ERRATUM

Use of Applied Biosystems 7900HT Sequence Detection System and Taqman Assay for Detection of Quinolone-Resistant *Neisseria gonorrhoeae*

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Volume 42, no. 7, p. 3281–3283, 2004. Page 3281, column 1, line 9 from bottom should read as follows. “. . . 1 μ l each of forward and reverse primers (0.2 μ M final concentration of each). . . .”

Page 3281. The contact information in the corresponding author footnote should read as follows. “Division of Infectious Diseases, Johns Hopkins University School of Medicine, 720 Rutland Ave., 1165 Ross, Baltimore, MD 21205. Phone: (410) 614-4480. Fax: (410) 614-9775. E-mail: jgiles4@jhmi.edu.”

Page 3281. Table 1 should appear as shown below.

TABLE 1. QRDR mutation patterns for gonococcal samples as determined by direct sequencing and real-time PCR detection with the ABI 7900HT sequence detection system

Sample type	QRDR mutation pattern ^a					
	<i>gyrA</i>		<i>parC</i>			
	aa 91 (Ser) ^b	aa 95 (Asp)	aa 86 (Asp)	aa 87 (Ser)	aa 88 (Ser)	aa 91 (Glu)
A (WT, Cip ⁵)	WT	WT	WT	WT	WT	WT
B (mutant, Cip ⁱ)	WT	GAC→AAC (Asp→Asn)	WT	WT	WT	WT
C (mutant, Cip ⁱ)	TCC→TTC (Ser→Phe)	GAC→GGC (Asp→Gly)	GAC→AAC (Asp→Asn)	WT	WT	WT

^a WT denotes the WT sequence at targeted loci.

^b The WT amino acid is listed in parentheses.

Page 3282, column 2: Lines 11–13 should read as follows. “. . . With widespread use of nucleic acid amplification tests (NAATs), antimicrobial resistance detection will require molecular methods, as has been described for other organisms (1, 7, 14, 16–18, 25).”

Page 3283: The Acknowledgments section was inadvertently omitted and should appear as shown below.

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Page 3283: Reference 1 should read as shown below.

1. Ahmad, S., E. E. Udo, and T. D. Chugh. 2000. Molecular diagnosis of antimicrobial resistance. *Leban. Med. J.* **48**:203–207.