

## AUTHOR'S CORRECTION

# Multi-Virulence-Locus Sequence Typing of *Listeria monocytogenes*

**Wei Zhang, Bhushan M. Jayarao, Stephen J. Knabel**

Department of Food Science and Department of Veterinary Science, The Pennsylvania State University, University Park, Pennsylvania 16802

Volume 70, no. 2, p. 913–920, 2004. Page 914, Table 1: For strains FSL J1-119, FSL J1-116, and FSL N3-013, the “Allelic profile” entry for *clpP* should be “4” and the “ST” entry should be “20.” For strain FSL R2-499, the “Allelic profile” entry for “*inlC*” should be “1” and the “ST” entry should be “1.”

Errors in the original allelic and sequence type assignment were due to sequencing errors on three synonymous single-nucleotide-polymorphic sites in *inlC* and *clpP* genes of the four strains in the original study. We have resequenced and confirmed all genes and strains and deposited updated sequences in GenBank under accession numbers [AY259744.1](#) through [AY259771.1](#) (*inlC*) and [AY259772.1](#) through [AY259799.1](#) (*clpP*).