**Mycobacterium avium subsp. paratuberculosis**

Infection in a Patient with HIV, Germany

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*Mycobacterium avium* subsp. *paratuberculosis* (MAP), the causative agent of Johne disease in ruminants, has been incriminated as the cause of Crohn disease in humans. We report the first case of human infection with MAP in a patient with HIV; infection was confirmed by obtaining isolates from several different specimen types.

Opportunistic infections caused by various *Mycobacterium* species are among the leading AIDS indicator diseases in HIV-positive patients (1). Infections with nontuberculous mycobacteria occur mainly in patients who have low CD4+ counts (<50 cells) or high virus counts (2); *Mycobacterium avium* complex is the most important mycobacterial species. *M. avium* complex includes the species *M. avium* and *M. intracellularare*, with *M. avium* consisting of *M. avium* subsp. *avium*, *M. avium* subsp. *sylvaticum*, and *M. avium* subsp. *paratuberculosis* (MAP). All these subspecies have identical 16S rRNA gene and 16S to 23S transcribed spacer sequences, as well as shared biochemical characteristics (3). However, MAP is dependent on mycobactin for its growth, whereas *M. avium* grows well on different solid media.

MAP is the causative agent of Johne disease, a chronic granulomatous ileitis occurring mainly in ruminants (4). MAP has been incriminated as the cause of Crohn disease in humans (5,6), although conflicting findings have been reported. However, culture-confirmed cases of MAP in human specimens remain rare (5,6).

**Case Report**

A 36-year-old HIV-positive man, who had been treated at our hospital since 1995 for HIV, hepatitis C, and hemophilia, had profuse diarrhea (6–8 episodes/day), fever as high as 39.9°C, and 10 kg of body weight loss in 5 weeks. Laboratory findings included hemoglobin 9.6 g/dL, pseudocholinesterase 2,099 U/L, HIV-DNA virus count 500 copies/mL, CD4+ lymphocyte count 29 x 10⁶/mL, and C-reactive protein 76 mg/L. Stained colon tissue samples, bone marrow punch, and liver biopsy showed abundant acid-fast bacilli. Endoscopic findings on colonoscopy were multiple polypoid lesions approximately 5 mm in size in the transverse and sigmoid colon.

Microbiologic analyses included culture for mycobacteria (liquid media: BACTEC 460TB or MGIT [Becton, Dickinson and Company, Cockeysville, MD] and solid media produced in-house, all media without supplementation of mycobactin) from at least 21 specimens (blood, urine, sputum, biopsy, feces) over a 3-year period. Of these, eight specimens (blood, feces, and biopsy) were positive for mycobacteria in liquid media after 6 to 16 weeks of incubation. Subcultures remained negative on Löwenstein-Jensen slants but after approximately 4 weeks became positive on mycobactin-supplemented Middlebrook slants with colorless dysgonic colonies. Microscopic examination of these colonies showed acid-fast bacilli (Figure 1).

For species identification, AccuProbe assays (Gen-Probe, San Diego, CA) for *M. avium* complex were performed on liquid media, all yielding strong positive results. However, repeated attempts to perform drug-susceptibility testing in the liquid BACTEC 460TB system were unsuccessful because of insufficient growth of the control. Since *M. avium* complex usually grows very well, the primary identification was questionable. Thus, polymerase chain reaction (PCR) for the amplification of a part of the mycobacterial gene coding for the ribosomal 16S RNA and additional sequencing was performed from two positive cultures (7). The resulting sequence was compared with those stored in the International Reference Center for Mycobacteria, Borstel, Germany; and University of Münster, Münster, Germany.

![Figure 1. Ziehl-Neelsen–stained micrograph of *Mycobacterium avium* subsp. *paratuberculosis* colonies growing on mycobactin-supplemented Middlebrook agar.](image-url)
Nucleotide Sequence Database (8), showing the signature sequence of *M. avium/M. paratuberculosis*, which is identical for both species and confirmed the AccuProbe results. For further differentiation between *M. avium* and MAP, PCR targeting the insertion sequence IS900 (primer: IS900-1: 5’-TGTTCGGGGCCGTCGCTTAG; IS900-2: 5’-CGTCCAGGCGCCAAGAGTAT), which is present only in MAP strains (9), was done with the two most recent positive cultures. This assay showed clearly positive results from the two cultures tested and the MAP type strain, while the *M. avium* strains remained negative (Figure 2).

Because acid-fast bacilli were identified in biopsy specimens, treatment was started with ethambutol, ciprofloxacin, clarithromycin, and rifabutin. Initially, no clinical improvement was observed, and the patient’s weight loss and daily fever of 39°C–40°C continued. When ciprofloxacin was replaced with levofloxacin, progression of the infection was not clearly positive and may have been hampered by the lack of clarithromycin and rifabutin, which were included in therapy. However, the patient’s response to treatment was not clearly positive and may have been hampered by his general poor health. This report suggests a pathogenic role of MAP for immunocompromised patients, raising the question of whether this strain so far has not been detected because of its limited growth, whether it has been misidentified as *M. avium*, or whether its occurrence in infections is low. However, herd prevalence of bovine paratuberculosis has been reported to range from 7% to 55% in Europe and to reach approximately 40% in stocks of >300 animals in the United States (4). Thus, consumption of inadequately pasteurized dairy products may be a possible risk for infection, especially for immunocompromised patients.

Conclusions

We describe the case of an HIV-infected patient who had a severe mycobacterial disorder thought to be caused by *M. avium* complex. Because growth was insufficient for susceptibility testing, the presence of MAP was assumed; however, the assumption was made after 2 years, because of difficulties in isolating MAP from human specimens (e.g., blood) in media not thought to enable its growth. Finally, the demonstration of the insertion sequence IS900, an assay not routinely performed in human diagnostic laboratories like ours, confirmed this hypothesis.

Acknowledgments

We thank Marie Thorel for providing the MAP type strain used as positive control for IS900 PCR and Frauke Schaefer for excellent technical assistance.

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References


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