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Epidemiology and Clinical Features of Pulmonary Nontuberculous Mycobacteriosis in Nagasaki, Japan

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Abstract

Background and Objectives
Recent reports indicate that the incidence of nontuberculous mycobacterial-lung disease (NTM-LD) is increasing. This study aimed to investigate the epidemiology and clinical features of NTM-LD patients in Nagasaki prefecture, Japan to identify the negative prognostic factors for NTM-LD in Japan.

Methods
The medical records of patients newly diagnosed with NTM-LD in eleven hospitals in Nagasaki prefecture between January 2001 and February 2010 were reviewed. Data regarding the annual population of each region and the incidence of all forms of tuberculosis were collected to assess geographic variations in NTM-LD incidence, isolates, and radiological features.

Results
A total 975 patients were diagnosed with NTM-LD. The incidence increased over the study period and reached 11.0 and 10.1 per 100,000 population in 2008 and 2009, respectively. M. intracellulare was the most common pathogen in the southern region, and M. avium most common in other regions. The most common radiographic pattern was the nodular-bronchiectatic pattern. Age >60 years, body mass index <18.5 kg/m², underlying
lung disease, and cavitary pattern were the negative prognostic factors at the 1-year follow-up.

Conclusions

The incidence of NTM-LD has been increasing in Nagasaki prefecture. The isolates and radiographic features of patients vary markedly by region.

Introduction

Nontuberculous mycobacteria (NTM) are ubiquitous organisms that cause diverse types of infectious disease in human organs, including lung, lymphatic, disseminated, skin, soft tissue, and bone disease [1,2]. Several reports have suggested that the incidence of nontuberculous mycobacterial lung disease (NTM-LD) has recently been increasing [1,3–6]. Although determining the precise incidence and prevalence of NTM-LD is difficult, the incidence in Japan was estimated at 5.9 per 100,000 population in 2005 [7], which was higher than that of other countries in the same year [4,6,8]. Although Mycobacterium species vary markedly by geographic region, Mycobacterium avium-intracellulare complex (MAC) has been reported as the dominant species in NTM-LD [5,9]. In Japan, M. avium is most commonly isolated from the respiratory tract, followed by M. intracellulare. Nevertheless, the species involved in NTM-LD has been found to vary even within Japan [7].

Previous studies reported the several risk factors for NTM infection such as gastroesophageal reflux disease, chronic obstructive pulmonary disease (COPD), cystic fibrosis, history of tuberculosis and immune defects or diseases such as HIV infection, corticosteroid use [1,10]. Negative prognostic factors for HIV-negative patients with NTM-LD are high comorbidity level, advanced age, male sex, fibrocavitary disease, body mass index (BMI) < 18.5 kg/m², and anemia [11,12]. This study aimed to investigate the epidemiology and clinical features of NTM-LD patients in Nagasaki prefecture to identify the risk factors and negative prognostic factors in Japan.

Patients and Methods

Participants

Nagasaki prefecture provides approximately 14,700 hospital beds for the medical care of a population of approximately 1,500,000. The participants were selected by review of the microbiological database at 11 hospitals (total 4,218 beds total) in the prefecture between January 2001 and February 2010. The participating facilities were Nagasaki University Hospital (862 beds), Nagasaki Municipal Hospital (414 beds), Nagasaki Municipal Medical Center (217 beds), Japanese Red Cross Nagasaki Genbaku Hospital (360 beds), Japanese Red Cross Nagasaki Genbaku Isahaya Hospital (140 beds), Isahaya Health Insurance General Hospital (333 beds), National Hospital Organization Nagasaki Medical Center (650 beds), Izumikawa Hospital (120 beds), Sasebo City General Hospital (594 beds), Hokusho Central Hospital (224 beds), and Nagasaki Goto Central Hospital (304 beds). The sputum, bronchial wash and tissue samples were cultured using Ogawa egg-based media and/or a Mycobacteria Growth Indicator Tube (MGIT; Becton Dickinson Microbiology Systems, Sparks, MD, USA). Isolated specimens were identified using polymerase chain reaction (PCR) or DNA-DNA hybridization (DDH) for M. avium and M. intracellulare; DDH for M. kansasii, M. abscessus, M. kansasii, M. gordonae, M. chelonae,
M. szulgai, M. simiae, M. peregrinum, M. scrofulaceum and M. terrae; and 16S ribosomal RNA sequencing for M. shimoidei [13,14]. Patients with at least two separate culture-positive sputum samples or one culture-positive bronchial washing or lung tissue sample during the study period were included. Patients with a single positive sputum culture or PCR positive-culture negative samples were excluded.

Diagnosis of NTM-LD

Diagnosis was performed by specialists in infectious disease and/or respiratory medicine according to 2007 American Thoracic Society (ATS)/Infectious Diseases Society of America (IDSA) criteria [1]. In accordance with the criteria, patients with suspected infection with contaminated environmental Mycobacterium species were excluded from study participation. Patients who had been diagnosed with NTM-LD prior to the current study were also excluded.

Epidemiological Study

Among all the patients who had presented at the 11 hospitals during the study period, 975 had been diagnosed with NTM-LD according to the 2007 ATS/IDSA criteria. After dividing the prefecture into five regions on the basis of administrative boundaries and medical care zones, the patients were classified into one region based on the residence listed in their medical records. To estimate the annual incidence of NTM-LD in each region, data regarding the annual population of each region were collected from the Nagasaki prefectural government, and data regarding the incidence of all forms of tuberculosis collected from the healthcare center of each region. In the analysis of the radiological features, 228 patients were excluded due to lack of accurate information about chest computed tomography (CT) scan (Fig 1). Based on the review of the results of chest CT scan, the remaining 747 patients were classified into four patterns according to their radiological characteristics: the nodular-bronchiectatic (NB), cavitory (CAV), hypersensitivity pneumonitis-like, or unclassifiable pattern. The NB pattern and CAV pattern were defined by chest CT scan showing multifocal bronchiectasis with multiple small nodules or cavitary opacities, respectively. If a patient had the characteristics of both the NB and CAV patterns, the patient was classified in accordance with the dominant pattern. Using these data, geographic variations in NTM-LD incidence, isolates, and radiological features were assessed.

Patient Characteristics and Prognostic Factors

Of the 975 patients, 374 were excluded from analysis due to lack of sufficient clinical information in their medical records (Fig 1). The clinical course of the disease was assessed after 1 year of diagnosis. Patient outcome was evaluated by review of (1) clinical, (2) radiological and (3) microbiological responses. Clinical response was defined as resolution of symptoms such as fever, weight loss, cough, sputum, hemoptysis and general fatigue. Radiological response was defined as the clearing or improvement of opacity on chest radiographs/CT. Microbiological response was determined by culturing of sputum. Patients who had shown improvement or no change in the 3 factors and had experienced no deterioration were defined as having achieved “stability”. Patients who had not met the criteria for “stability” definition and had subsequently died from all causes were classified as having experienced “deterioration”.

Statistical Analysis

The data are presented as mean ± SD values. Statistical significance was evaluated using the chi-square test or the two-tailed Mann-Whitney test, with the level of significance set at
P < 0.05. All analyses were performed with GraphPad Prism 5.0 for Macintosh (GraphPad Software, La Jolla, CA, USA).

Ethics Statement
This study was approved by the Ethical Committee of the Nagasaki University Hospital. As this study was retrospective, and data were analyzed anonymously, the need to obtain patient consent was waived by the committee.

Results
Fig 2A shows the numbers of NTM-LD patients newly diagnosed each year between 2001 and 2009, the population of Nagasaki prefecture, the incidence of all tuberculosis cases, and the estimated incidence of NTM-LD between 2001 and 2009. As can be observed, 975 patients were newly diagnosed with NTM-LD. Regarding the incidence by year, the NTM-LD incidence gradually increased over the entire study period and especially in 2008 and 2009, when it
reached 11.0 and 10.1 per 100,000 population, respectively. The incidence was the highest in the southeastern region of the prefecture, reaching 18.2 and 21.2 per 100,000 population in 2008 and 2009, respectively. Regarding the incidence by age, men and women >60 years accounted for 83.8% of all patients, and the greatest increase in incidence occurred in men and women >60 years and in women 45–59 years (Fig 2B and 2C).

Table 1 shows that the most frequently identified pathogen was *M. intracellulare* (n = 432, 44.3%), followed by *M. avium* (n = 415, 42.6%), *M. abscessus* (n = 30, 3.1%), *M. avium-intracellulare* complex (n = 24, 2.5%), *M. kansasii* (n = 20, 2.1%), *M. gordonae* (n = 20, 2.1%), and other pathogens (n = 34, 3.5%). Although the incidence of infection with *M. avium* and *M. intracellulare* particularly increased over the study period (Fig 3), the distribution of the isolates varied throughout the prefecture, with *M. intracellulare* accounting for 77.5% of all the isolates detected in the southern region (Fig 4). The most common radiographic pattern was the NB pattern, which was observed in 591 cases (79.1%), followed by the CAV pattern, which was observed in 134 cases (17.9%). No patients were found to have the hypersensitivity-like pattern.
The percentage of patients with each pattern was found to vary by region, especially in the southern region, where CAV pattern was observed in 29.4% of the patients (Fig 5).

Of the 975 patients, the clinical information of 601 patients with a mean age of 71.2 ± 11.2 years was analyzed. Among these, 182 (30.3%) patients were male with a mean age of 73.0 ± 10.9 years and 419 (69.7%) female with a mean age of 70.5 ± 11.2 years. Regarding disease state, the underlying respiratory disease was identified in 89 (48.9%) male and 91 (21.7%) female patients (Table 2) and systemic complications (diabetes, collagen disease, malignant disease other than lung cancer, chronic kidney disease, and chronic hepatic disease) in 52 (28.6%) male and 87 (20.8%) female patients. The underlying disease in 53 (8.9%) patients was treated by the administration of corticosteroids and/or immunosuppressants.

As observed in Table 3, which shows the causes of immunodeficiency, no patient was infected with HIV. The NB pattern was the most common radiologic pattern, as observed in 479 (79.7%) of the 601 patients, followed by the CAV pattern in 94 (15.6%), and an unclassifiable pattern in 28 (4.7%). The number of patients with the NB pattern increased over the study period, whereas the number with the CAV pattern remained unchanged. Regarding treatment, 279 (46.4%) patients had been treated by the administration of 1–5 anti-mycobacterial drugs within 3 months after diagnosis for a mean duration of 14.8 months, 214 (33.9%) patients had been observed clinically without being administered medication for 1 year, and treatment data were unavailable for 108 (18.0%) after 1 year (Fig 1). The analysis of prognostic factors was performed with the 205 patients who had been followed up for at least 1 year after diagnosis. Of the 86 patients (42.0%) who had improved, 75 (87.2%) had been treated with anti-mycobacterial drugs and 11 (12.8%) had been treated with no medication. Poor prognostic factors were found to be age >60 years, BMI<18.5 kg/m², underlying lung disease, and radiographic CAV pattern (Table 4).

| Table 1. Isolates detected in 975 NTM-LD patients. |
|-----------------------------------------------|
| **Isolate** | **Number (%)** |
|-----------------|----------------|
| *M. intracellulare* | 432 (44.3) |
| *M. avium* | 415 (42.6) |
| *M. abscessus* | 30 (3.1) |
| MAC | 24 (2.5) |
| *M. kansasii* | 20 (2.1) |
| *M. gordonae* | 20 (2.1) |
| *M. cheloneae* | 7 (0.7) |
| *M. fortuitum* | 3 (0.3) |
| *M. szulgai* | 3 (0.3) |
| *M. simiae* | 2 (0.2) |
| *M. peregrinum* | 2 (0.2) |
| *M. scrofulaceum* | 2 (0.2) |
| *M. terrae* | 2 (0.2) |
| *M. shimoidei* | 1 (0.1) |
| PMI except for MAC | 7 (0.7) |
| Not identified | 5 (0.5) |

MAC, *Mycobacterium avium-intracellulare* complex; PMI, polymicrobial infection

*Identified using 16S rRNA.

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Discussion

NTM-LD has been increasing over the past few decades in many areas of the world [1,3–5,15]. The results of the present study revealed that the annual incidence of NTM-LD in Nagasaki prefecture, Japan exceeded 10.0 per 100,000 population in 2008 and 2009. This incidence is much higher than that of other developed countries over the past decade, which has been reported as 2.7–5.6/100,000 in the United States [4,16,17], 0.72–0.74/100,000 in France [18], 0.9–2.9/100,000 in the United Kingdom [19], 2.2–3.2/100,000 in Australia [6], 1.06–2.00/100,000 in Taiwan [8], and 1.08/100,000 in Denmark. The dynamics of NTM-LD may be affected by various factors, including behavioral [12], cultural, genetic, environmental characteristics [20], soil characteristics, income levels, and population densities [21,22]. In this study, men and women >60 years old accounted for 83.8% of all participants, and the proportion of this age group increased over the study period. This finding and those of previous reports [6,12,19] suggest that the increasing incidence of NTM-LD could be associated with the aging of society and the increasing number of patients with comorbidities. In addition, previous reports indicate that cross-immunity and the protection provided by infection with *M. tuberculosis* and use of the bacillus Calmette-Guerin (BCG) vaccine is associated with NTM infection [15,23–25]. Our study also showed that the decreasing rate of tuberculosis is inversely associated with the increase in NTM-LD. However, cross-immunity might not be a key component, since almost all Japanese have been vaccinated with BCG for over half a century, and the southern prefecture, which was found to have a high incidence of NTM-LD in this study, has been a moderately endemic area of tuberculosis (a constant incidence of 28.9–40.4/100,000) over the past decade.

*Mycobacterium* species are ubiquitous organisms present in water, soil, and room dust [26]. In this study, *M. avium* was the most commonly detected species in most regions of Nagasaki prefecture, although *M. intracellulare* was the most commonly detected in the southern region. This variation may be related to exposure to certain substances present in the soil that have been associated with NTM infection [27,28]. In Australia, Chou et al. reported that the presence of *M. intracellulare* infection was correlated with a shallower soil depth and agricultural activities that lead to soil disturbance [22]. As the southern region of Nagasaki prefecture is an
agricultural zone with soil well drained by an active volcano, in which the percentage of cultivated area to total area is twice as high as the average of the prefecture (26.5% vs. 12.3%), the soil and agricultural activities performed within it might have led to differences in the dominant isolate and high incidence in this region compared to that in other regions. Additionally, in the southern region, the frequency of radiographic CAV pattern was higher than other areas. As described above, high prevalence of tuberculosis might have led to high rate of CAV pattern.

In general, effective treatment of NTM-LD requires long-term treatment with multiple antimycobacterial drugs. ATS/IDSA guidelines recommend that patients with NTM-LD should be treated until negative cultures are obtained for at least 1 year [1]. In addition to sputum culture, assessment of symptom subscore [29], clinical symptoms [30], and radiological findings [30,31] are useful in predicting clinical response. As short-term response to initial treatment has been found to reflect long-term outcome [32], this study evaluated 1-year response by assessment of clinical symptoms, radiological findings, and microbiological responses. The factors of age >60 years, BMI <18.5 kg/m², underlying lung disease, and radiographic CAV pattern were found to be poor prognostic factors, which accords with the findings of Hayashi et al. [11], who found advanced age, systemic and/or respiratory comorbidities, non-NB radiographic features, BMI <18.5 kg/m², anemia, hypoalbuminemia, and erythrocyte sedimentation rate ≥50 mm/h.

**Fig 4. Geographic variation in isolates detected in NTM-LD patients.** Nine patients were residents of another prefecture. Map was obtained from the Geospatial Information Authority of Japan (http://www.gsi.go.jp/ENGLISH/index.html). MAC, *Mycobacterium avium*-intracellulare complex.
Fig 5. Geographic variation in radiographic patterns in NTM-LD patients. Four patients were residents of another prefecture. NB, nodular-bronchiectatic pattern; CAV, cavitary pattern.

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Table 2. Underlying respiratory diseases of NTM-LD.

| Underlying respiratory diseases                      | Total    | Male     | Female   | P value |
|------------------------------------------------------|----------|----------|----------|---------|
| Underlying lung disease                              | 189 (31.4) | 90 (49.5) | 99 (23.4) | <0.0001 |
| History of tuberculosis                             | 81 (13.5)  | 35 (19.2) | 46 (11.0) | 0.0065  |
| COPD                                                 | 35 (5.8)   | 22 (12.1) | 13 (3.1)  | <0.0001 |
| Lung cancer                                          | 33 (5.5)   | 17 (9.3)  | 16 (3.8)  | 0.0063  |
| Interstitial lung disease                            | 24 (4.0)   | 13 (7.1)  | 11 (2.6)  | 0.0094  |
| Fungal infection                                     | 15 (2.5)   | 9 (4.9)   | 6 (1.4)   | 0.0112  |
| History of lung surgery                              | 13 (2.2)   | 9 (4.9)   | 4 (1.0)   | 0.002   |
| Bronchial asthma                                     | 11 (1.8)   | 4 (2.2)   | 7 (1.7)   | 0.6578  |
| Chronic bronchitis                                   | 9 (1.5)    | 5 (2.7)   | 4 (1.0)   | 0.0964  |
| Silicosis                                            | 8 (1.3)    | 8 (4.4)   | 0 (0.0)   | <0.0001 |
| Bronchiectasis                                       | 2 (0.3)    | 1 (0.5)   | 1 (0.2)   | 0.5433  |
| Pulmonary sarcoidosis                                | 2 (0.3)    | 1 (0.5)   | 1 (0.2)   | 0.5433  |

COPD, Chronic Obstructive Pulmonary Disease.

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to be negative prognostic factors for all-cause mortality of HIV-negative NTM-LD patients. It also accords with those of Andréjak et al. [12], who found high comorbidity level, age >65 years, and male sex to be predictors of all-cause mortality for NTM-LD patients. Although it remains unclear whether treatment of NTM-LD should be started immediately after diagnosis or be delayed, these findings may be helpful in determining when to start treatment.

This study faced several limitations that should be considered when reviewing the findings. Although a relatively large number of hospitals (11 hospitals) in Nagasaki prefecture were examined, not all NTM-LD cases in Nagasaki could be investigated. Second, the patients whose sputum culture was NTM positive only once were excluded from further analysis. These limitations might have led to underestimation of the incidence of NTM-LD. Third, the long-term prognosis is not shown in this retrospective study since we could not obtain the sufficient data. Additionally, selection bias may have been a factor in selecting the duration of treatment and follow-up, specifically the selection of relatively brief treatment and follow-up periods. However, as described above, short-term response for initial treatment has been reported to reflect long-term outcome, so we believe that this study of prognostic factors 1 year after diagnosis provides useful information.

In conclusion, the incidence of NTM-LD has been increasing over the past decade in Nagasaki, Japan, a prefecture in which the isolates and radiographic patterns have varied markedly by region. Among all possible factors, older age, BMI <18.5 kg/m², underlying lung disease,
and radiographic CAV pattern appear to be negative prognostic factors in the deterioration of patients with the disease.

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Author Contributions
Conceived and designed the experiments: SI SN YY. Performed the experiments: SI. Analyzed the data: SI SN YY YK Y. Fukuda HI ES KY YH KH Y. Futsuki YI KF NS SK. Contributed reagents/materials/analysis tools: SI SN YY. Wrote the paper: SI SN.

References
1. Griffith DE, Aksamit T, Brown-Elliott BA, Catanzaro A, Daley C, Gordin F, et al. (2007) An Official ATS/IDSA Statement: Diagnosis, Treatment, and Prevention of Nontuberculous Mycobacterial Diseases. American Journal of Respiratory and Critical Care Medicine 175: 367–416. PMID:17277290
2. Wolinsky E (1979) Nontuberculous mycobacteria and associated diseases. American Review of Respiratory Disease 119: 107–159. PMID:426334
3. Iseman MD, Marras TK (2008) The importance of nontuberculous mycobacterial lung disease. American Journal of Respiratory and Critical Care Medicine 178: 999–1000. doi:10.1164/rccm.200808-1258ED PMID:18987345
4. Bodle EE, Cunningham JA, Della-Latta P, Schluger NW, Saiman L (2008) Epidemiology of nontuberculous mycobacteria in patients without HIV infection, New York City. Emerging Infectious Diseases 14: 390–396. doi: 10.3201/eid1403.061143 PMID:18325252
5. Prevots DR, Shaw PA, Strickland D, Jackson LA, Raebel MA, Biosky MA, et al. (2010) Nontuberculous mycobacterial lung disease prevalence at four integrated health care delivery systems. American Journal of Respiratory and Critical Care Medicine 182: 970–976. doi:10.1164/rccm.201002-0310OC PMID:20538958
6. Thomson RM; NTM working group at Queensland TB Control Centre and Queensland Mycobacterial Reference Laboratory (2010) Changing epidemiology of pulmonary nontuberculous mycobacteria infections. Emerging Infectious Diseases 16: 1576–1583. doi:10.3201/eid1610.091201 PMID:20875283
7. Kajiki A (2011) Non-tuberculous mycobacteriosis. What has been coming out. Kekkaku 86: 113–125. PMID:21404655
8. Chen C-Y, Chen H-Y, Chou C-H, Huang C-T, Lai C-C, Hsueh PR (2012) Pulmonary infection caused by nontuberculous mycobacteria in a medical center in Taiwan, 2005–2008. Diagnostic Microbiology and Infectious Disease 72: 47–51. doi: 10.1016/j.diagmicrobio.2011.09.009 PMID:22015322
9. Simons S, van Ingen J, Hsueh P-R, Van Hung N, Dekhuijzen PNR, Boeree MJ, et al. (2011) Nontuberculous mycobacteria in respiratory tract infections, eastern Asia. Emerging Infectious Diseases 17: 343–349. doi:10.3201/eid1703.100604 PMID:21392422
10. Kim RD, Greenberg DE, Ehrmantraut ME, Guide SV, Ding L, Shea Y, et al. (2008) Pulmonary nontuberculous mycobacterial disease: prospective study of a distinct preexisting syndrome. American Journal of Respiratory and Critical Care Medicine 178: 1066–1074. doi:10.1164/rccm.200805-0786OC PMID:18703788
11. Hayashi M, Takayanagi N, Kanauchi T, Miyahara Y, Yanagisawa T, Sugita Y (2012) Prognostic factors of 634 HIV-negative patients with Mycobacterium avium complex lung disease. American Journal of Respiratory and Critical Care Medicine 185: 575–583. doi:10.1164/rccm.201107-1203OC PMID:22199005
12. Andréjak C, Thomsen VØ, Johansen IS, Riis A, Benfield TL, Duhaert P, et al. (2010) Nontuberculous pulmonary mycobacteriosis in Denmark: incidence and prognostic factors. American Journal of Respiratory and Critical Care Medicine 181: 514–521. doi:10.1164/rccm.200905-0778OC PMID:20007929
13. Kusunoki S, Ezaki T, Tamesada M, Hatana Y, Asano K, Hashimoto Y, et al. (1991) Application of colormetric microdilution plate hybridization for rapid genetic identification of 22 Mycobacterium species. Journal of Clinical Microbiology 29: 1596–1603. PMID:1761680
14. Kazumi Y, Maeda S, Sugawara I (2006) Identification of mycobacteria by sequencing of rpoB gene and 16S rRNA. Kekkaku 81: 551–558. PMID: 17037387
15. Marras TK, Chedore P, Ying AM, Jamieson F (2007) Isolation prevalence of pulmonary non-tuberculous mycobacteria in Ontario, 1997 2003. Thorax 62: 661–666. PMID: 17311842
16. Winthrop KL, McNelley E, Kendall B, Marshall-Olson A, Morris C, Cassidy M, et al. (2010) Pulmonary nontuberculous mycobacterial disease prevalence and clinical features: an emerging public health disease. American Journal of Respiratory and Critical Care Medicine 182: 977–982. doi: 10.1164/rccm.201003-050OC PMID: 20508209
17. Cassidy PM, Hedberg K, Saulson A, McNelley E, Winthrop KL (2009) Nontuberculous mycobacterial disease prevalence and risk factors: a changing epidemiology. Clinical Infectious Diseases 49: e124–e129. doi: 10.1086/648443 PMID: 19911942
18. Dailloux M, Abalain ML, Laurain C, Lebrun L, Loos-Ayav C, Lozniewski A, et al. (2006) Respiratory infections associated with nontuberculous mycobacteria in non-HIV patients. European Respiratory Journal 28: 1211–1215. PMID: 17138678
19. Moore JE, Kruishaar ME, Ormerod LP, Drobniewski F, Abubakar I (2010) Increasing reports of non-tuberculous mycobacteria in England, Wales and Northern Ireland, 1995–2006. BMC Public Health 10: 612. doi: 10.1186/1471-2458-10-612 PMID: 20950421
20. Adjemian J, Olivier KN, Seitz AE, Holland SM, Prevots DR (2012) Prevalence of nontuberculous mycobacterial lung disease in U.S. Medicare beneficiaries. American Journal of Respiratory and Critical Care Medicine 185: 881–886. doi: 10.1164/rccm.201111-2016OC PMID: 23212016
21. Adjemian J, Olivier KN, Seitz AE, Falkinham JO, Holland SM, Prevots DR (2012) Spatial clusters of nontuberculous mycobacterial lung disease in the United States. American Journal of Respiratory and Critical Care Medicine 186: 553–558. doi: 10.1164/rccm.201205-0913OC PMID: 22773732
22. Chou MP, Clements ACA, Thomson RM (2014) A spatial epidemiological analysis of nontuberculous mycobacterial infections in Queensland, Australia. BMC Infectious Diseases 14: 279. doi: 10.1186/1471-2334-14-279 PMID: 24885916
23. Trnka L, Danková D, Svandová E (1994) Six years’ experience with the discontinuation of BCG vaccination. 4. Protective effect of BCG vaccination against the Mycobacterium avium intracellulare complex. Tubercle and Lung Disease 75: 348–352. PMID: 7841428
24. Romanus V, Hallander HO, Wählén P, Olander-Nielsen AM, Magnusson PH, Juhlin I (1995) Atypical mycobacteria in extrapolmonary disease among children. Incidence in Sweden from 1969 to 1990, related to changing BCG-vaccination coverage. Tubercle and Lung Disease 76: 300–310. PMID: 7579311
25. Hersh AL, Tala-Heikkilä M, Tala E, Tosteson ANA, Fordham von Reyn C (2003) A cost-effectiveness analysis of universal versus selective immunization with Mycobacterium bovis bacille Calmette-Guérin in Finland. International Journal of Tuberculosis and Lung Disease 7: 22–29. PMID: 12701831
26. Falkinham JO (2009) Surrounded by mycobacteria: nontuberculous mycobacteria in the human environment. Journal of Applied Microbiology 107: 356–367. doi: 10.1111/j.1365-2672.2009.04161.x PMID: 19228258
27. Reed C, Reyn von CF, Chamblee S, Ellerbrock TV, Johnson JW, Marsh BJ, et al. (2006) Environmental risk factors for infection with Mycobacterium avium complex. American Journal of Epidemiology 164: 32–40. PMID: 16675537
28. Fujita K, Ito Y, Hirai T, Maekawa K, Imai S, Tatsumi S, et al. (2012) Genetic relatedness of Mycobacterium avium-intracellulare complex isolates from patients with pulmonary MAC disease and their residential soils. Clinical Microbiology and Infection 19: 537–41. doi: 10.1111/j.1365-2672.2012.03929.x PMID: 22712883
29. Singer J, Thorne A, Khorasheh S, Raboud JM, Wu AW, Salit I, et al. (2000) Symptomatic and health status outcomes in the Canadian randomized MAC treatment trial (CTN010), Canadian HIV Trials Network Protocol 010 Study Group. International Journal of STD and AIDS 11: 212–219. PMID: 10772083
30. Kobashi Y, Matsushima T (2004) Comparison of clinical features in patients with pulmonary Mycobacterium avium complex (MAC) disease treated before and after proposal for guidelines. Journal of Infection and Chemotherapy 10: 25–30. PMID: 14991914
31. Fujiuchi S, Matsumoto H, Yamazaki Y, Nakao S, Takahashi M, Satoh K, et al. (2003) Analysis of chest CT in patients with Mycobacterium avium complex pulmonary disease. Respiration 70: 76–81. PMID: 12584395
32. Fujikane T, Fujiuchi S, Yamazaki Y, Sato M, Yamamoto Y, Takeda A, et al. (2005) Efficacy and outcomes of clarithromycin treatment for pulmonary MAC disease. International Journal of Tuberculosis and Lung Disease 9: 1281–1287. PMID: 16333938