AUTHOR'S CORRECTION

Evolution of the Core Genome of *Pseudomonas syringae*, a Highly Clonal, Endemic Plant Pathogen

Sara F. Sarkar and David S. Guttman

Department of Botany, University of Toronto, Toronto, Ontario, Canada

Volume 70, no. 4, p. 1999–2012, 2004. It has come to our attention that the PtoDC3000 and PsyB728A strain names became switched for some of the analyses in this paper. This error specifically affects the placement of the two strain names in the *rpoD* split decomposition graph in Fig. 3 and the MLST analysis of recombination for the *rpoD* locus presented on p. 2006.

Page 2005, Fig. 3, bottom panel: The strain names PtoDC3000 and PsyB728A should be switched.

Pages 2006–2007: The entire "MLST analyses" section should be deleted.

Page 2011, column 1, lines 19–26: "The MLST-based recombination analyses are . . . for a data set of the present size" should be deleted.

As discussed in the original paper, these analyses were largely inconclusive. The error did not carry over into any other analyses and in no way changes the conclusions of the work.