

Diversity in a Variable-Number Tandem Repeat from <i>

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Francisella tularensis Strain Typing Using Multiple-Locus, Variable-Number Tandem Repeat Analysis. Journal of Clinical Microbiology, 2001, 39, 3186-3192.	1.8	150
2	A tandem repeats database for bacterial genomes: application to the genotyping of Yersinia pestis and Bacillus anthracis. BMC Microbiology, 2001, 1, 2.	1.3	222
3	Differentiation of Strains of Xylella fastidiosa by a Variable Number of Tandem Repeat Analysis. Applied and Environmental Microbiology, 2001, 67, 4091-4095.	1.4	97
4	Extensive Allelic Variation among Francisella tularensis Strains in a Short-Sequence Tandem Repeat Region. Journal of Clinical Microbiology, 2001, 39, 3140-3146.	1.8	66
5	Identification and Characterization of Variable-Number Tandem Repeats in the Yersinia pestis Genome. Journal of Clinical Microbiology, 2001, 39, 3179-3185.	1.8	198
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9	Variable Tandem Repeat Vca of Vibrio cholerae. Molecular Biology, 2002, 36, 874-878.	0.4	3
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13	Tandem Tetramer-Based Microsatellite Fingerprinting for Typing of Proteus mirabilis Strains. Journal of Clinical Microbiology, 2003, 41, 1673-1680.	1.8	5
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16	Genotyping, Orientalis-like <i>Yersinia pestis</i>, and Plague Pandemics. Emerging Infectious Diseases, 2004, 10, 1585-1592.	2.0	166
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