

ERRATUM

Multilocus Sequence Typing and Evolutionary Relationships among the Causative Agents of Melioidosis and Glanders, *Burkholderia pseudomallei* and *Burkholderia mallei*

Daniel Godoy, Gaynor Randle, Andrew J. Simpson, David M. Aanensen, Tyrone L. Pitt, Reimi Kinoshita, and Brian G. Spratt

Department of Infectious Disease Epidemiology, Faculty of Medicine, Imperial College London, St. Mary's Hospital, London W2 1PG, Molecular Infectious Diseases Group, Weatherall Institute of Molecular Medicine, John Radcliffe Hospital, Oxford OX3 9DS, Department of Medical Microbiology, The Royal Free Hospital, London NW3 2QG, and Laboratory of Hospital Infection, Central Public Health Laboratory, London NW9 5HT, United Kingdom, and Veterinary Hospital, Ocean Park Corporation, and Departments of Pathology and Microbiology, Faculty of Medicine, University of Hong Kong, Hong Kong

Vol. 41, no. 5, p. 2068–2079, 2003. Page 2069, column 2, lines 6–16 should read as follows: “. . .The following pairs of primers were used for the amplification of the housekeeping gene fragments: ace-up (5'-CGGCGCTTCTCAAACGATA-3') and ace-dn (5'-GAATCGCCTTCACCATGTC-3'), gltB-up (5'ACGCTCGCGATCGCGATGAA-3') and gltB-dn (5'-TTCAGCACGAGCGTCTGCTG-3'), gmhD-up (5'-GCAGTTCCTGTATGCGTC-3') and gmhD-dn (5'-GAAGCACTGGTACTTGCC-3'), lepA-up (5'-CATATTCGCAATTTCTCGATC-3') and lepA-dn (5'-CACGAGCATCACGACGCCG-3'), lipA-up (5'-GGCACCGCGACGTTTCATG-3') and lipA-dn (5'-GACCATCAGGCCCGATTTTCG-3'), narK-up (5'-CTACTCGTGCGCTGGGAT-3') and narK-dn (5'-GACGATGAACGGCACCCAC-3'), and ndh-up (5'-AGTCGCGACGTTCTACAC-3') and ndh-dn (5'-CGAGTTCAGACGAGATA-3'). . .” The correct primer sequences are available at the *Burkholderia pseudomallei* MLST website (<http://bpseudomallei.mlst.net>).

Page 2071, column 2, line 1: “in 1942 from a horse in China” should read “in 1944 from a human in Burma.”