

Mycobacterium tuberculosis and Mycobacterium avium Coinfection in an Immunocompetent Patient

Review began 06/15/2024
Review ended 06/21/2024
Published 06/25/2024

© Copyright 2024

Karki et al. This is an open access article distributed under the terms of the Creative Commons Attribution License CC-BY 4.0., which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Sailesh Karki ¹, Sagar Pandey ¹, Nabin K C ¹, Arjun Mainali ¹, Muhammad N. Pasha ², Harish J. Patel ²

¹. Internal Medicine, One Brooklyn Health/Interfaith Medical Center, Brooklyn, USA ². Pulmonary and Critical Care Medicine, One Brooklyn Health, Brooklyn, USA

Corresponding author: Sailesh Karki, sailkarki@gmail.com

Abstract

Despite the increasing incidence of simultaneous mycobacterial and non-mycobacterial tuberculosis (TB) infection, little literature is available exploring the topic. Here, we present a case of a 22-year-old female diagnosed with pulmonary TB for four months with simultaneous multiple sputum cultures positive for non-tuberculous mycobacteria (NTM). Computed tomography of the chest without contrast reported linear areas of scarring involving both lung apices, more prominent on the left side. The patient completed intensive phase treatment for TB and is currently on isoniazid and rifampin with a referral to an infectious disease specialist for recommendations on treatment of *Mycobacterium avium* regimen in view of azithromycin allergy (intense cough and rash). While the coexistence of NTM is commonly attributed to colonization, differentiating colonization from disease is crucial considering the long duration of treatment, potential drug toxicity, risk of drug resistance, and significant cost of treatment. Clinical, microbiological, and radiological evidence should be considered for diagnosis of TB and NTM coinfection and expert consultation should be sought in formulating the treatment plan.

Categories: Internal Medicine, Infectious Disease, Pulmonology

Keywords: mycobacterium tuberculosis, mycobacterium avium complex, nontuberculous mycobacteria (ntm), azithromycin allergy, coinfection, tuberculosis

Introduction

Mycobacterium avium complex (MAC) is comprised of multiple non-tuberculous mycobacteria (NTM) that cause pulmonary disease, skin and soft tissue infections, musculoskeletal infections, disseminated disease, catheter-associated disease, and lymphadenitis [1]. Two major clinical manifestations of *Mycobacterium avium* pulmonary disease have been identified. The fibrocavitary subtype is rapidly progressive and usually develops in middle-aged male smokers [2]. In contrast, the nodular bronchiectatic subtype classically develops in non-smoker females and is also known as Lady Windermere syndrome [1,3]. Disseminated, multi-organ involvement occurs in immunocompromised patients, including patients taking immunosuppressive medications and people with acquired immunodeficiency syndrome (AIDS) [4]. The incidence of NTM has been increasing in developed countries, with a simultaneous decrease in *Mycobacterium tuberculosis* (MTB) infection. This has been partially attributed to improved diagnostic methods for NTM [5]. The coinfection of MTB and *Mycobacterium avium* in an immunocompetent patient in a developed country is rare and presents a diagnostic and clinical challenge. This case report discusses a unique presentation of such a coinfection in a young immunocompetent female.

Case Presentation

A 22-year-old female diagnosed with pulmonary tuberculosis for four months was referred to the pulmonary clinic by the patient's primary care physician due to multiple sputum cultures positive for MAC. The patient reported that the symptoms started as night sweats and sore throat. The symptoms had been started two months before the initial diagnosis of pulmonary tuberculosis. However, she denied any history of fever, cough, hemoptysis, weight loss, chest tightness, or positive contact history. She denied smoking, consuming alcohol, or using any illicit drug. Past medical history was significant for a diagnosis of latent tuberculosis at 15 years of age, which was treated with eight months of isoniazid and vitamin B6. The patient had no significant travel history. On examination, the patient's vital signs were stable. Bilateral bronchial breath sounds were heard on chest auscultation. The rest of the examination findings were within normal limits. Chest X-ray (CXR) showed bilateral apical lobe scarring/atelectasis (Figure 1). Computed tomography of the chest without contrast corroborated the CXR findings and reported linear areas of scarring involving both lung apices, more prominent on the left side (Figure 2).

How to cite this article

Karki S, Pandey S, K C N, et al. (June 25, 2024) Mycobacterium tuberculosis and Mycobacterium avium Coinfection in an Immunocompetent Patient. Cureus 16(6): e63108. DOI 10.7759/cureus.63108

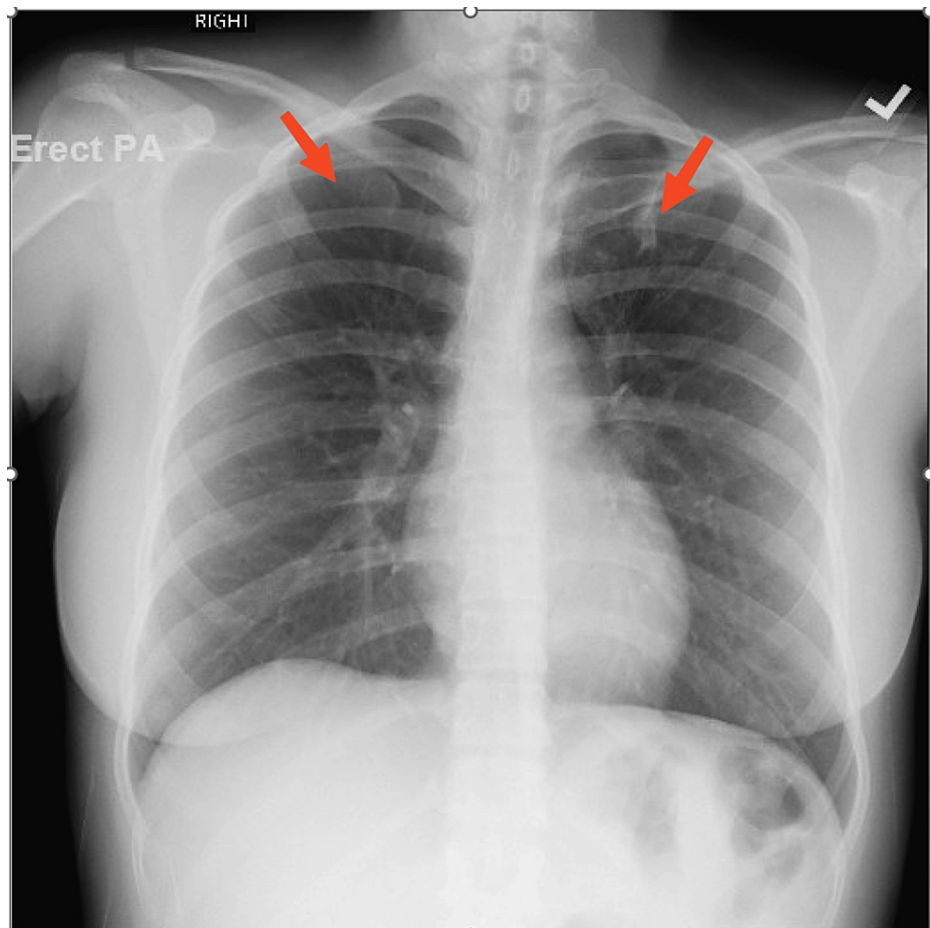


FIGURE 1: Chest X-ray demonstrating apical lobe scarring/atelectasis (red arrows).



FIGURE 2: CT of the chest without contrast demonstrating linear areas of scarring involving both lung apices and more prominent on the left side (red arrow).

The patient was diagnosed with pulmonary tuberculosis when she was admitted to the hospital for upper respiratory tract symptoms, which were not resolved on antibiotics. Workup was done for pulmonary tuberculosis, including sputum cultures and bronchoscopy sputum specimens, which tested positive for tuberculosis. The patient was, however, tested negative for HIV. She was started on intensive phase treatment for pulmonary tuberculosis, i.e., isoniazid, rifampin, pyrazinamide, and ethambutol, along with pyridoxine. Ethambutol and pyrazinamide had to be dose adjusted because of adverse effects to the medications. The patient was in the continuation phase of treatment with isoniazid and rifampin when she presented to the pulmonary clinic. On presentation to our clinic, she had decreased night sweats but persistent sore throat. Multiple sets of sputum and bronchial fluid specimens from bronchoscopy tested positive for mycobacterial tuberculosis and nontubercular mycobacteria (MAC). Lastly, the patient reported an allergy to azithromycin in the form of intense cough and rashes. Therefore, the patient was referred to an infectious disease specialist for expert opinion on the treatment regimen for the patient for coinfection of MTB with MAC in the setting of macrolide allergy.

Discussion

MAC organisms are ubiquitous in many environmental sites and are present in water, soil, and animals. The organism is thought to be acquired by inhalation or ingestion. There is no evidence for human-to-human transmission [6]. Differentiating colonization from disease is crucial in MAC considering the long duration of anti-microbial treatment, potential drug toxicity, and significant cost of treatment [7]. Detection of MAC in more than one sputum sample is recommended to rule out the possibility of contamination. Moreover, clinical and radiographic presentation has to be considered by the clinician, in addition to microbiological identification of MAC to avoid misinterpreting colonization as a true infection [1,7]. Our patient had an atypical presentation with a sore throat and night sweats. Additionally, radiography demonstrating apical scarring and tree bud opacities and multiple positive MAC results were supportive of MAC clinical disease.

Fujita et al. demonstrated concurrent infections with MAC involving microorganisms like methicillin-sensitive *Staphylococcus aureus* (MSSA), *Pseudomonas aeruginosa*, and *Aspergillus*. There was an increased prevalence of concurrent infection in those with chronic obstructive pulmonary disease (COPD) [8]. In a subsequent study, the coinfections were shown not to have any impact on the therapeutic efficacy of MAC [9]. Another study looked into the impact of concomitant infection by fungal species like *Aspergillus fumigatus*, *Histoplasma capsulatum*, and *Cryptococcus neoformans*. Patients with fungal co-infections were found to have a poorer prognosis [10]. Further studies are required to determine the impact of MTB coinfection on the prognosis of patients with MAC.

Coinfection with *Mycobacterium avium* may have important implications in selecting anti-microbial therapy. Bazzi et al. described a case where coinfection with MAC resulted in an incorrect interpretation of MTB as rifampin-resistant. This occurred because the test (GeneXpert) simultaneously identified the

rifampicin resistance gene of *M. avium* and detected the tuberculosis genes of MTB [11]. This points to the need for a multiplex polymerase chain reaction (PCR)-based mycobacterial detection system in areas with a high prevalence of MAC [12]. Treatment of MAC also has implications for lung function. In a study evaluating the coinfection of MAC with MTB, anti-MAC therapy was found to improve pulmonary function tests in patients who had abnormal pulmonary function at baseline [13].

Conclusions

Mycobacterium tuberculosis and *Mycobacterium avium* coinfection is increasing in incidence. Differentiating colonization from the disease of NTM is critical for the optimal treatment of patients, decreasing side effects, and reducing the risk of drug resistance.

Additional Information

Author Contributions

All authors have reviewed the final version to be published and agreed to be accountable for all aspects of the work.

Concept and design: Sailesh Karki, Sagar Pandey, Nabin K C, Arjun Mainali, Muhammad N. Pasha, Harish J. Patel

Drafting of the manuscript: Sailesh Karki, Sagar Pandey

Critical review of the manuscript for important intellectual content: Sailesh Karki, Sagar Pandey, Nabin K C, Arjun Mainali, Muhammad N. Pasha, Harish J. Patel

Supervision: Nabin K C, Arjun Mainali, Harish J. Patel

Disclosures

Human subjects: Consent was obtained or waived by all participants in this study. **Conflicts of interest:** In compliance with the ICMJE uniform disclosure form, all authors declare the following: **Payment/services info:** All authors have declared that no financial support was received from any organization for the submitted work. **Financial relationships:** All authors have declared that they have no financial relationships at present or within the previous three years with any organizations that might have an interest in the submitted work. **Other relationships:** All authors have declared that there are no other relationships or activities that could appear to have influenced the submitted work.

References

1. Akram SM, Attia FN: *Mycobacterium avium* Complex. StatPearls Publishing, Treasure Island, FL; 2024.
2. Griffith DE, Aksamit T, Brown-Elliott BA, et al.: An official ATS/IDSA statement: diagnosis, treatment, and prevention of nontuberculous mycobacterial diseases. *Am J Respir Crit Care Med*. 2007, 175:367-416. [10.1164/rccm.200604-571ST](https://doi.org/10.1164/rccm.200604-571ST)
3. Tortoli E: Clinical manifestations of nontuberculous mycobacteria infections. *Clin Microbiol Infect*. 2009, 15:906-10. [10.1111/j.1469-0691.2009.03014.x](https://doi.org/10.1111/j.1469-0691.2009.03014.x)
4. Disseminated *Mycobacterium avium* complex disease. (2019). Accessed: September 24, 2023: <https://clinicalinfo.hiv.gov/en/guidelines/hiv-clinical-guidelines-adult-and-adolescent-opportunistic-infections/diss....>
5. Chin KL, Sarmiento ME, Alvarez-Cabrera N, Norazmi MN, Acosta A: Pulmonary non-tuberculous mycobacterial infections: current state and future management. *Eur J Clin Microbiol Infect Dis*. 2020, 39:799-826. [10.1007/s10096-019-03771-0](https://doi.org/10.1007/s10096-019-03771-0)
6. *Mycobacterium avium* complex - an overview. (2023). Accessed: September 24, 2023: <https://www.sciencedirect.com/topics/biochemistry-genetics-and-molecular-biology/mycobacterium-avium-complex>.
7. Catanzaro A: Diagnosis, differentiating colonization, infection, and disease. *Clin Chest Med*. 2002, 23:599-601. [10.1016/S0272-5231\(02\)00020-5](https://doi.org/10.1016/S0272-5231(02)00020-5)
8. Fujita K, Ito Y, Hirai T, Kubo T, Togashi K, Ichihara S, Mishima M: Prevalence and risk factors for chronic co-infection in pulmonary *Mycobacterium avium* complex disease. *BMJ Open Respir Res*. 2014, 1:e000050. [10.1136/bmjresp-2014-000050](https://doi.org/10.1136/bmjresp-2014-000050)
9. Urabe N, Sakamoto S, Shimanuki Y, et al.: Impact of chronic co-infection in pulmonary *Mycobacterium avium* complex disease after treatment initiation. *BMC Pulm Med*. 2022, 22:157. [10.1186/s12890-022-01947-7](https://doi.org/10.1186/s12890-022-01947-7)
10. Joao I, Bujdaková H, Jordao L: Opportunist coinfections by nontuberculous mycobacteria and fungi in immunocompromised patients. *Antibiotics (Basel)*. 2020, 9:771. [10.3390/antibiotics9110771](https://doi.org/10.3390/antibiotics9110771)
11. Bazzi AM, Abulhamayel Y, Rabaan AA, Al-Tawfiq JA: The impact of the coexistence of *Mycobacterium avium* with *Mycobacterium tuberculosis* on the result of GeneXpert and MGIT susceptibility test. *J Infect Public Health*. 2020, 13:827-9. [10.1016/j.jiph.2020.01.006](https://doi.org/10.1016/j.jiph.2020.01.006)
12. Abdeldaim G, Svensson E, Blomberg J, Herrmann B: Duplex detection of the *Mycobacterium tuberculosis* complex and medically important non-tuberculosis mycobacteria by real-time PCR based on the *rnpB* gene.

- APMIS. 2016, 124:991-5. [10.1111/apm.12598](https://doi.org/10.1111/apm.12598)
13. Khan Z, Miller A, Bachan M, Donath J: Mycobacterium avium complex (MAC) lung disease in two inner city community hospitals: recognition, prevalence, co-infection with Mycobacterium tuberculosis (MTB) and pulmonary function (PF) improvements after treatment. Open Respir Med J. 2010, 4:76-81. [10.2174/1874306401004010076](https://doi.org/10.2174/1874306401004010076)