>Biological Chemistry</i> , 2013, 394, 415-420.	1.2	52
10	Forensic age estimation for pelvic X-ray images using deep learning. <i>European Radiology</i> , 2019, 29, 2322-2329.	2.3	51
11	Null Genotypes of GSTM1 and GSTT1 Contribute to Risk of Cervical Neoplasia: An Evidence-Based Meta-Analysis. <i>PLoS ONE</i> , 2011, 6, e20157.	1.1	49
12	Semen-specific miRNAs: Suitable for the distinction of infertile semen in the body fluid identification?. <i>Forensic Science International: Genetics</i> , 2018, 33, 161-167.	1.6	49
13	Development and application of a nonbinary SNP-based microhaplotype panel for paternity testing involving close relatives. <i>Forensic Science International: Genetics</i> , 2020, 46, 102255.	1.6	48
14	Identifying novel microhaplotypes for ancestry inference. <i>International Journal of Legal Medicine</i> , 2019, 133, 983-988.	1.2	47
15	Association of transforming growth factor- β 21 gene polymorphisms with genetic susceptibility to nasopharyngeal carcinoma. <i>Clinica Chimica Acta</i> , 2007, 380, 165-169.	0.5	46
16	A comparative study of insertion/deletion polymorphisms applied among Southwest, South and Northwest Chinese populations using InvestigatorA® DIPplex. <i>Forensic Science International: Genetics</i> , 2016, 21, 10-14.	1.6	45
17	Genetic diversity of 21 autosomal STR loci in the Han population from Sichuan province, Southwest China. <i>Forensic Science International: Genetics</i> , 2017, 31, e33-e35.	1.6	41
18	Interleukin-10 gene promoter polymorphisms and the risk of nasopharyngeal carcinoma. <i>Tissue Antigens</i> , 2007, 70, 12-17.	1.0	40

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19	IL-8 251A/T polymorphism is associated with decreased cancer risk among population-based studies: Evidence from a meta-analysis. <i>European Journal of Cancer</i> , 2010, 46, 1333-1343.	1.3	37
20	Hippocampus-dependent spatial memory impairment due to molar tooth loss is ameliorated by an enriched environment. <i>Archives of Oral Biology</i> , 2016, 61, 1-7.	0.8	37
21	Genetic Polymorphisms of the DNA Repair Gene and Risk of Nasopharyngeal Carcinoma. <i>DNA and Cell Biology</i> , 2007, 26, 491-496.	0.9	36
22	The IL-16 gene polymorphisms and the risk of the systemic lupus erythematosus. <i>Clinica Chimica Acta</i> , 2009, 403, 223-225.	0.5	36
23	Genetic polymorphism of Interleukin-16 and risk of nasopharyngeal carcinoma. <i>Clinica Chimica Acta</i> , 2009, 409, 132-135.	0.5	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. <i>Medicine (United States)</i> , 2015, 94, e1289.	0.4	36
25	Association Between Single-Nucleotide Polymorphisms in Pre-miRNAs and the Risk of Asthma in a Chinese Population. <i>DNA and Cell Biology</i> , 2011, 30, 919-923.	0.9	35
26	Characteristics of eight X-STR loci for forensic purposes in the Chinese population. <i>International Journal of Legal Medicine</i> , 2011, 125, 127-131.	1.2	35
27	Genotyping polymorphic microhaplotype markers through the Illumina® MiSeq platform for forensics. <i>Forensic Science International: Genetics</i> , 2019, 39, 1-7.	1.6	35
28	Identification of serum biomarkers for nasopharyngeal carcinoma by proteomic analysis. <i>Cancer</i> , 2008, 112, 544-551.	2.0	31
29	Screening and confirmation of microRNA markers for distinguishing between menstrual and peripheral blood. <i>Forensic Science International: Genetics</i> , 2017, 30, 24-33.	1.6	31
30	<i>Moringa oleifera</i> Lam and its Therapeutic Effects in Immune Disorders. <i>Frontiers in Pharmacology</i> , 2020, 11, 566783.	1.6	31
31	CTLA4 and CD86 gene polymorphisms and susceptibility to chronic obstructive pulmonary disease. <i>Human Immunology</i> , 2010, 71, 1141-1146.	1.2	28
32	The association between ATM D1853N polymorphism and breast cancer susceptibility: a meta-analysis. <i>Journal of Experimental and Clinical Cancer Research</i> , 2010, 29, 117.	3.5	28
33	Association between SNPs in pre-miRNA and risk of chronic obstructive pulmonary disease. <i>Clinical Biochemistry</i> , 2011, 44, 813-816.	0.8	28
34	Circular Ribonucleic Acid Expression Profile in Mouse Cortex after Traumatic Brain Injury. <i>Journal of Neurotrauma</i> , 2019, 36, 1018-1028.	1.7	28
35	Association of IL-1B Gene Polymorphisms with Nasopharyngeal Carcinoma in a Chinese Population. <i>Clinical Oncology</i> , 2008, 20, 207-211.	0.6	27
36	Association of single nucleotide polymorphisms in interleukin 12 (IL-12A and -B) with asthma in a Chinese population. <i>Human Immunology</i> , 2011, 72, 603-606.	1.2	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. <i>Clinical and Experimental Medicine</i> , 2018, 18, 79-87.	1.9	27
38	The xeroderma pigmentosum group C gene polymorphisms and genetic susceptibility of nasopharyngeal carcinoma. <i>Acta Oncologica</i> , 2008, 47, 379-384.	0.8	26
39	Association Between pri-miR-218 Polymorphism and Risk of Hepatocellular Carcinoma in a Han Chinese Population. <i>DNA and Cell Biology</i> , 2012, 31, 761-765.	0.9	26
40	Mutational analysis of 33 autosomal short tandem repeat (STR) loci in southwest Chinese Han population based on trio parentage testing. <i>Forensic Science International: Genetics</i> , 2016, 23, 86-90.	1.6	25
41	Genetic portrait of 27 Y-STR loci in the Tibetan ethnic population of the Qinghai province of China. <i>Forensic Science International: Genetics</i> , 2018, 34, e18-e19.	1.6	25
42	Two-person DNA mixture interpretation based on a novel set of SNP-STR markers. <i>Forensic Science International: Genetics</i> , 2018, 37, 37-45.	1.6	25
43	Expression difference of miR-10b and miR-135b between the fertile and infertile semen samples (p). <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e257-e259.	0.1	24
44	Population genetics for 17 Y-STR loci(AmpFISTR®Y-filer™) in Luzhou Han ethnic group. <i>Forensic Science International: Genetics</i> , 2013, 7, e23-e26.	1.6	23
45	Interactions of miR-34b/c and TP53 Polymorphisms on the Risk of Intracranial Aneurysm. <i>Clinical and Developmental Immunology</i> , 2012, 2012, 1-7.	3.3	22
46	Evaluation of the microhaplotype markers in kinship analysis. <i>Electrophoresis</i> , 2019, 40, 1091-1095.	1.3	22
47	Single nucleotide polymorphisms of VEGF gene and Psoriasis risk. <i>Journal of Dermatological Science</i> , 2008, 49, 263-265.	1.0	21
48	An investigation of a set of DIP-STR markers to detect unbalanced DNA mixtures among the southwest Chinese Han population. <i>Forensic Science International: Genetics</i> , 2017, 31, 34-39.	1.6	21
49	Association of Matrix Metalloproteinases 1, 7, and 9 Gene Polymorphisms with Genetic Susceptibility to Colorectal Carcinoma in a Han Chinese Population. <i>DNA and Cell Biology</i> , 2010, 29, 657-661.	0.9	20
50	Multi-Indel: A Microhaplotype Marker Can Be Typed Using Capillary Electrophoresis Platforms. <i>Frontiers in Genetics</i> , 2020, 11, 567082.	1.1	19
51	No association between epidermal growth factor and epidermal growth factor receptor polymorphisms and nasopharyngeal carcinoma. <i>Cancer Genetics and Cytogenetics</i> , 2008, 185, 69-73.	1.0	18
52	5 miRNA expression analyze in post-mortem interval (PMI) within 48h. <i>Forensic Science International: Genetics Supplement Series</i> , 2013, 4, e190-e191.	0.1	18
53	Development of a SNP-STRs multiplex for forensic identification. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e598-e600.	0.1	18
54	Postmortem interval determination using mRNA markers and DNA normalization. <i>International Journal of Legal Medicine</i> , 2020, 134, 149-157.	1.2	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. <i>Forensic Science International: Genetics</i> , 2020, 49, 102371.	1.6	16
56	Association of Tumor Necrosis Factor Gene Polymorphisms with Susceptibility to Dilated Cardiomyopathy in a Han Chinese Population. <i>DNA and Cell Biology</i> , 2010, 29, 625-628.	0.9	15
57	NGS technology makes microhaplotype a potential forensic marker. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e233-e234.	0.1	15
58	Microhaplotype identified and performed in genetic investigation using PCR-SSCP. <i>Forensic Science International: Genetics</i> , 2017, 28, e1-e7.	1.6	15
59	The association between dilated cardiomyopathy and RTN4 3'UTR insertion/deletion polymorphisms. <i>Clinica Chimica Acta</i> , 2009, 400, 21-24.	0.5	14
60	CD86 +1057 G/A Polymorphism and the Risk of Colorectal Cancer. <i>DNA and Cell Biology</i> , 2010, 29, 381-386.	0.9	14
61	Interactions of interleukin-12A and interleukin-12B polymorphisms on the risk of intracranial aneurysm. <i>Molecular Biology Reports</i> , 2012, 39, 11217-11223.	1.0	14
62	Forensic parameters of 19 X-STR polymorphisms in two Chinese populations. <i>International Journal of Legal Medicine</i> , 2017, 131, 975-977.	1.2	14
63	Association of CD40 C/T polymorphism in the 5'-untranslated region and chronic obstructive pulmonary disease. <i>Clinica Chimica Acta</i> , 2009, 408, 56-59.	0.5	13
64	The Association Between Interleukin-23 Receptor Gene Polymorphisms and Systemic Lupus Erythematosus. <i>DNA and Cell Biology</i> , 2010, 29, 79-82.	0.9	13
65	FLfinder: A novel software for the microhaplotype marker. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e622-e624.	0.1	13
66	Influences of different RT-qPCR methods on forensic body fluid identification by microRNA. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e295-e297.	0.1	13
67	Establishing a second-tier panel of 18 ancestry informative markers to improve ancestry distinctions among Asian populations. <i>Forensic Science International: Genetics</i> , 2019, 41, 159-167.	1.6	13
68	mRNA and microRNA stability validation of blood samples under different environmental conditions. <i>Forensic Science International: Genetics</i> , 2021, 55, 102567.	1.6	13
69	Population genetic analysis of 30 insertion-deletion (INDEL) loci in a Qinghai Tibetan group using the Investigator DIPplex Kit. <i>International Journal of Legal Medicine</i> , 2019, 133, 1039-1041.	1.2	12
70	Intermittent fasting enhances hippocampal NPY expression to promote neurogenesis after traumatic brain injury. <i>Nutrition</i> , 2022, 97, 111621.	1.1	12
71	SNP-STR polymorphism: A sensitive compound marker for forensic genetic applications. <i>Forensic Science International: Genetics Supplement Series</i> , 2013, 4, e206-e207.	0.1	11
72	A Novel SNP-STR System Based on a Capillary Electrophoresis Platform. <i>Frontiers in Genetics</i> , 2021, 12, 636821.	1.1	11

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73	Association of ADAM33 Polymorphisms and Susceptibility to Psoriasis. DNA and Cell Biology, 2010, 29, 435-439.	0.9	10
74	Association Between IRF-5 Polymorphisms and Risk of Acute Coronary Syndrome. DNA and Cell Biology, 2010, 29, 19-23.	0.9	10
75	Association of TNF- α Gene Promoter Polymorphisms With Susceptibility of Cervical Cancer in Southwest China. Laboratory Medicine, 2011, 42, 287-290.	0.8	10
76	A new approach to detect a set of SNP α -SNP markers: Combining ARMS α -PCR with SNaPshot technology. Electrophoresis, 2020, 41, 1189-1197.	1.3	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.	1.2	10
78	Assess the diversity of gut microbiota among healthy adults for forensic application. Microbial Cell Factories, 2022, 21, 46.	1.9	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.	0.9	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.	0.5	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.	1.6	9
82	Rapidly mutating Y-STRs study in Chinese Yi population. International Journal of Legal Medicine, 2019, 133, 45-50.	1.2	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.	1.2	8
84	Expression of basigin in the early phase of acute myocardial ischemia in rats. Molecular Medicine Reports, 2013, 7, 1494-1500.	1.1	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.	1.2	7
86	Two X-Chromosome STR Loci DXS6803 and XS6793 Frequency Data in Chinese Population. Journal of Forensic Sciences, 2004, 49, 1-2.	0.9	7
87	Optogenetics for Understanding and Treating Brain Injury: Advances in the Field and Future Prospects. International Journal of Molecular Sciences, 2022, 23, 1800.	1.8	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.	0.1	6
89	mRNA degradation pattern analysis in post-mortem normalized using the DNA. Forensic Science International: Genetics Supplement Series, 2013, 4, e266-e267.	0.1	6
90	Genotyping microhaplotype markers through massively parallel sequencing. Forensic Science International: Genetics Supplement Series, 2017, 6, e314-e316.	0.1	6

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91	Estimate the heterozygote balance of microhaplotype marker with massively parallel sequencing. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e375-e376.	0.1	6
92	Genetic polymorphism of 21 non-CODIS STR loci in Chengdu Han population and its interpopulation analysis between 25 populations in China. <i>Legal Medicine</i> , 2018, 31, 14-16.	0.6	6
93	A functional variant in the flanking region of priâ€tâ€f contributes to colorectal cancer risk in a Chinese population. <i>Journal of Cellular Physiology</i> , 2019, 234, 15717-15725.	2.0	6
94	CR-GAN: Automatic craniofacial reconstruction for personal identification. <i>Pattern Recognition</i> , 2022, 124, 108400.	5.1	6
95	Scientific Evidences of Calorie Restriction and Intermittent Fasting for Neuroprotection in Traumatic Brain Injury Animal Models: A Review of the Literature. <i>Nutrients</i> , 2022, 14, 1431.	1.7	6
96	An overview of SNP-SNP microhaplotypes in the 26 populations of the 1000 Genomes Project. <i>International Journal of Legal Medicine</i> , 2022, 136, 1211-1226.	1.2	6
97	Genetic data for 30 insertion/deletion polymorphisms in six Chinese populations with Qiagen Investigator DIPplex Kit. <i>Forensic Science International: Genetics Supplement Series</i> , 2013, 4, e268-e269.	0.1	5
98	Microhaplotype: Ability of personal identification and being ancestry informative marker. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e442-e444.	0.1	5
99	SNP-STR analysis for non-invasive paternity test for fetus. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e413-e414.	0.1	5
100	Systemic Lupus Erythematosus (SLE) Risk Factors: Novel Proteins Detected From Familial SLE Using Proteomics. <i>Laboratory Medicine</i> , 2009, 40, 408-411.	0.8	4
101	A comparison of malpractice lawsuits mediated and judged in court in China. <i>Journal of Clinical Forensic and Legal Medicine</i> , 2018, 54, 109-113.	0.5	4
102	Population genetic analysis of a 21-plex DIP panel in seven Chinese ethnic populations. <i>International Journal of Legal Medicine</i> , 2018, 132, 145-147.	1.2	4
103	Developmental validation of the Microreaderâ„¢ 20A ID system. <i>Electrophoresis</i> , 2019, 40, 3099-3107.	1.3	4
104	Detection of cellâ€free fetal DNA in maternal plasma using two types of compound markers. <i>Electrophoresis</i> , 2021, 42, 1158-1167.	1.3	4
105	A fully automated sex estimation for proximal femur X-ray images through deep learning detection and classification. <i>Legal Medicine</i> , 2022, 57, 102056.	0.6	4
106	A primary investigation on SNPs associated with eyelid traits of Chinese Han Adults. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e669-e670.	0.1	3
107	Genetic polymorphisms of 17 Y-chromosomal STRs in the Chengdu Han population of China. <i>International Journal of Legal Medicine</i> , 2017, 131, 967-968.	1.2	3
108	Developing eight SNP-STR markers for DNA mixture detection. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e351-e352.	0.1	3

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109	The expression of 10 candidate specific microRNA markers for human body fluid identification in animal buccal swabs. <i>Forensic Science International</i> , 2019, 300, e44-e49.	1.3	3
110	Validation of the Microreader 28A ID System: A 6â€œ dye multiplex amplification assay for forensic application. <i>Electrophoresis</i> , 2021, 42, 1928-1935.	1.3	3
111	Sequence Polymorphisms of the Mitochondrial DNA Control Region in 105 Chinese Han Population. <i>Journal of Forensic Sciences</i> , 2003, 48, 1-5.	0.9	3
112	Allele Frequency Distribution of Two X-Chromosomal STR Loci in Han Population in China. <i>Journal of Forensic Sciences</i> , 2004, 49, 1-2.	0.9	3
113	Allele Frequency Distribution of STR Loci D5S2848 in Four Populations. <i>Journal of Forensic Sciences</i> , 2004, 49, 1-2.	0.9	3
114	Set of 15 SNP-SNP Markers for Detection of Unbalanced Degraded DNA Mixtures and Noninvasive Prenatal Paternity Testing. <i>Frontiers in Genetics</i> , 2021, 12, 800598.	1.1	3
115	Association between BMP4 gene polymorphisms and eyelid traits in Chinese Han population. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e355-e356.	0.1	2
116	Multiplex DNA methylation profiling by ARMS-PCR for body fluid identification. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 820-822.	0.1	2
117	DNA-based eyelid trait prediction in Chinese Han population. <i>International Journal of Legal Medicine</i> , 2021, 135, 1743-1752.	1.2	2
118	The effect of infertile semen on the mRNAâ€œbased body fluid identification. <i>Electrophoresis</i> , 2021, 42, 1614-1622.	1.3	2
119	Feasibility of using probabilistic methods to analyse microRNA quantitative data in forensically relevant body fluids: a proof-of-principle study. <i>International Journal of Legal Medicine</i> , 2021, 135, 2247-2261.	1.2	2
120	Effect of infertile semen samples on mRNA-based body fluid identification by KLK3 and PRM1. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 507-508.	0.1	2
121	Allele Frequency Distributions for 9 STR Loci of Tibetan Population in Chinese Tibet. <i>Journal of Forensic Sciences</i> , 2005, 50, 1-2.	0.9	2
122	Allele frequency distribution of two X-chromosomal STR loci in the Han population in China. <i>International Congress Series</i> , 2004, 1261, 145-147.	0.2	1
123	Construction and characterization of monoclonal antibodies specific for the R transactivator 185 of Epstein-Barr virus. <i>Journal of Virological Methods</i> , 2007, 144, 12-16.	1.0	1
124	The species specific of 3 microRNA markers in saliva. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e674-e676.	0.1	1
125	Screening and confirmation of microRNA markers for distinguishing between menstrual and peripheral blood. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e353-e355.	0.1	1
126	Mutation Study of 28 Autosomal STR Loci in Southwest Chinese Han Population. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e298-e299.	0.1	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.	0.1	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.	0.1	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.	1.2	1
130	STR Loci D19S400's Allele Frequency Distribution in Ten Populations. Journal of Forensic Sciences, 2005, 50, 1-1.	0.9	1
131	Application of MHanalyzer software in the study of microhaplotypes in forensics. Forensic Science International: Genetics Supplement Series, 2019, 7, 271-273.	0.1	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.	0.9	0
134	A 21-plex DIP panel's application in multinational Chinese population. Forensic Science International: Genetics Supplement Series, 2015, 5, e537-e538.	0.1	0
135	A novel system for forensic SNP analysis through PCR's ligase detection reaction. Forensic Science International: Genetics Supplement Series, 2015, 5, e231-e232.	0.1	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.	0.1	0
137	Degradation of AIF in mouse heart tissue for estimating postmortem interval (PMI). Forensic Science International: Genetics Supplement Series, 2017, 6, e575-e576.	0.1	0
138	Allele Frequency Distribution of STR Loci D5S1486 in Three Populations. Journal of Forensic Sciences, 2005, 50, 1-2.	0.9	0
139	Allele Frequency Distributions for 15 STR Loci in Chinese Chengdu Han Population. Journal of Forensic Sciences, 2005, 50, 1-2.	0.9	0
140	Allele Frequency Distribution of STR Loci D5S814 in Four Populations. Journal of Forensic Sciences, 2005, 50, 1-2.	0.9	0
141	Allele Frequency Distribution of STR D5S819 in Four Populations. Journal of Forensic Sciences, 2005, 50, 1-2.	0.9	0
142	Allele Frequency Distribution of STR Loci D11S1390 and D11S2008 in Two Populations. Journal of Forensic Sciences, 2005, 50, 1-1.	0.9	0
143	Allele Frequency Distribution of STR Loci D5S2845 in Four Populations. Journal of Forensic Sciences, 2005, 50, 1-2.	0.9	0
144	A new proposed nomenclature for microhaplotypes. Forensic Science International: Genetics Supplement Series, 2019, 7, 813-815.	0.1	0

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145	What makes your "eyes" look different?. Forensic Science International: Genetics Supplement Series, 2019, 7, 105-106.	0.1	0
146	Allele distributions for D21 S1435 and D21S2055 loci in two Chinese populations. Journal of Forensic Sciences, 2002, 47, 667-8.	0.9	0
147	Allele frequency distribution of STR loci D5S814 in four populations. Journal of Forensic Sciences, 2005, 50, 226-7.	0.9	0
148	STR loci D19S400's allele frequency distribution in ten populations. Journal of Forensic Sciences, 2005, 50, 725.	0.9	0