

ERRATA

A Simple RNA Probe System for Analysis of *Listeria monocytogenes* Polymerase Chain Reaction Products

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Volume 59, no. 9, p. 2796, column 2, lines 4 and 5: "For the amplification of a 730-bp fragment spanning nucleotides 602 to 1332" should read "For amplification of a 730-bp fragment spanning nucleotides 680 to 1411."

Page 2796, column 2, lines 36 and 37: "The template was a 321-bp fragment spanning nucleotides 861 to 1156" should read "The template was a 321-bp fragment spanning nucleotides 939 to 1234."

Mobile Bacteria and Transport of Polynuclear Aromatic Hydrocarbons in Porous Media

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Volume 59, no. 10, p. 3309, equation 4. The equation should read as follows:

$$R = \frac{(1 + K_d^{\text{cell}}[\text{cell}]) + (K_d^s \rho/\eta)}{1 + \{K_d^{\text{cell}}[\text{cell}]/(1 + K_{\text{cell}}^s \rho/\eta)\}} \quad (4)$$

Quantitative Reverse Sample Genome Probing of Microbial Communities and Its Application to Oil Field Production Waters

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Volume 59, no. 12, p. 4101, first sentence of the Abstract. This sentence should read as follows: "This paper presents a protocol for quantitative analysis of microbial communities by reverse sample genome probing in which (i) whole community DNA is isolated and labeled in the presence of a known amount of an added internal standard and (ii) the resulting spiked reverse genome probe is hybridized with a master filter on which denatured genomic DNAs from bacterial standards isolated from the target environment were spotted in large amounts (up to 1,500 ng) in order to improve detection sensitivity."