

Hongling Zhang

List of Publications by Year in descending order

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Version: 2024-02-01

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papers

286
citations

1039406

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citing authors

#	ARTICLE	IF	CITATIONS
1	Population genetic data and mutations of 22 autosomal STR loci in Guizhou Han population. <i>Forensic Science International: Genetics</i> , 2017, 29, e29-e30.	1.6	33
2	Genetic diversity, structure and forensic characteristics of Hmong-Mien-speaking Miao revealed by autosomal insertion/deletion markers. <i>Molecular Genetics and Genomics</i> , 2019, 294, 1487-1498.	1.0	32
3	Population genetics, diversity and forensic characteristics of Tai-Kadai-speaking Bouyei revealed by insertion/deletions markers. <i>Molecular Genetics and Genomics</i> , 2019, 294, 1343-1357.	1.0	32
4	Genomic Insights Into the Admixture History of Mongolic- and Tungusic-Speaking Populations From Southwestern East Asia. <i>Frontiers in Genetics</i> , 2021, 12, 685285.	1.1	23
5	Genetic profile of 17 Y chromosome STRs in the Guizhou Han population of southwestern China. <i>Forensic Science International: Genetics</i> , 2016, 25, e6-e7.	1.6	21
6	Genetic profile of 23 Y chromosomal STR loci in Guizhou Shui population, southwest China. <i>Forensic Science International: Genetics</i> , 2017, 28, e16-e17.	1.6	17
7	Forensic Features and Population Genetic Structure of Dong, Yi, Han, and Chuanqing Human Populations in Southwest China Inferred From Insertion/Deletion Markers. <i>Frontiers in Genetics</i> , 2020, 11, 360.	1.1	17
8	Genetic structure and forensic characterisation of 19 X-chromosomal STR loci in Guizhou Sui population. <i>Annals of Human Biology</i> , 2019, 46, 246-253.	0.4	16
9	Population genetic analysis of 36 Y-chromosomal STRs yields comprehensive insights into the forensic features and phylogenetic relationship of Chinese Tai-Kadai-speaking Bouyei. <i>PLoS ONE</i> , 2019, 14, e0224601.	1.1	15
10	Forensic genetic polymorphisms and population structure of the Guizhou Bouyei people based on 19 X-STR loci. <i>Annals of Human Biology</i> , 2019, 46, 574-580.	0.4	12
11	Genetic structure and forensic characterisation of 36 Y-chromosomal STR loci in Hmong-Mien-speaking Miao population. <i>Annals of Human Biology</i> , 2020, 47, 541-548.	0.4	11
12	Male-Dominated Migration and Massive Assimilation of Indigenous East Asians in the Formation of Muslim Hui People in Southwest China. <i>Frontiers in Genetics</i> , 2020, 11, 618614.	1.1	11
13	Fine-Scale Population Admixture Landscape of Tai-Kadai-Speaking Maonan in Southwest China Inferred From Genome-Wide SNP Data. <i>Frontiers in Genetics</i> , 2022, 13, 815285.	1.1	10
14	Genome-wide analysis of unrecognised ethnic group Chuanqing people revealing a close affinity with Southern Han Chinese. <i>Annals of Human Biology</i> , 2020, 47, 465-471.	0.4	9
15	The mitochondrial DNA control region sequences from the Chinese Miao population of southeastern China. <i>Annals of Human Biology</i> , 2019, 46, 606-609.	0.4	7
16	Population genetic data of 22 autosomal STRs in the Guizhou Miao population, southwestern China. <i>Forensic Science International: Genetics</i> , 2018, 32, e7-e8.	1.6	6
17	The mitochondrial DNA control region sequences from the Chinese sui population of southwestern China. <i>Annals of Human Biology</i> , 2021, 48, 635-640.	0.4	4
18	Population genetic data of 22 autosomal STRs in Guizhou Bouyei population, Southwestern China. <i>Forensic Science International: Genetics</i> , 2018, 33, e11-e12.	1.6	3

#	ARTICLE	IF	CITATIONS
19	X-chromosomal STRs for genetic composition analysis of Guizhou Dong group and its phylogenetic relationships with other reference populations. <i>Annals of Human Biology</i> , 2021, , 1-9.	0.4	2
20	Developmental Validation of a Rapidly Mutating Y-STR Panel Labeled by Six Fluoresceins for Forensic Research. <i>Frontiers in Genetics</i> , 2022, 13, 777440.	1.1	0