

# Shengjie Nie

## List of Publications by Year in descending order

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Version: 2024-02-01

24  
papers

545  
citations

840585

11  
h-index

642610

23  
g-index

28  
all docs

28  
docs citations

28  
times ranked

579  
citing authors

#	ARTICLE	IF	CITATIONS
1	A global analysis of Y-chromosomal haplotype diversity for 23 STR loci. <i>Forensic Science International: Genetics</i> , 2014, 12, 12-23.	1.6	214
2	Genetic analysis of 20 autosomal STR loci in the Miao ethnic group from Yunnan Province, Southwest China. <i>Forensic Science International: Genetics</i> , 2017, 28, e28-e29.	1.6	27
3	Population data for 20 autosomal STR loci in the Yi ethnic minority from Yunnan Province, Southwest China. <i>Forensic Science International: Genetics</i> , 2017, 28, e43-e44.	1.6	24
4	The CRHR1 Gene Contributes to Genetic Susceptibility of Aggressive Behavior Towards Others in Chinese Southwest Han Population. <i>Journal of Molecular Neuroscience</i> , 2014, 52, 481-486.	1.1	22
5	Population data of 23 autosomal STR loci in the Chinese Han population from Guangdong Province in southern China. <i>International Journal of Legal Medicine</i> , 2018, 132, 133-135.	1.2	21
6	Genetic variation of 20 autosomal STR loci in three ethnic groups (Zhuang, Dai and Hani) in the Yunnan province of southwestern China. <i>Forensic Science International: Genetics</i> , 2017, 31, e41-e42.	1.6	20
7	DNA Methylation Analysis of the NR3C1 Gene in Patients with Schizophrenia. <i>Journal of Molecular Neuroscience</i> , 2020, 70, 1177-1185.	1.1	17
8	Genetic polymorphisms of 20 autosomal STR loci in the Vietnamese population from Yunnan Province, Southwest China. <i>International Journal of Legal Medicine</i> , 2017, 131, 661-662.	1.2	16
9	The massive assimilation of indigenous East Asian populations in the origin of Muslim Hui people inferred from paternal Y chromosome. <i>American Journal of Physical Anthropology</i> , 2019, 169, 341-347.	2.1	16
10	Population data and mutation rates of 20 autosomal STR loci in a Chinese Han population from Yunnan Province, Southwest China. <i>International Journal of Legal Medicine</i> , 2018, 132, 1083-1085.	1.2	15
11	Sex-dependent association of mineralocorticoid receptor gene (NR3C2) DNA methylation and schizophrenia. <i>Psychiatry Research</i> , 2020, 292, 113318.	1.7	11
12	Genetic polymorphisms of 24 Y-STR loci in Hani ethnic minority from Yunnan Province, Southwest China. <i>International Journal of Legal Medicine</i> , 2017, 131, 1235-1237.	1.2	10
13	Genetic analysis of 24 Y-STR loci in the Miao ethnic minority from Yunnan Province, southwestern China. <i>Forensic Science International: Genetics</i> , 2017, 28, e30-e32.	1.6	9
14	Glucocorticoid receptor gene (NR3C1) is hypermethylated in adult males with aggressive behaviour. <i>International Journal of Legal Medicine</i> , 2021, 135, 43-51.	1.2	8
15	Genetic polymorphism investigation of 16 X-STR loci in the Bai ethnic minority in Yunnan Province, Southwest China. <i>International Journal of Legal Medicine</i> , 2022, 136, 543-545.	1.2	5
16	Association of mineralocorticoid receptor gene (NR3C2) hypermethylation in adult males with aggressive behavior. <i>Behavioural Brain Research</i> , 2021, 398, 112980.	1.2	5
17	Forensic features for Yunnan Lisu ethnic minority and phylogenetic structure exploration among 26 Chinese populations. <i>International Journal of Legal Medicine</i> , 2019, 133, 103-104.	1.2	4
18	Methylation of the MAOA promoter is associated with schizophrenia. <i>Annals of Translational Medicine</i> , 2020, 8, 864-864.	0.7	4

#	ARTICLE	IF	CITATIONS
19	Forensic genetic polymorphisms of 16 X-STR loci in the Yunnan Miao population and their relationship to other Chinese groups. <i>Legal Medicine</i> , 2021, 53, 101961.	0.6	4
20	Sex-dependent DNA hypermethylation of SLC6A4 in patients with schizophrenia. <i>Neuroscience Letters</i> , 2022, 769, 136394.	1.0	4
21	A NR3C2 haplotype increases the risk of alcoholism in schizophrenic patients in Han Chinese population. <i>Psychiatry Research</i> , 2015, 229, 1057-1058.	1.7	3
22	Genetic polymorphisms of 16 X-STR loci in the Hani population from Southwest China. <i>Forensic Sciences Research</i> , 2022, 7, 196-201.	0.9	3
23	Genetic polymorphisms of 16 X-STR loci analyzed in the Han population of Yunnan Province, Southwest China. <i>Legal Medicine</i> , 2022, 54, 101974.	0.6	1
24	Insights into AIM-InDel diversities in Yunnan Miao and Hani ethnic groups of China for forensic and population genetic purposes. <i>Hereditas</i> , 2022, 159, 22.	0.5	1