**INTRODUCTION**

*Mycobacterium chimaera (M. chimaera)* is a slow-growing nontuberculous mycobacteria that has recently become a global public health concern due to an invasive outbreak following cardiac surgery. This outbreak has been attributed to a specific brand of contaminated heater-cooler units (HCUs) used during surgery. Highly related genomes of *M. chimaera* have been isolated from geographically distant infections and from the manufacturer and are consistent with contamination from a single point source. Given the rarity of postoperative *M. chimaera* infection, its typically indolent course, and lack of physician awareness of the outbreak, it is often recognized very late or misdiagnosed.

**CASE REPORT**

A 53-year-old woman with a history of bicuspid aortic valve and ascending aortic aneurysm presented with constitutional symptoms 2 years after undergoing bovine pericardial aortic valve and aortic root replacement. An infectious source was ruled out on multiple occasions with negative blood cultures, imaging studies (computed tomography scan of chest, abdomen, and pelvis), and transesophageal echocardiogram. The patient had persistent cytopenias, marked splenomegaly, transaminitis, and constitutional symptoms including weight loss and fevers that led to expanding our differential diagnosis. She was noted to have non–parathyroid-hormone-related hypercalcemia and an elevated serum angiotensin converting enzyme level. A bone marrow biopsy revealed noncaseating granulomas (Figure 1). At this point, a decision was made to treat her for presumed sarcoidosis, and she was started on immunosuppressants. Although constitutional symptoms initially improved, she developed fever and back pain 6 months later, leading to further workup. Spine magnetic resonance imaging revealed T10-T11 vertebral body inflammation concerning for infection, and a subsequent bone biopsy showed acid-fast bacilli. Culture of the specimen and blood cultures grew *M. chimaera*, and the patient was started on a macrolide-based, multidrug, antimicrobial regimen.

**DISCUSSION**

After the 2011 diagnosis of disseminated *M. chimaera* infection in a Swiss patient post cardiothoracic surgery, several similar cases have been recognized—all linked to contaminated HCUs of a specific brand. The infections are presumably spread during cardiac surgery by airborne transmission of aerosolized bacteria from the water tanks of these units, and it can take months to
years to become clinically apparent. Symptoms are nonspecific and typically present as constitutional symptoms.\textsuperscript{2} \textit{M. chimaera} is less virulent than other nontuberculous mycobacteria and is usually associated with pulmonary infections in immunosuppressed patients and patients with pre-existing lung conditions. Disseminated extrapulmonary infection—including prosthetic valve endocarditis, prosthetic vascular graft infection, sternotomy wound infection, and mediastinitis—has been seen in patients whose surgery involved a contaminated HCU. Embolic and immunologic sequelae of disseminated infection have also been reported, such as splenomegaly, arthritis, osteomyelitis, cytopenia due to bone marrow involvement, chorioretinitis, hepatitis, nephritis, and myocarditis.\textsuperscript{2}

It is noteworthy that nontubercular mycobacterial infections can cause both caseating and noncaseating granuloma. Since the clinical features and indolent course of \textit{M. chimaera} infection can mimic sarcoidosis,\textsuperscript{3} it should be suspected in patients who have undergone cardiac surgery and present with unexplained fever and evidence of granulomatous disease. The presence of extrapulmonary disease and bone-marrow involvement, infrequent with sarcoidosis,\textsuperscript{4} may indicate the alternate diagnosis of \textit{M. chimaera}—an important distinction given the potential consequences of inappropriate immunosuppressant therapy in an infected patient.

The diagnosis of \textit{M. chimaera} infection is challenging because mycobacterial cultures are not part of the routine microbiological workup for cardiovascular infections. Identification of \textit{M. chimaera} requires special cultures, and results are slow and have low sensitivity except when infected tissue is obtained by invasive sampling. Differentiation of mycobacteria at the species level requires specialized DNA sequence-based testing, and no direct nucleic-acid amplification or metagenomics assay is available for rapid detection of \textit{M. chimaera}. Whole-genome sequencing has shown some utility in identifying iatrogenic \textit{M. chimaera} infection, and further efforts are underway.\textsuperscript{5} A clarithromycin-based, multidrug, antimicrobial regimen is commonly used for treatment. Even so, eradication of \textit{M. chimaera} is difficult, and outcomes have been poor despite long-term antimycobacterial and surgical therapies.\textsuperscript{2,3}

Healthcare-associated infection due to \textit{M. chimaera} is likely still under-recognized, and a high level of suspicion in the at-risk population should be maintained to facilitate more rapid diagnosis and prevent inappropriate treatment of this potentially devastating condition.

\textbf{Conflict of Interest Disclosure:}  
The authors have completed and submitted the \textit{Methodist DeBakey Cardiovascular Journal} Conflict of Interest Statement and none were reported.

\textbf{Keywords:}  
mycobacterium chimaera, sarcoidosis, cardiac surgery, heater-cooler units

\textbf{REFERENCES}


