

Manohar R Furtado

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

1,842
citations

516710

16
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642732

23
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all docs

23
docs citations

23
times ranked

2270
citing authors

#	ARTICLE	IF	CITATIONS
1	Mutability of Y-Chromosomal Microsatellites: Rates, Characteristics, Molecular Bases, and Forensic Implications. <i>American Journal of Human Genetics</i> , 2010, 87, 341-353.	6.2	324
2	Progress toward an efficient panel of SNPs for ancestry inference. <i>Forensic Science International: Genetics</i> , 2014, 10, 23-32.	3.1	211
3	SNPs for a universal individual identification panel. <i>Human Genetics</i> , 2010, 127, 315-324.	3.8	194
4	Comparative genomics of the bacterial genus <i>Listeria</i> : Genome evolution is characterized by limited gene acquisition and limited gene loss. <i>BMC Genomics</i> , 2010, 11, 688.	2.8	174
5	Genome sequencing reveals diversification of virulence factor content and possible host adaptation in distinct subpopulations of <i>Salmonella enterica</i> . <i>BMC Genomics</i> , 2011, 12, 425.	2.8	133
6	Comparative Analysis of Genome Sequences Covering the Seven <i>Cronobacter</i> Species. <i>PLoS ONE</i> , 2012, 7, e49455.	2.5	130
7	Comprehensive mutation analysis of 17 Y-chromosomal short tandem repeat polymorphisms included in the AmpFISTR [®] Yfiler [®] PCR amplification kit. <i>International Journal of Legal Medicine</i> , 2009, 123, 471-482.	2.2	121
8	A Whole-Genome Single Nucleotide Polymorphism-Based Approach To Trace and Identify Outbreaks Linked to a Common <i>Salmonella enterica</i> subsp. <i>enterica</i> Serovar Montevideo Pulsed-Field Gel Electrophoresis Type. <i>Applied and Environmental Microbiology</i> , 2011, 77, 8648-8655.	3.1	100
9	Analysis of alternatively spliced human immunodeficiency virus type-1 mRNA species, one of which encodes a novel TAT-ENV fusion protein. <i>Virology</i> , 1991, 185, 258-270.	2.4	64
10	Developmental Validation of the Quantifiler [®] Duo DNA Quantification Kit for Simultaneous Quantification of Total Human and Human Male DNA and Detection of PCR Inhibitors in Biological Samples*. <i>Journal of Forensic Sciences</i> , 2009, 54, 305-319.	1.6	64
11	Developmental Validation of the PrepFiler [®] Forensic DNA Extraction Kit for Extraction of Genomic DNA from Biological Samples*. <i>Journal of Forensic Sciences</i> , 2009, 54, 599-607.	1.6	62
12	Identification and Characterization of Novel <i>Salmonella</i> Mobile Elements Involved in the Dissemination of Genes Linked to Virulence and Transmission. <i>PLoS ONE</i> , 2012, 7, e41247.	2.5	61
13	Mini-haplotypes as lineage informative SNPs and ancestry inference SNPs. <i>European Journal of Human Genetics</i> , 2012, 20, 1148-1154.	2.8	45
14	Expanding data and resources for forensic use of SNPs in individual identification. <i>Forensic Science International: Genetics</i> , 2012, 6, 646-652.	3.1	43
15	A Sample Extraction Method for Faster, More Sensitive PCR-Based Detection of Pathogens in Blood Culture. <i>Journal of Molecular Diagnostics</i> , 2012, 14, 120-129.	2.8	26
16	Genomic characterization of <i>Salmonella</i> Cerro ST367, an emerging <i>Salmonella</i> subtype in cattle in the United States. <i>BMC Genomics</i> , 2014, 15, 427.	2.8	19
17	Extraction of high quality DNA from biological materials and calcified tissues. <i>Forensic Science International: Genetics Supplement Series</i> , 2009, 2, 159-160.	0.3	15
18	Multiplexed SNP detection panels for human identification. <i>Forensic Science International: Genetics Supplement Series</i> , 2009, 2, 538-539.	0.3	13

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19	Evaluation of Applied Biosystems MicroSEQ [®] Real-Time PCR System for Detection of Salmonella spp. in Food. <i>Journal of AOAC INTERNATIONAL</i> , 2011, 94, 1106-1116.	1.5	11
20	Rapid Detection of Salmonella in Pet Food: Design and Evaluation of Integrated Methods Based on Real-Time PCR Detection. <i>Journal of Food Protection</i> , 2012, 75, 347-352.	1.7	10
21	Efficient expression of small RNA polymerase III genes from a novel Simian virus 40 vector and their effect on viral gene expression. <i>Nucleic Acids Research</i> , 1989, 17, 1159-1176.	14.5	5
22	A dedicated automated system for extraction, quantification and STR amplification of forensic evidence samples. <i>Forensic Science International: Genetics Supplement Series</i> , 2009, 2, 64-65.	0.3	1