ERRATUM

Quantification of *Hyphomicrobium* Populations in Activated Sludge from an Industrial Wastewater Treatment System as Determined by 16S rRNA Analysis

A. C. LAYTON, P. N. KARANTH, C. A. LAJOIE, A. J. MEYERS, I. R. GREGORY, R. D. STAPLETON, D. E. TAYLOR, AND G. S. SAYLER

Center for Environmental Biotechnology, Department of Chemical Engineering, and Department of Microbiology and Department of Ecology and Evolutionary Biology, The University of Tennessee, Knoxville, Tennessee 37996, and Tennessee Eastman Division, Eastman Chemical Company, Kingsport, Tennessee 37662

Volume 66, no. 3, p. 1167–1174, 2000. P. 1169, Table 2, column 3: the oligonucleotide probe sequence for universal probe 1390 should read “GACGGGCAGGTTGACAA.”