

Fei Guo

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

624
citations

686830

13
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642321

23
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23
all docs

23
docs citations

23
times ranked

446
citing authors

#	ARTICLE	IF	CITATIONS
1	Mitochondrial genome sequencing with short overlapping amplicons on MiSeq FGx system. <i>Forensic Sciences Research</i> , 2022, 7, 142-153.	0.9	5
2	Genetic polymorphisms of 21 STR loci of GoldenGate DNA ID 22NC kit in five ethnic groups of China. <i>Forensic Sciences Research</i> , 2019, 4, 348-350.	0.9	3
3	StatsX v2.0: the interactive graphical software for population statistics on X-STR. <i>International Journal of Legal Medicine</i> , 2019, 133, 39-44.	1.2	36
4	Allele frequencies of 15 autosomal STR loci in the Yi ethnic minority from Yunnan Province in the far southwest of China. <i>Forensic Science International: Genetics</i> , 2017, 26, e23-e24.	1.6	9
5	Population genetics for 17 Y-STR loci in Hui ethnic minority from Liaoning Province, Northeast China. <i>Forensic Science International: Genetics</i> , 2017, 28, e36-e37.	1.6	18
6	Population genetics for 17 Y-STR loci in Northern Han Chinese from Liaoning Province, Northeast China. <i>Forensic Science International: Genetics</i> , 2017, 29, e35-e37.	1.6	8
7	Population genetic data for 27 Y-STR loci in the Zhuang ethnic minority from Guangxi Zhuang Autonomous Region in the south of China. <i>Forensic Science International: Genetics</i> , 2017, 27, 182-183.	1.6	28
8	Genetic variation of 17 autosomal STR loci in the Zhuang ethnic minority from Guangxi Zhuang Autonomous Region in the south of China. <i>Forensic Science International: Genetics</i> , 2017, 28, e51-e52.	1.6	13
9	Massively parallel sequencing of forensic STRs and SNPs using the Illumina® ForenSeq® DNA Signature Prep Kit on the MiSeq FGx® Forensic Genomics System. <i>Forensic Science International: Genetics</i> , 2017, 31, 135-148.	1.6	110
10	Genetic polymorphism of 17 autosomal STR loci in the Lahu ethnic minority from Yunnan Province, Southwest China. <i>Forensic Science International: Genetics</i> , 2017, 31, e52-e53.	1.6	12
11	Allele frequencies of 17 autosomal STR loci in the Va ethnic minority from Yunnan Province, Southwest China. <i>International Journal of Legal Medicine</i> , 2017, 131, 1251-1252.	1.2	4
12	Genetic variation of 17 autosomal STR loci in the Dong ethnic minority from Guangxi Zhuang Autonomous Region, South China. <i>International Journal of Legal Medicine</i> , 2017, 131, 1537-1538.	1.2	5
13	Population genetic data for 12 X-STR loci in the Northern Han Chinese and StatsX package as tools for population statistics on X-STR. <i>Forensic Science International: Genetics</i> , 2017, 26, e1-e8.	1.6	24
14	Evaluation of the Early Access STR Kit v1 on the Ion Torrent PGM® platform. <i>Forensic Science International: Genetics</i> , 2016, 23, 111-120.	1.6	47
15	Next generation sequencing of SNPs using the HID-Ion AmpliSeq® Identity Panel on the Ion Torrent PGM® platform. <i>Forensic Science International: Genetics</i> , 2016, 25, 73-84.	1.6	63
16	Strategies for complete mitochondrial genome sequencing on Ion Torrent PGM® platform in forensic sciences. <i>Forensic Science International: Genetics</i> , 2016, 22, 11-21.	1.6	46
17	Population genetics for 17 Y-STR loci in Korean ethnic minority from Liaoning Province, Northeast China. <i>Forensic Science International: Genetics</i> , 2016, 22, e9-e11.	1.6	12
18	Population genetics for 17 Y-STR loci in Mongolian ethnic minority from Liaoning Province, Northeast China. <i>Forensic Science International: Genetics</i> , 2015, 17, 153-154.	1.6	23

#	ARTICLE	IF	CITATIONS
19	Population genetics of 17 Y-STR loci in Xibe ethnic minority from Liaoning Province, Northeast China. Forensic Science International: Genetics, 2015, 16, 86-87.	1.6	24
20	Population data for 15 autosomal STR loci in the Miao ethnic minority from Guizhou Province, Southwest China. Forensic Science International: Genetics, 2015, 16, e3-e4.	1.6	37
21	Population data of 17 Y-STR loci in Nanyang Han population from Henan Province, Central China. Forensic Science International: Genetics, 2014, 13, 145-146.	1.6	17