

ADDENDUM

Adeleke A, Pruckler J, Benson R, Rowbotham T, Halablab M, Fields B. *Legionella*-like amoebal pathogens—phylogenetic status and possible role in respiratory disease. *Emerging Infectious Diseases* 1996;3:225-30.

Since the publication of the above article, accession numbers have been allocated to the following sequences:

LLAP -1 U64034

LLAP -8 U64035

Information about the phylogeny of *S. lyticum* and LLAP-3 can be found in references 20 and 21, respectively. The derivation of sequence data (accession numbers: LLAP-4x97357, LLAP-6x97359, LLAP-10x97363, LLAP-11x97362, LLAP-12x97366J) referred to in our publication, can be found in Birtles RJ, Rowbotham TJ, Raoult D, Harrison TG. Phylogenetic diversity of intra-amoebal legionellae as revealed by 16S rRNA gene sequencing. *Microbiology* 1996; 142:3525-30. In this study, which was carried out concurrently with the study described in the article by Adeleke et al. 16S rRNA sequence data were also obtained for LLAP1 (accession number x97355), LLAP2 (x97356), LLAP7 (x97365), LLAP8 (x97361), LLAP9 (x97360), and the *L. lytica* strains L2 (x97364) and LLAP3 (x97358).