## Rebecca Just

## List of Publications by Year in descending order

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430754 395590 1,280 33 18 33 h-index citations g-index papers 33 33 33 1038 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Modeling allelic analyte signals for aSTRs in NGS DNA profiles. Journal of Forensic Sciences, 2021, 66, 1234-1245.	0.9	8
2	Validation of NGS for mitochondrial DNA casework at the FBI Laboratory. Forensic Science International: Genetics, 2020, 44, 102151.	1.6	48
3	Variability and additivity of read counts for aSTRs in NGS DNA profiles. Forensic Science International: Genetics, 2020, 48, 102351.	1.6	5
4	Sequence-based autosomal STR characterization in four US populations using PowerSeqâ,,¢ Auto/Y system. Forensic Science International: Genetics, 2020, 48, 102311.	1.6	11
5	An examination of STR nomenclatures, filters and models for MPS mixture interpretation. Forensic Science International: Genetics, 2020, 48, 102319.	1.6	10
6	Automation of high volume MPS mixture interpretation using CaseSolver. Forensic Science International: Genetics Supplement Series, 2019, 7, 14-15.	0.1	2
7	Use of the LUS in sequence allele designations to facilitate probabilistic genotyping of NGS-based STR typing results. Forensic Science International: Genetics, 2018, 34, 197-205.	1.6	29
8	A closer look at Verogen's Forenseqâ,,¢ DNA Signature Prep kit autosomal and Yâ€5TR data for streamlined analysis of routine reference samples. Electrophoresis, 2018, 39, 2685-2693.	1.3	34
9	Performance and concordance of the ForenSeqâ,,¢ system for autosomal and Y chromosome short tandem repeat sequencing of reference-type specimens. Forensic Science International: Genetics, 2017, 28, 1-9.	1.6	76
10	Internal validation of STRmixâ,,¢ for the interpretation of single source and mixed DNA profiles. Forensic Science International: Genetics, 2017, 29, 126-144.	1.6	74
11	The mitochondrial landscape of African Americans: An examination of more than 2500 control region haplotypes from 22 U.S. locations. Forensic Science International: Genetics, 2016, 22, 139-148.	1.6	10
12	Mitochondrial DNA heteroplasmy in the emerging field of massively parallel sequencing. Forensic Science International: Genetics, 2015, 18, 131-139.	1.6	118
13	Full mtGenome reference data: Development and characterization of 588 forensic-quality haplotypes representing three U.S. populations. Forensic Science International: Genetics, 2015, 14, 141-155.	1.6	78
14	Short tandem repeat typing on the 454 platform: Strategies and considerations for targeted sequencing of common forensic markers. Forensic Science International: Genetics, 2014, 12, 107-119.	1.6	68
15	Questioning the prevalence and reliability of human mitochondrial DNA heteroplasmy from massively parallel sequencing data. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E4546-7.	3.3	25
16	Development of forensic-quality full mtGenome haplotypes: Success rates with low template specimens. Forensic Science International: Genetics, 2014, 10, 73-79.	1.6	18
17	Characterization of a modified amplification approach for improved STR recovery from severely degraded skeletal elements. Forensic Science International: Genetics, 2012, 6, 578-587.	1.6	18
18	Assessing the potential of next generation sequencing technologies for missing persons identification efforts. Forensic Science International: Genetics Supplement Series, 2011, 3, e447-e448.	0.1	19

#	Article	IF	CITATIONS
19	Short tandem repeat sequencing on the 454 platform. Forensic Science International: Genetics Supplement Series, 2011, 3, e357-e358.	0.1	18
20	Evaluation of automatable silica-based extraction methods for low quantity samples. Forensic Science International: Genetics Supplement Series, 2011, 3, e504-e505.	0.1	4
21	Titanic's unknown child: The critical role of the mitochondrial DNA coding region in a re-identification effort. Forensic Science International: Genetics, 2011, 5, 231-235.	1.6	17
22	mtGenome reference population databases and the future of forensic mtDNA analysis. Forensic Science International: Genetics, 2011, 5, 222-225.	1.6	33
23	The Use of Mitochondrial DNA Single Nucleotide Polymorphisms to Assist in the Resolution of Three Challenging Forensic Cases. Journal of Forensic Sciences, 2009, 54, 887-891.	0.9	32
24	Mitochondrial control region sequences from an African American population sample. Forensic Science International: Genetics, 2009, 4, e45-e52.	1.6	15
25	Identification of West Eurasian mitochondrial haplogroups by mtDNA SNP screening: Results of the 2006–2007 EDNAP collaborative exercise. Forensic Science International: Genetics, 2008, 2, 61-68.	1.6	13
26	Mitochondrial control region sequences from a U.S. "Hispanic―population sample. Forensic Science International: Genetics, 2008, 2, e19-e23.	1.6	15
27	Complete mitochondrial genome sequences for 265 African American and U.S. "Hispanic―individuals. Forensic Science International: Genetics, 2008, 2, e45-e48.	1.6	38
28	The application of mtDNA SNPs to a forensic case. Forensic Science International: Genetics Supplement Series, 2008, 1, 295-297.	0.1	4
29	Development and expansion of high-quality control region databases to improve forensic mtDNA evidence interpretation. Forensic Science International: Genetics, 2007, 1, 154-157.	1.6	49
30	DNA Identification of "Earthquake McGoon―50â€fYears Postmortem. Journal of Forensic Sciences, 2007, 52, 1115-1118.	0.9	37
31	Effective strategies for forensic analysis in the mitochondrial DNA coding region. International Journal of Legal Medicine, 2006, 120, 27-32.	1.2	34
32	Single nucleotide polymorphisms over the entire mtDNA genome that increase the power of forensic testing in Caucasians. International Journal of Legal Medicine, 2004, 118, 137-146.	1.2	195
33	A multiplex allele-specific primer extension assay for forensically informative SNPs distributed throughout the mitochondrial genome. International Journal of Legal Medicine, 2004, 118, 147-157.	1.2	125