David H Warshauer

List of Publications by Year in descending order

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840585 1058333 14 693 11 14 citations h-index g-index papers 15 15 15 622 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	High-quality and high-throughput massively parallel sequencing of the human mitochondrial genome using the Illumina MiSeq. Forensic Science International: Genetics, 2014, 12, 128-135.	1.6	155
2	Single nucleotide polymorphism typing with massively parallel sequencing for human identification. International Journal of Legal Medicine, 2013, 127, 1079-1086.	1.2	112
3	STRait Razor: A length-based forensic STR allele-calling tool for use with second generation sequencing data. Forensic Science International: Genetics, 2013, 7, 409-417.	1.6	94
4	High sensitivity multiplex short tandem repeat loci analyses with massively parallel sequencing. Forensic Science International: Genetics, 2015, 16, 38-47.	1.6	69
5	STRait Razor v2.0: The improved STR Allele Identification Tool – Razor. Forensic Science International: Genetics, 2015, 14, 182-186.	1.6	55
6	Increasing the discrimination power of ancestry- and identity-informative SNP loci within the ForenSeqâ,,¢ DNA Signature Prep Kit. Forensic Science International: Genetics, 2018, 36, 60-76.	1.6	41
7	An evaluation of the transfer of saliva-derived DNA. International Journal of Legal Medicine, 2012, 126, 851-861.	1.2	38
8	Massively parallel sequencing of 68 insertion/deletion markers identifies novel microhaplotypes for utility in human identity testing. Forensic Science International: Genetics, 2016, 25, 198-209.	1.6	29
9	Novel Y-chromosome Short Tandem Repeat Variants Detected Through the Use of Massively Parallel Sequencing. Genomics, Proteomics and Bioinformatics, 2015, 13, 250-257.	3.0	28
10	Sequencing the hypervariable regions of human mitochondrial DNA using massively parallel sequencing: Enhanced data acquisition for DNA samples encountered in forensic testing. Legal Medicine, 2015, 17, 123-127.	0.6	28
11	Massively parallel sequencing of forensically relevant single nucleotide polymorphisms using TruSeqâ,,¢ forensic amplicon. International Journal of Legal Medicine, 2015, 129, 31-36.	1.2	23
12	Validation of the PLEX-IDTM mass spectrometry mitochondrial DNA assay. International Journal of Legal Medicine, 2013, 127, 277-286.	1.2	8
13	Empirical testing of a 23-AIMs panel of SNPs for ancestry evaluations in four major US populations. International Journal of Legal Medicine, 2016, 130, 891-896.	1.2	8
14	Deep Sequencing Provides Comprehensive Multiplex Capabilities. Forensic Science International: Genetics Supplement Series, 2013, 4, e334-e335.	0.1	5