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Retrospective analysis of patients with non-tuberculous mycobacteria from a primary hospital in Southeast China

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To achieve a comprehensive understanding of the characteristics of patients with non-tuberculous mycobacteria (NTM), patients with NTM between January 2016 and June 2019 were recruited from a primary hospital. NTM were identified based on the MBP64 protein assay. The clinical records and laboratory assay results were retrospectively reviewed. A total of 204 patients with NTM were included in the final analysis. The patients with multiple isolations were more likely accompanied with chronic obstructive pulmonary disease (COPD) ($p = 0.029$) and arthritis ($p = 0.049$), but showed a lower percentage of positive T-spot results ($p = 0.022$). In addition, patients with multiple isolations showed a higher rate of positive acid-fast staining results and their symptom duration was more likely longer than 30 days ($p = 0.019$). Patients with a positive response in T-spot assay showed a higher proportion of nodular manifestation on computed tomography (CT) than those with a negative response. Compared with male patients with NTM, female patients showed lower rates of positive acid-fast staining results ($p = 0.03$), but were more likely accompanied with COPD ($p < 0.0001$). The positive acid-fast staining results were closely associated with pulmonary cavities and tuberculosis antibody. Patients with different NTM isolation frequencies were closely associated with coexisting diseases and examination results.

As environmental microbes, non-tuberculous mycobacteria (NTM) can colonize mainly in the lungs and other sites by direct exposure, causing diseases when the balance between the host defense and bacteria is disrupted. The prevalence of NTM diseases has increased in recent years and the 5-year morbidity has been reported to be as high as 17.8% in Korea¹. Although the clinical significance of NTM isolates remains to be evaluated², NTM diseases cause substantial economic burden³.

Given the fact that NTM-related diseases are not mandatory to be treated, patients with NTM are always mis-diagnosed as other chronic diseases such as tuberculosis or cancer^{4,5}. Their common symptoms such as cough and expectoration result in difficulties in diagnosis, leading to inappropriate diagnosis and treatment. In fact, in countries with a high incidence rate of tuberculosis such as China, once a patient shows positive results in acid-fast staining assay and T-spot assay, he/she is likely to be diagnosed with tuberculosis and undergo tuberculosis treatment^{6,7}. When species identification tests are unavailable or not conducted, patients are not suspected to have NTM infection until they are not susceptible to anti-mycobacterial drugs. Thus, the prevalence and significance of NTM infections are mostly underestimated in China.

Based on the principles of NTM diagnosis proposed by ATS/IDSA, repeated isolation in sputum is required to confirm a pulmonary infection; however, two or more sample collections were mostly unavailable in hospitals in China as NTM surveillance is not mandatory⁸. Sample isolation frequency may help to the discern colonized isolates from the pathogenic ones. Our previous work showed a significant difference between patients with MTBC and those with NTM, but the exact contributions of the isolated NTM to diseases were not determined⁹. The potential difference in patient characteristics for different NTM isolation frequencies remained largely unknown and required to be evaluated urgently¹⁰.

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| Features | Values | Single isolation (n = 169) | Multiple isolation (n = 35) | P value |
|-----------------------------------------|-------------------|-------------------------------|--------------------------------|---------------|
| Gender | Male | 94 (55.6) | 17 (48.6) | 0.446 |
| | Female | 75 (44.4) | 18 (51.4) | |
| Age | <60 | 40 (23.7) | 4 (11.4) | 0.128* |
| | 60~80 | 100 (59.2) | 27 (77.1) | |
| | ≥80 | 29 (17.2) | 4 (11.4) | |
| Occupation | Non-farmer | 30 (17.8) | 4 (11.4) | 0.361 |
| | farmer | 139 (82.2) | 31 (88.6) | |
| Smoke history | Yes | 54 (32.0) | 13 (37.1) | 0.552 |
| | No | 115 (68.0) | 22 (62.9) | |
| Tuberculosis history | Yes | 30 (17.8) | 9 (25.7) | 0.276 |
| | No | 139 (82.2) | 26 (74.3) | |
| Surgeon history | Yes | 55 (32.5) | 21 (60.0) | 0.002 |
| | No | 114 (67.5) | 14 (40.0) | |
| Acid fast assay | Positive | 35 (22.4) | 18 (51.4) | 0.01 |
| | Negative | 121 (77.6) | 17 (48.6) | |
| CT | Nodular | 81 (47.9) | 18 (51.4) | 0.706 |
| | Non-nodular | 88 (52.1) | 17 (48.6) | |
| Cavities | Yes | 13 (7.7) | 7 (20.0) | 0.053* |
| | No | 156 (92.3) | 28 (80.0) | |
| T. spot | Positive | 30 (44.8) | 3 (15.8) | 0.022 |
| | Negative | 37 (55.2) | 16 (84.2) | |
| Tuberculosis antibody | Positive | 20 (15.2) | 7 (23.3) | 0.278 |
| | Negative | 112 (84.8) | 23 (76.7) | |
| COPD | Yes | 50 (29.6) | 17 (48.6) | 0.029 |
| | No | 119 (70.4) | 18 (51.4) | |
| Bronchiectasis | Yes | 39 (23.1) | 8 (22.9) | 0.978 |
| | No | 130 (76.9) | 27 (77.1) | |
| Diabetes | Yes | 11 (6.5) | 5 (14.3) | 0.159* |
| | No | 158 (93.5) | 30 (85.7) | |
| Tumor | Yes | 13 (7.7) | 3 (8.6) | 0.741* |
| | No | 156 (92.3) | 32 (91.4) | |
| Arthritis | Yes | 5 (3.0) | 4 (11.4) | 0.049 |
| | No | 164 (97.0) | 31 (88.6) | |
| Diagnosis when discharged from hospital | Tuberculosis | 49 (30.0) | 10 (28.6) | 0.005* |
| | NTM | 9 (5.3) | 9 (25.7) | |
| | Non-mycobacterial | 111 (65.7) | 16 (45.7) | |
| Duration of symptoms (days) | ≥30 | 89 (52.7) | 26 (74.3) | 0.019 |
| | <30 | 80 (47.3) | 9 (25.7) | |

Table 1. The characteristics of patients with single species and multiple species of non-tuberculosis mycobacteria. *Fisher exact test. The bold values mean significantly with p value less than 0.05.

In this study, the characteristics of patients with NTM were retrospectively analyzed based on the data from a primary hospital located in central Zhejiang Province, in an attempt to determine the potential risk factors associated with NTM diseases. Our results could help clinical staff to comprehensively understand the characteristics of patients with NTM.

Materials and Methods

The strains were isolated from inpatients admitted at a primary hospital between January 2016 and June 2019. Samples were analyzed via acid-fast staining assay and mycobacterial culture using BACTEC Mycobacterial Growth Indicator Tube 320 systems (BD, USA), simultaneously. When the culture results were positive, the morphology was confirmed by positive acid-fast staining response. Then, the mycobacterial strains were tested via MPB64 protein assay (Hangzhou Genesis, Hangzhou, China) and identified as NTM by a negative response or MTBC by a positive response¹¹. The clinical records of inpatients with corresponding NTM isolates were anonymously reviewed and informed consent was obtained from all participants. All experimental protocols were approved by the Ethics Committee of Dongyang People's Hospital Ethics Committee and Institutional Review Board. All methods were carried out in accordance with relevant guidelines and regulations.

Basic information including gender, age, occupation, smoke history, tuberculosis history, and surgery history was collected. Laboratory examinations included acid-fast staining assay followed by mycobacterial culture,

| Features | Values | T. spot Positive (n = 33) | T. spot Negative (n = 53) | P |
|-----------------------------------------|-------------------|---------------------------|---------------------------|---------------|
| Gender | Male | 23 (69.7) | 29 (54.7) | 0.167 |
| | Female | 10 (30.3) | 24 (45.3) | |
| Age | <60 | 11 (33.3) | 8 (15.1) | 0.034 |
| | 60~80 | 16 (48.5) | 40 (75.5) | |
| | ≥80 | 6 (18.2) | 5 (9.4) | |
| Occupation | Non-farmer | 5 (15.2) | 6 (11.3) | 0.742* |
| | Farmer | 28 (84.8) | 47 (88.7) | |
| Smoke history | Yes | 14 (42.4) | 19 (35.8) | 0.542 |
| | No | 19 (57.6) | 34 (64.2) | |
| Tuberculosis history | Yes | 3 (9.1) | 13 (24.5) | 0.074 |
| | No | 30 (90.9) | 40 (75.5) | |
| Surgeon history | Yes | 12 (36.4) | 22 (41.5) | 0.635 |
| | No | 21 (63.6) | 31 (58.5) | |
| Acid fast assay | Negative | 22 (68.8) | 33 (62.3) | 0.544 |
| | Positive | 10 (31.3) | 20 (37.7) | |
| CT | Nodular | 25 (75.8) | 24 (45.3) | 0.006 |
| | Non-nodular | 8 (24.2) | 29 (54.7) | |
| Cavities | Yes | 3 (9.1) | 10 (18.9) | 0.354* |
| | No | 30 (90.9) | 43 (81.1) | |
| Tuberculosis antibody | Positive | 5 (16.7) | 8 (15.1) | 1* |
| | Negative | 25 (83.3) | 45 (84.9) | |
| COPD | Yes | 7 (21.2) | 16 (30.2) | 0.36 |
| | No | 26 (78.8) | 37 (69.8) | |
| Bronchiectasis | Yes | 2 (6.1) | 11 (20.8) | 0.119* |
| | No | 31 (93.9) | 42 (79.2) | |
| Diabetes | Yes | 3 (9.1) | 4 (7.5) | 1* |
| | No | 30 (90.9) | 49 (92.5) | |
| Tumor | Yes | 5 (15.2) | 2 (3.8) | 0.101* |
| | No | 28 (84.8) | 51 (96.2) | |
| Diagnosis when discharged from hospital | Tuberculosis | 22 (66.7) | 16 (30.2) | 0.004* |
| Duration of symptoms (days) | NTM | 2 (6.1) | 9 (17.0) | |
| | Non-mycobacterial | 9 (27.3) | 28 (52.8) | |
| Duration of symptoms (days) | ≥30 | 17 (51.5) | 27 (50.9) | 1 |
| | <30 | 16 (48.5) | 26 (49.1) | |

Table 2. The characteristics of patients with positive and negative response of T. spot assay. *Fisher exact test. The bold values mean significantly with p value less than 0.05.

T-spot assay, and tuberculosis antibody assay. The results of computed tomography (CT) of the lung were also reviewed and the presence of nodules or cavities was checked. The coexisting illnesses such as COPD, bronchiectasis, cancer, diabetes, and arthritis were included in the evaluation.

Patients with a single isolation of NTM were referred to as the ones from whom NTM was isolated only once despite the number of samples being sourced and at different intervals at the hospital. Patients with multiple NTM isolations were referred to as the ones from whom NTM was isolated two or more times based on multiple clinical sample sources. If the MPB64 protein assay showed inconsistent results in the samples of patients with multiple isolations, the corresponding patients were excluded from the analysis. When the data for a single patient was found in different years, the latest information was collected and examination results throughout the researched years were reviewed. The suspected or assured diagnosis at discharge was closely associated with later treatment or healthcare; thus, the diagnosis could be classified into tuberculosis, non-tuberculous mycobacterium disease, and non-mycobacterial infection. The duration of symptoms refers to the time from symptom onset to the time when the patient visited the hospital; thus, the symptoms may not be occurring continuously.

Statistical analysis. All data from the inpatients were categorized into different groups and the differences were analyzed by X^2 test or fisher exact test using 20.0 version SPSS software (IBM, USA). The p values less than 0.05 were considered statistically significant between the analyzed groups.

Results

A total of 292 NTM strains were identified by MPB64 assay. After excluding repeated samples from the same patients and patients without available clinical records, 204 inpatients were finally analyzed in this study.

| Features | Values | Male (n = 111) | Female (n = 93) | P |
|-----------------------------------------|-------------------|----------------|-----------------|-------------------|
| Age | <60 | 22 (19.8) | 19 (20.4) | 0.728 |
| | 60~80 | 65 (58.6) | 58 (62.4) | |
| | ≥80 | 24 (21.6) | 16 (17.2) | |
| Occupation | Non-farmer | 23 (20.7) | 11 (11.8) | 0.09 |
| | Farmer | 88 (79.3) | 82 (88.2) | |
| Smoke history | Yes | 66 (59.5) | 1 (1.1) | <0.0001 |
| | No | 45 (40.5) | 92 (98.9) | |
| Tuberculosis history | Yes | 23 (20.7) | 16 (17.2) | 0.525 |
| | No | 88 (79.3) | 77 (82.8) | |
| Surgeon history | Yes | 39 (35.1) | 37 (39.8) | 0.494 |
| | No | 72 (64.9) | 56 (60.2) | |
| Acid fast assay | Negative | 67 (65.7) | 71 (79.8) | 0.03 |
| | Positive | 35 (34.3) | 18 (20.2) | |
| Tuberculosis antibody | Positive | 17 (19.8) | 10 (13.2) | 0.26 |
| | Negative | 69 (80.2) | 66 (86.8) | |
| CT | Nodular | 54 (48.6) | 45 (48.4) | 0.97 |
| | Non-nodular | 57 (51.4) | 48 (51.6) | |
| Cavities | Yes | 14 (12.6) | 6 (6.5) | 0.141 |
| | No | 97 (87.4) | 87 (93.5) | |
| COPD | Yes | 42 (37.8) | 25 (26.9) | 0.097 |
| | No | 69 (62.2) | 68 (73.1) | |
| Bronchiectasis | Yes | 12 (10.8) | 35 (37.6) | <0.0001 |
| | No | 99 (89.2) | 58 (62.4) | |
| Diabetes | Yes | 10 (9.0) | 6 (6.5) | 0.499 |
| | No | 101 (91.0) | 87 (93.5) | |
| Tumor | Yes | 12 (10.8) | 4 (4.3) | 0.085 |
| | No | 99 (89.2) | 89 (95.7) | |
| Diagnosis when discharged from hospital | Tuberculosis | 12 (10.8) | 22 (23.6) | <0.0001 |
| | NTM | 37 (33.3) | 5 (5.4) | |
| | Non-mycobacterial | 62 (55.9) | 66 (71.0) | |
| Duration of symptoms (days) | ≥30 | 60 (54.1) | 55 (59.1) | 0.466 |
| | <30 | 51 (45.9) | 38 (40.9) | |

Table 3. The characteristics of patients with non-tuberculous mycobacteria between male and female. *Fisher exact test. The bold values mean significantly with p value less than 0.05.

Considering that the isolation frequency from sputum was significantly important for the diagnosis of pulmonary NTM, the differences between the patients with single and multiple isolations were analyzed (Table 1). Compared with patients with single isolation, those with multiple NTM isolations showed a significantly higher percentage of surgery history, whereas the two populations were similar in terms of gender, age, occupation, and smoking history. Among the laboratory assay results used for tuberculosis, the patients with multiple isolations showed a significantly higher frequency of positive acid-fast staining results ($p = 0.01$), but showed a lower percentage of positive T-spot results ($p = 0.022$). In addition, patients with multiple isolations were more likely accompanied with COPD ($p = 0.029$) and arthritis ($p = 0.049$), but not with diabetes, cancer, or bronchiectasis. When discharged from the hospital, significantly more patients with multiple isolations were diagnosed with NTM-related diseases. Finally, patients with a single isolation had experienced the symptoms for a shorter duration (less than 30 days) before visiting the hospital.

Because T-spot assay was of significance to distinguish individuals with latent or active TB from healthy individuals, it could also help distinguish between active TB patients and NTM patients among mycobacterial culture-positive patients. The risk factors of a positive T-spot response in patients with NTM are shown in Table 2. Patients with a positive T-spot response showed a higher frequency of nodular manifestations on CT compared with patients with a negative response ($p = 0.006$); further, a higher proportion of patients with a negative response were aged between 60 and 80 years ($p = 0.034$).

Table 3 shows differences between genders in terms of characteristics of patients with NTM. Approximately 20% female patients showed positive acid-fast staining results, which is significantly less compared with the male patients (34.3%; $p = 0.03$). Moreover, 37.6% of female patients were accompanied with bronchiectasis; this was significantly higher than the percentage in male patients (10.8%, $p < 0.0001$). The percentage of patients diagnosed with NTM at hospital discharge was higher in males than in females (33.3% vs 5.4%, $p < 0.0001$).

Imaging manifestations in CT scans were important in the diagnosis of pulmonary infections including pulmonary tuberculosis and NTM diseases. In addition, multiple small nodules were common manifestations of pulmonary MAC diseases. Table 4 shows the characteristics of patients with NTM and nodules. Patients with

| Features | Values | Nodular (n = 99) | Non-nodular (n = 105) | |
|-----------------------------------------|-------------------|------------------|-----------------------|--------------|
| Age | <60 | 28 (28.3) | 13 (12.4) | 0.015 |
| | 60~80 | 52 (52.5) | 71 (67.6) | |
| | ≥80 | 19 (19.2) | 21 (20.0) | |
| Occupation | Non-farmer | 15 (15.2) | 19 (18.1) | 0.573 |
| | Farmer | 84 (84.8) | 86 (81.9) | |
| Smoke history | Yes | 32 (32.3) | 35 (33.3) | 0.878 |
| | No | 67 (67.7) | 70 (66.7) | |
| Tuberculosis history | Yes | 20 (20.2) | 19 (18.1) | 0.702 |
| | No | 79 (79.8) | 86 (81.9) | |
| Surgeon history | Yes | 36 (36.4) | 40 (38.1) | 0.798 |
| | No | 63 (63.6) | 65 (61.9) | |
| Acid fast assay | Negative | 65 (71.4) | 73 (73.0) | 0.809 |
| | Positive | 26 (28.6) | 27 (27.0) | |
| Tuberculosis antibody | Positive | 14 (17.5) | 13 (15.9) | 0.779 |
| | Negative | 66 (82.5) | 69 (84.1) | |
| Cavities | Yes | 12 (12.1) | 8 (7.6) | 0.28 |
| | No | 87 (87.9) | 97 (92.4) | |
| COPD | Yes | 36 (36.4) | 40 (38.1) | 0.798 |
| | No | 63 (63.4) | 65 (61.9) | |
| Bronchiectasis | Yes | 33 (33.3) | 26 (24.8) | 0.177 |
| | No | 66 (66.7) | 79 (75.2) | |
| Diabetes | Yes | 8 (8.1) | 8 (7.6) | 0.902 |
| | No | 91 (91.9) | 97 (92.4) | |
| Tumor | Yes | 10 (13.1) | 6 (5.7) | 0.244 |
| | No | 89 (86.9) | 99 (94.3) | |
| Diagnosis when discharged from hospital | Tuberculosis | 35 (35.4) | 24 (22.9) | 0.107 |
| | NTM | 6 (6.1) | 11 (10.5) | |
| | Non-mycobacterial | 58 (58.6) | 70 (66.7) | |
| Duration of symptoms (days) | ≥30 | 55 (55.6) | 60 (57.1) | 0.819 |
| | <30 | 44 (44.4) | 45 (42.9) | |

Table 4. The characteristics of patients with non-tuberculous mycobacteria between nodular and non-nodular. *Fisher exact test. The bold values mean significantly with p value less than 0.05.

nodules were slightly younger than those without nodules. Other characteristics including demographic features, laboratory examinations, and coexisting diseases were comparable between these two groups.

Further, the percentage of tuberculosis history, cavities in lung, and positive response in tuberculosis antibody assay was significantly higher in patients with a positive acid-fast staining result than in those with a negative result (Table 5). In addition, patients with positive acid-fast staining results were more likely to be diagnosed with mycobacterial infections including tuberculosis and NTM diseases at hospital discharge.

Discussions

Although the prevalence of NTM increased and its clinical significance in infections begins to be established, a comprehensive understanding of the clinical manifestation and risk factors for NTM infections remain unclear. In this study, the characteristics of inpatients with NTM and the potential risk factors of NTM patients were comprehensively analyzed. We found that the isolation frequency of NTM was associated with laboratory examinations (such as T-spot assay and acid-fast staining assay), coexisting diseases (such as COPD and arthritis), and clinical manifestations (e.g., symptom duration). Further, there were significant differences among patients with NTM in terms of gender, T-spot assay results, and acid-fast staining assay results.

Repeated exposure to NTM in the environment was the main mode of transmission of NTM, and a least of two sputum samples were needed to confirm a diagnosis of pulmonary NTM^{8,12}. Moreover, the treatment was not mandatory in patients from whom NTM was isolated, and was determined by the severity of infection as evaluated by the doctors. As reported in a study from Singapore, we found that the patients with more than one isolate were more likely to be accompanied with COPD¹⁰. A higher percentage of surgery history in multiple isolates suggested that NTM infection may have been hospital acquired¹³. Repeat isolation of NTM from the same patient provided valuable information for the diagnosis of NTM diseases and suggest a persistent infection (duration of symptoms longer than 30 days in this study). There is a huge data to suggest that chronicity of several NTMs is associated with genetic defects in IFN- γ /IL-12 axis, especially in patients with disseminated NTM diseases¹⁴. Besides, the patients with NTM infection commonly express high levels of anti-IFN- γ autoantibodies and suffer from recurrent infection^{15,16}.

| Features | Values | Acid-fast positive (n = 53) | Acid-fast negative (n = 138) | P |
|-----------------------------------------|-------------------|-----------------------------|------------------------------|-------------------|
| Age | <60 | 12 (22.6) | 26 (18.8) | 0.42 |
| | 60~80 | 28 (52.8) | 87 (63.0) | |
| | ≥80 | 13 (24.5) | 25 (18.1) | |
| Occupation | Non-farmer | 9 (17.0) | 23 (16.7) | 0.958 |
| | Farmer | 44 (83.0) | 115 (83.3) | |
| Smoke history | Yes | 19 (35.8) | 44 (31.9) | 0.602 |
| | No | 34 (64.2) | 94 (68.1) | |
| Tuberculosis history | Yes | 18 (34.0) | 19 (13.8) | 0.002 |
| | No | 35 (66.0) | 119 (86.2) | |
| Surgeon history | Yes | 23 (43.4) | 50 (36.2) | 0.362 |
| | No | 30 (56.6) | 88 (63.8) | |
| Tuberculosis antibody | Positive | 14 (32.6) | 12 (10.5) | 0.001 |
| | Negative | 29 (67.4) | 102 (89.5) | |
| Cavities | Yes | 14 (26.4) | 5 (3.6) | <0.001 |
| | No | 39 (73.6) | 133 (96.4) | |
| COPD | Yes | 20 (37.7) | 46 (33.3) | 0.567 |
| | No | 33 (62.3) | 92 (66.7) | |
| Bronchiectasis | Yes | 14 (26.4) | 31 (22.5) | 0.564 |
| | No | 39 (73.6) | 107 (77.5) | |
| Diabetes | Yes | 6 (11.3) | 8 (5.8) | 0.218* |
| | No | 47 (88.7) | 130 (94.2) | |
| Tumor | Yes | 4 (7.5) | 10 (7.2) | 1* |
| | No | 49 (92.5) | 128 (92.8) | |
| Diagnosis when discharged from hospital | Tuberculosis | 28 (52.8) | 27 (19.6) | <0.001* |
| discharged from hospital | NTM | 13 (24.5) | 4 (2.9) | |
| | Non-mycobacterial | 12 (22.6) | 107 (77.5) | |
| Duration of symptoms (days) | ≥30 | 34 (64.2) | 78 (56.5) | 0.338 |
| | <30 | 19 (35.8) | 60 (43.5) | |

Table 5. The characteristics of patients with non-tuberculous mycobacteria between smear positive and smear negative. *Fisher exact test. The bold values mean significantly with p value less than 0.05.

NTM was more likely to infect older females as reported previously^{1,17,18}; this may suggest that estrogen has a protective role against NTM¹⁸. However, in other studies including our study, no significant difference of distribution was found between genders^{2,10}, and the prevalence of NTM in younger individuals should also be given attention⁶. The difference of prevalence between genders may be due to the heterogeneity of the infection-causing microorganism among different regions² or the distribution of ages between genders¹⁹. In our study, female patients with NTM showed a significantly higher percentage of bronchiectasis than male patients, similar to the results reported by Zhang et al in Singapore¹⁰. In addition, authors from the same study found that a higher proportion of male patients had COPD, but this difference was not significant in our study¹⁰. Although several studies have shown that structural lung diseases such as COPD and bronchiectasis predispose people to NTM, the gender may play a potential role in determining the risk of coexisting diseases in patients with NTM. This phenomenon is urgently needed to be proven in a larger population and the mechanisms should be explored in the future.

The laboratory examinations for NTM were limited. For the diagnosis of MAC diseases, the potential of anti-GPL (glycopeptidolipid) IgA in the serum has been researched and has indicated encouraging results in terms of specificity and sensitivity^{20,21}. However, considering the diversity in species composition and largely unknown epidemiology of NTM in China, diagnosis using anti-GPL IgA needs to be investigated in the future. The molecular identification including genes (rpoB, hsp65, 16S rRNA and ITS region) sequencing rely on a large number of the strains, and should be performed based on a positive culture²²⁻²⁴. LPA and GeneXpert assays (detecting MTB/RIF) could differentiate MTB and NTM with considerable specificity and sensitivity in a broad range of sample sources, but it has not been applied commonly in primary hospitals because of requiring special equipment²⁵. Thus, the acid-fast staining assay of samples before culture and immunological assays such as IFN- γ release assay and T-spot assay would provide timely information for the diagnosis of active or latent tuberculosis^{26,27}. However, patients with NTM have shown a high proportion of positive T-spot results in our previous study and another study^{9,28}. In this study, a positive T-spot assay result in patients with NTM was associated with nodular manifestation in CT, indicating that a potential relation between nodule formation and immunological response may exist in these patients. In addition, positive results in acid-fast staining assay in patients with NTM suggest a large amount of isolates in individuals, resulting in a severe response in pulmonary infections, shown as cavities in the lung^{29,30}. The risk factors determined in laboratory examinations would help us evaluate

the progress of infection in patients with NTM. The limitations of this study include the possibility of colonization and infection both in patients with single isolations. This leads to a difficulty in establishing an association between the isolated strains and the diseases the patient suffers; however, this is a common phenomenon in China where NTM infections have not been paid enough attention, especially in primary hospitals and rural regions. In the future, patients from whom NTM strains are isolated should be recruited and checked to determine whether the isolates exist persistently.

Conclusions

Although the NTM related infection has been increasingly reported worldwide, most NTM cases are derived from the patients suspected of tuberculosis^{31,32}. Routine diagnosis of tuberculosis is inadequate for accurate diagnosis of NTM infection, which leads to underestimated prevalence^{33,34}. When molecular methods for species identification are non-available in resources limited region, comprehensive analysis of patients with NTM strains will shed light on realizing the importance of NTM infections in primary hospitals. In addition, laboratory examination results are closely associated with clinical or radiological manifestations in this study, indicating a comprehensive analysis of risk factors was necessary in diagnosis of NTM related diseases.

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References

- Park, S. C. *et al.* Prevalence, incidence, and mortality of nontuberculous mycobacterial infection in Korea: a nationwide population-based study. *BMC Pulm Med.* **19**, 140 (2019).
- Mortazavi, Z. *et al.* Evaluating the clinical significance of nontuberculous mycobacteria isolated from respiratory samples in Iran: an often overlooked disease. *Infect Drug Resist.* **12**, 1917–1927 (2019).
- Goring, S. M. *et al.* The cost of Mycobacterium avium complex lung disease in Canada, France, Germany, and the United Kingdom: a nationally representative observational study. *BMC Health Serv Res.* **18**, 700 (2018).
- Ringshausen, F. C. *et al.* Prevalence of Nontuberculous Mycobacterial Pulmonary Disease, Germany, 2009–2014. *Emerg Infect Dis.* **22**, 1102–1105 (2016).
- Koh W. J. Nontuberculous Mycobacteria-Overview. *Microbiol Spectr.* **5** (2017).
- Mbeha, B., Mine, M., Motswaledi, M. S. & Dewar, J. Nontuberculous Mycobacteria, Botswana, 2011–2014. *Emerg Infect Dis.* **25**, 1401–1403 (2019).
- Hernandez-Garduno E. & Elwood R. K. Increasing incidence of nontuberculous mycobacteria, Taiwan, 2000–2008. *Emerg Infect Dis.* **16**, 1047; author reply 1047–1048(2010).
- Griffith, D. E. *et al.* An official ATS/IDSA statement: diagnosis, treatment, and prevention of nontuberculous mycobacterial diseases. *Am J Respir Crit Care Med.* **175**, 367–416 (2007).
- Pan, X. *et al.* Differences in clinical and laboratory features between patients with non-tuberculous Mycobacterium and Mycobacterium tuberculosis complex. *Infect Dis (Lond).* **51**, 613–614 (2019).
- Zhang, Z. X., Cherng, B. P. Z., Sng, L. H. & Tan, Y. E. Clinical and microbiological characteristics of non-tuberculous mycobacteria diseases in Singapore with a focus on pulmonary disease, 2012–2016. *BMC Infect Dis.* **19**, 436 (2019).
- Liu, W. *et al.* Phenotypic and genotypic characterization of pyrazinamide resistance among multidrug-resistant Mycobacterium tuberculosis clinical isolates in Hangzhou, China. *Clin Microbiol Infect.* **24**, 1016 e1011–1016 e1015 (2018).
- Ramos, A. L., Carvalho, T. & Guimaraes, J. T. The importance of multiple samples in mycobacterial recovery: A 10-year retrospective study. *Int J Mycobacteriol.* **8**, 175–179 (2019).
- Desai, A. N. & Hurtado, R. M. Infections and outbreaks of nontuberculous mycobacteria in hospital settings. *Curr Treat Options Infect Dis.* **10**, 169–181 (2018).
- Namkoong, H., Hasegawa, N. & Betsuyaku, T. Susceptibility genes for nontuberculous mycobacterial disease. *Nihon Rinsho Meneki Gakkai Kaishi.* **40**, 60–67 (2017).
- Kham-Ngam, I. *et al.* Epidemiology of and risk factors for extrapulmonary nontuberculous mycobacterial infections in Northeast Thailand. *PeerJ.* **6**, e5479 (2018).
- Krisnawati, D. I. *et al.* Functional neutralization of anti-IFN-gamma autoantibody in patients with nontuberculous mycobacteria infection. *Sci Rep.* **9**, 5682 (2019).
- Marras, T. K. *et al.* Nontuberculous mycobacterial lung infections in Ontario, Canada: clinical and microbiological characteristics. *Lung.* **188**, (289–299 (2010).
- Mirsaedi, M. & Sadikot, R. T. Gender susceptibility to mycobacterial infections in patients with non-CF bronchiectasis. *Int J Mycobacteriol.* **4**, 92–96 (2015).
- Lee, H. *et al.* Epidemiology of Nontuberculous Mycobacterial Infection, South Korea, 2007–2016. *Emerg Infect Dis.* **25**, 569–572 (2019).
- Kitada, S. *et al.* Levels of Antibody against Glycopeptidolipid Core as a Marker for Monitoring Treatment Response in Mycobacterium avium Complex Pulmonary Disease: a Prospective Cohort Study. *J Clin Microbiol.* **55**, 884–892 (2017).
- Shibata, Y. *et al.* Diagnostic test accuracy of anti-glycopeptidolipid-core IgA antibodies for Mycobacterium avium complex pulmonary disease: systematic review and meta-analysis. *Sci Rep.* **6**, 29325 (2016).
- Appak, O., Turkel, S., Esen, N. & Ozkutuk, A. A. Comparison of polymerase chain reaction-restriction enzyme analysis method and DNA sequence analysis results in the identification of non-tuberculous mycobacteria. *Acta Microbiol Immunol Hung.* **65**, 515–527 (2018).
- Brown-Elliott B. A. *et al.* Comparison of Two Commercial Matrix-Assisted Laser Desorption/Ionization-Time of Flight Mass Spectrometry (MALDI-TOF MS) Systems for Identification of Nontuberculous Mycobacteria. *Am J Clin Pathol.* (2019).
- Maleki, M. R., Kafil, H. S., Harzandi, N. & Moaddab, S. R. Identification of nontuberculous mycobacteria isolated from hospital water by sequence analysis of the hsp65 and 16S rRNA genes. *J Water Health.* **15**, 766–774 (2017).
- Deng, S. *et al.* Accuracy of Commercial Molecular Diagnostics for the Detection of Pulmonary Tuberculosis in China: A Systematic Review. *Sci Rep.* **9**, 4553 (2019).
- Ai, L. *et al.* Clinical value of interferon-gamma release assay in the diagnosis of active tuberculosis. *Exp Ther Med.* **18**, 1253–1257 (2019).
- Marks, S. M. *et al.* Estimates of Testing for Latent Tuberculosis Infection and Cost, United States, 2013. *Public Health Rep.* **134**, 522–527 (2019).
- Wang, M. S., Wang, J. L. & Wang, X. F. The performance of interferon-gamma release assay in nontuberculous mycobacterial diseases: a retrospective study in China. *BMC Pulm Med.* **16**, 163 (2016).

29. Kim, S. J. *et al.* Characteristics associated with progression in patients with of nontuberculous mycobacterial lung disease: a prospective cohort study. *BMC Pulm Med.* **17**, 5 (2017).
30. Lee, Y. *et al.* CT findings of pulmonary non-tuberculous mycobacterial infection in non-AIDS immunocompromised patients: a case-controlled comparison with immunocompetent patients. *Br J Radiol.* **86**, 20120209 (2013).
31. Davari, M. *et al.* Genetic Diversity and Prevalence of Nontuberculous Mycobacteria Isolated from Clinical Samples in Tehran, Iran. *Microb Drug Resist.* **25**, 264–270 (2019).
32. Xu, J. *et al.* Prevalence and risk factors of pulmonary nontuberculous mycobacterial infections in the Zhejiang Province of China. *Epidemiol Infect.* **147**, e269 (2019).
33. Guglielmetti, L. *et al.* Human infections due to nontuberculous mycobacteria: the infectious diseases and clinical microbiology specialists' point of view. *Future Microbiol.* **10**, 1467–1483 (2015).
34. Zhang, Z. *et al.* Differences in risk factors and drug susceptibility between *Mycobacterium avium* and *Mycobacterium intracellulare* lung diseases in China. *Int J Antimicrob Agents.* **45**, 491–495 (2015).

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Author contributions

J.S. collected and analyzed the patient data. S.Y. performed the management of the patients involved in this study. X.W. and S.J. were major contributor in and performance and reviewing the laboratory examinations. P.X. contributed the data analysis and manuscript writing. All authors read and approved the final manuscript.

Competing interests

The authors declare no competing interests.

Additional information

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