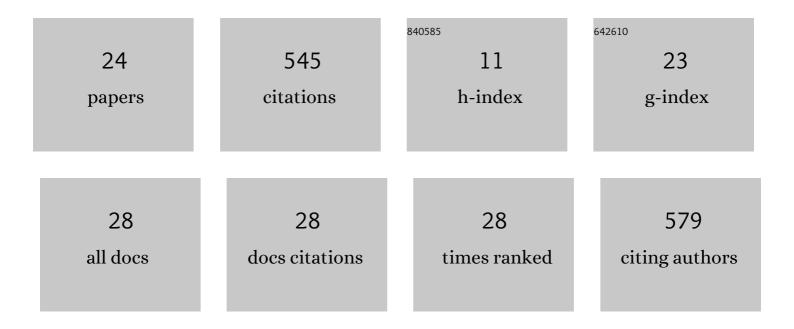
Shengjie Nie

List of Publications by Year in descending order

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SHENCUE NIE

#	Article	IF	CITATIONS
1	Genetic polymorphism investigation of 16 X-STR loci in the Bai ethnic minority in Yunnan Province, Southwest China. International Journal of Legal Medicine, 2022, 136, 543-545.	1.2	5
2	Genetic polymorphisms of 16 X-STR loci in the Hani population from Southwest China. Forensic Sciences Research, 2022, 7, 196-201.	0.9	3
3	Genetic polymorphisms of 16 X-STR loci analyzed in the Han population of Yunnan Province, Southwest China. Legal Medicine, 2022, 54, 101974.	0.6	1
4	Sex-dependent DNA hypermethylation of SLC6A4 in patients with schizophrenia. Neuroscience Letters, 2022, 769, 136394.	1.0	4
5	Insights into AIM-InDel diversities in Yunnan Miao and Hani ethnic groups of China for forensic and population genetic purposes. Hereditas, 2022, 159, 22.	0.5	1
6	Glucocorticoid receptor gene (NR3C1) is hypermethylated in adult males with aggressive behaviour. International Journal of Legal Medicine, 2021, 135, 43-51.	1.2	8
7	Association of mineralocorticoid receptor gene (NR3C2) hypermethylation in adult males with aggressive behavior. Behavioural Brain Research, 2021, 398, 112980.	1.2	5
8	Forensic genetic polymorphisms of 16 X-STR loci in the Yunnan Miao population and their relationship to other Chinese groups. Legal Medicine, 2021, 53, 101961.	0.6	4
9	Sex-dependent association of mineralocorticoid receptor gene (NR3C2) DNA methylation and schizophrenia. Psychiatry Research, 2020, 292, 113318.	1.7	11
10	Methylation of the MAOA promoter is associated with schizophrenia. Annals of Translational Medicine, 2020, 8, 864-864.	0.7	4
11	DNA Methylation Analysis of the NR3C1 Gene in Patients with Schizophrenia. Journal of Molecular Neuroscience, 2020, 70, 1177-1185.	1.1	17
12	The massive assimilation of indigenous East Asian populations in the origin of Muslim Hui people inferred from paternal Y chromosome. American Journal of Physical Anthropology, 2019, 169, 341-347.	2.1	16
13	Forensic features for Yunnan Lisu ethnic minority and phylogenetic structure exploration among 26 Chinese populations. International Journal of Legal Medicine, 2019, 133, 103-104.	1.2	4
14	Population data of 23 autosomal STR loci in the Chinese Han population from Guangdong Province in southern China. International Journal of Legal Medicine, 2018, 132, 133-135.	1.2	21
15	Population data and mutation rates of 20 autosomal STR loci in a Chinese Han population from Yunnan Province, Southwest China. International Journal of Legal Medicine, 2018, 132, 1083-1085.	1.2	15
16	Genetic polymorphisms of 24 Y-STR loci in Hani ethnic minority from Yunnan Province, Southwest China. International Journal of Legal Medicine, 2017, 131, 1235-1237.	1.2	10
17	Population data for 20 autosomal STR loci in the Yi ethnic minority from Yunnan Province, Southwest China. Forensic Science International: Genetics, 2017, 28, e43-e44.	1.6	24
18	Genetic analysis of 20 autosomal STR loci in the Miao ethnic group from Yunnan Province, Southwest China. Forensic Science International: Genetics, 2017, 28, e28-e29.	1.6	27

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#	Article	IF	CITATIONS
19	Genetic analysis of 24 Y-STR loci in the Miao ethnic minority from Yunnan Province, southwestern China. Forensic Science International: Genetics, 2017, 28, e30-e32.	1.6	9
20	Genetic polymorphisms of 20 autosomal STR loci in the Vietnamese population from Yunnan Province, Southwest China. International Journal of Legal Medicine, 2017, 131, 661-662.	1.2	16
21	Genetic variation of 20 autosomal STR loci in three ethnic groups (Zhuang, Dai and Hani) in the Yunnan province of southwestern China. Forensic Science International: Genetics, 2017, 31, e41-e42.	1.6	20
22	A NR3C2 haplotype increases the risk of alcoholism in schizophrenic patients in Han Chinese population. Psychiatry Research, 2015, 229, 1057-1058.	1.7	3
23	The CRHR1 Gene Contributes to Genetic Susceptibility of Aggressive Behavior Towards Others in Chinese Southwest Han Population. Journal of Molecular Neuroscience, 2014, 52, 481-486.	1.1	22
24	A global analysis of Y-chromosomal haplotype diversity for 23 STR loci. Forensic Science International: Genetics, 2014, 12, 12-23.	1.6	214