Jianhui Xie

List of Publications by Year in descending order

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623734 642732 37 599 14 23 h-index citations g-index papers 40 40 40 738 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Identification and assessment of a subset of Y-SNPs with recurrent mutation for forensic purpose. Forensic Science International, 2022, 334, 111270.	2.2	2
2	Application of machine learning for ancestry inference using multi-InDel markers. Forensic Science International: Genetics, 2022, 59, 102702.	3.1	3
3	Evaluation of one-step RT-PCR multiplex assay for body fluid identification. International Journal of Legal Medicine, 2021, 135, 1727-1735.	2.2	8
4	Calculation of the Paternity Index for STR with tri-allelic patterns in paternity testing. Forensic Science International, 2021, 324, 110832.	2.2	5
5	Variants in linkage status at D5S818 detected by multiple STR kits comparison and Sanger sequencing. Molecular Genetics & D5S818 detected by multiple STR kits comparison and Sanger sequencing.	1.2	2
6	Identification and Characterization of Nine Novel X-Chromosomal Short Tandem Repeats on Xp21.1, Xq21.31, and Xq23 Regions. Frontiers in Genetics, 2021, 12, 784605.	2.3	3
7	Assessing Autosomal InDel Loci With Multiple Insertions or Deletions of Random DNA Sequences in Human Genome. Frontiers in Genetics, 2021, 12, 809815.	2.3	2
8	Genetic analysis of type 2 tri-allelic pattern at TPOX locus in the Chinese Han population. Molecular Genetics and Genomics, 2020, 295, 933-939.	2.1	3
9	Development of a multiplex system for the identification of forensically relevant body fluids. Forensic Science International: Genetics, 2020, 47, 102312.	3.1	21
10	Genetic analysis of tri-allelic patterns at the CODIS STR loci. Molecular Genetics and Genomics, 2020, 295, 1263-1268.	2.1	6
11	A 16-plex Y-SNP typing system based on allele-specific PCR for the genotyping of Chinese Y-chromosomal haplogroups. Legal Medicine, 2020, 46, 101720.	1.3	9
12	Characterizing Y-STRs in the Evaluation of Population Differentiation Using the Mean of Allele Frequency Difference between Populations. Genes, 2020, 11, 566.	2.4	2
13	Forensic age estimation based on the pigmentation in the costal cartilage from human mortal remains. Legal Medicine, 2019, 40, 32-36.	1.3	6
14	Evaluation of 12 Multi-InDel markers for forensic ancestry prediction in Asian populations. Forensic Science International: Genetics, 2019, 43, 102155.	3.1	17
15	Using miRNAs and circRNAs to estimate PMI in advanced stage. Legal Medicine, 2019, 38, 51-57.	1.3	34
16	Characterization of tissue-specific biomarkers with the expression of circRNAs in forensically relevant body fluids. International Journal of Legal Medicine, 2019, 133, 1321-1331.	2.2	38
17	Characterization of the extra copy of TPOX locus with tri-allelic pattern. BMC Genetics, 2019, 20, 18.	2.7	8
18	C-terminus of heat shock protein 60 can activate macrophages by lectin-like oxidized low-density lipoprotein receptor 1. Biochemical and Biophysical Research Communications, 2019, 508, 1113-1119.	2.1	9

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19	Evaluation of the inclusion of circular RNAs in mRNA profiling in forensic body fluid identification. International Journal of Legal Medicine, 2018, 132, 43-52.	2.2	35
20	Genetic analysis of 29 Y-STR loci in the Chinese Han population from Shanghai. Forensic Science International: Genetics, 2018, 32, e1-e4.	3.1	27
21	Null alleles and sequence variations at primer binding sites of STR loci within multiplex typing systems. Legal Medicine, 2018, 30, 10-13.	1.3	12
22	Metabolic profiling of femoral muscle from rats at different periods of time after death. PLoS ONE, 2018, 13, e0203920.	2.5	16
23	Applying massively parallel sequencing to paternity testing on the Ion Torrent Personal Genome Machine. Forensic Science International: Genetics, 2017, 31, 155-159.	3.1	18
24	Association of rs1122608 with Coronary Artery Disease and Lipid Profile: A Meta-analysis. Archives of Medical Research, 2016, 47, 315-320.	3.3	4
25	Mutation analysis of 19 autosomal short tandem repeats in Chinese Han population from Shanghai. International Journal of Legal Medicine, 2016, 130, 1439-1444.	2.2	15
26	Fluorescence- and magnetic-activated cell sorting strategies to separate spermatozoa involving plural contributors from biological mixtures for human identification. Scientific Reports, 2016, 6, 36515.	3.3	19
27	Deregulation of RGS17 Expression Promotes Breast Cancer Progression. Journal of Cancer, 2015, 6, 767-775.	2.5	18
28	Identification and characterization of the highly polymorphic locus D14S739 in the Han Chinese population. Croatian Medical Journal, 2015, 56, 482-489.	0.7	7
29	Genetic analysis of 17 Y-STR loci in Han population from Gansu province, northwestern China. Forensic Science International: Genetics, 2015, 19, 134-135.	3.1	7
30	Identification of mammalian species using the short and highly variable regions of mitochondrial DNA, 2015, 26, 550-554.	0.6	7
31	Deletion mapping of the regions with AMELY from two Chinese males. Legal Medicine, 2014, 16, 290-292.	1.3	6
32	Genetic distribution on 20 STR loci from the Han population in Shanghai, China. Forensic Science International: Genetics, 2014, 9, e30-e31.	3.1	36
33	Development of Highly Sensitive and Specific mRNA Multiplex System (XCYR1) for Forensic Human Body Fluids and Tissues Identification. PLoS ONE, 2014, 9, e100123.	2.5	43
34	The C-type lectin-like receptors of Dectin-1 cluster in natural killer gene complex. Glycoconjugate Journal, 2012, 29, 273-284.	2.7	13
35	Lectin-Like Oxidized Low-Density Lipoprotein Receptor-1 Delivers Heat Shock Protein 60-Fused Antigen into the MHC Class I Presentation Pathway. Journal of Immunology, 2010, 185, 2306-2313.	0.8	52
36	Laminarin-mediated targeting to Dectin-1 enhances antigen-specific immune responses. Biochemical and Biophysical Research Communications, 2010, 391, 958-962.	2.1	42

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37	Human Dectin-1 isoform E is a cytoplasmic protein and interacts with RanBPM. Biochemical and Biophysical Research Communications, 2006, 347, 1067-1073.	2.1	31