Forensic characterization of 15 autosomal STRs in four pand genetic relationships with neighboring populations

Scientific Reports 8, 4673

DOI: 10.1038/s41598-018-22975-6

Citation Report

#	Article	IF	CITATIONS
1	The Ong Be language-speaking population in Hainan Island: genetic diversity, phylogenetic characteristics and reflections on ethnicity. Molecular Biology Reports, 2019, 46, 4095-4103.	1.0	11
2	Population genetics of 15 autosomal STR loci in the Han population of Ili Kazakh Autonomous Prefecture, Northwestern China. Annals of Human Genetics, 2019, 83, 318-324.	0.3	3
3	Allele and haplotype frequencies of 19 X-STRs in the Kyrgyz and Han populations from Kizilsu Prefecture. Forensic Science International: Genetics, 2019, 40, e259-e261.	1.6	6
4	Population data of 23 Y STRs from Manchu population of Liaoning Province, Northeast China. International Journal of Legal Medicine, 2019, 133, 785-788.	1.2	12
5	Genetic analysis of 12 X-STRs for forensic purposes in Liaoning Manchu population from China. Gene, 2019, 683, 153-158.	1.0	10
6	Phylogenetic relationship and genetic history of Central Asian Kazakhs inferred from Y-chromosome and autosomal variations. Molecular Genetics and Genomics, 2020, 295, 221-231.	1.0	4
7	Genetic characterization of 15 autosomal STRs in the interior Sindhi population of Pakistan and their phylogenetic relationship with other populations. International Journal of Immunogenetics, 2020, 47, 149-157.	0.8	2
8	Population genetic portrait of Pakistani Lahore-Christians based on 32 STR loci. Scientific Reports, 2020, 10, 18960.	1.6	3
9	Genetic structure and forensic characteristics of the Korean population revealed by GoldenEye 20A. Annals of Human Biology, 2020, 47, 560-563.	0.4	1
11	Genetic structure of the ethnic Lao groups from mainland Southeast Asia revealed by forensic microsatellites. Annals of Human Genetics, 2020, 84, 357-369.	0.3	7
12	Genetic polymorphism study on 21 autosomal STR loci of populations from six geographical regions in the Kingdom of Thailand. Forensic Science International: Genetics, 2020, 47, 102297.	1.6	0
13	Updated population genetic data of 15 autosomal STR loci in a Shandong Han population from East China and genetic relationships among 26 Chinese populations. Annals of Human Biology, 2020, 47, 472-477.	0.4	2
14	Genetic structure and forensic characteristics of the Kyrgyz population from Kizilsu Kirghiz autonomous prefecture based on autosomal DIPs. International Journal of Legal Medicine, 2022, 136, 539-541.	1.2	3
15	Forensic features and genetic structure of 20 autosomal STR loci in the Han population of Ningde City, Southeastern China. Annals of Human Biology, 2021, 48, 56-61.	0.4	1
16	Useful autosomal STR marker sets for forensic and paternity applications in the Central Indian population. Annals of Human Biology, 2021, 48, 37-48.	0.4	10
17	Polymorphic investigation and interpopulation genetic differentiation analyses of 20 STR loci in Inner Mongolia Han population. Gene Reports, 2021, 25, 101373.	0.4	O
18	Evaluation of diallelic STR markers with inter-population allelic database for their usefulness in paternity trios in the Central Indian population. Annals of Human Biology, 2021, 48, 605-613.	0.4	2
19	The Heart of Silk Road "Xinjiang,―lts Genetic Portray, and Forensic Parameters Inferred From Autosomal STRs. Frontiers in Genetics, 2021, 12, 760760.	1.1	4

#	Article	IF	CITATIONS
20	Genetic polymorphism and forensic application of 23 autosomal STR loci in the Han population of Panjin City, Liaoning Province, Northeastern China. Annals of Human Biology, 2022, 49, 254-259.	0.4	1
21	Genetic characterization of the highlander Tibetan population from Qinghai-Tibet Plateau revealed by X chromosomal STRs. PLoS ONE, 2022, 17, e0271769.	1.1	4
22	Systematic selections and forensic application evaluations of 111 individual identification SNPs in the Chinese Inner Mongolia Manchu group. Frontiers in Genetics, $0, 13, .$	1.1	0