Mycobacterial Infections (H Bach, Section Editor)

CrossMark

Treatment of Non-Tuberculous Mycobacterial Lung Disease

Julie V. Philley, MD¹ Mary Ann DeGroote, MD^{2,3} Jennifer R. Honda, PhD^{4,5} Michael M. Chan, PharmD^{6,7} Shannon Kasperbauer, MD^{4,2} Nicholas D. Walter, MD, PhD^{8,5} Edward D. Chan, MD^{4,8,5,*}

Address

¹Division of Pulmonary and Critical Care Medicine, University of Texas Health Science Center, University of Texas, Tyler, TX, USA

²Division of Infectious Diseases, University of Colorado Anschutz Medical Campus, Aurora, CO, USA

³Department of Microbiology, Immunology, and Pathology, Colorado State University, Fort Collins, CO, USA

*^{,4}Department of Medicine and Academic Affairs, National Jewish Health, D509, Neustadt Building, 1400 Jackson St, Denver, CO, 80206, USA

Email: ChanE@NJHealth.org

⁵Division of Pulmonary Sciences and Critical Care Medicine, University of Colorado Anschutz Medical Campus, Aurora, CO, USA

⁶University of Colorado School of Pharmacy, Aurora, CO, USA

⁷PharmcareUSA, Denver, CO, USA

⁸Department of Medicine, Denver Veterans Affairs Medical Center, Denver, CO, USA

Published online: 11 October 2016 © Springer Science+Business Media New York (outside the USA) 2016

The original version of this article was revised: The alignment of entries in Tables 5, 6, and 7, was incorrect. Table 5: Under the "Drug" column, it should be "± ciprofloxacin", under "Dose and schedule" column, it should be "250-750 mg BID", and under the "Known adverse effects" column, it should be "See moxifloxacin above" Table 6: "± Amikacin" should be moved to the right under the "Drug" column, "10-25 mg/kg IV or IM TIW for the first 2-3 months" should be moved under the "Cose and schedule" column, and "See above" should be moved under the "Known adverse effects" column. Table 7: The text for the last row is missing and is actually misplaced in the footnote: So for the last row, under the "Drug" column should be "One double-strength tablet BID", and under the "Known adverse effects" column adverse effects.

This article is part of the Topical Collection on *Mycobacterial Infections*

Keywords Antibiotic · Atypical mycobacteria · Bronchiectasis · *Mycobacterium abscessus* complex · *Mycobacterium avium* complex · Pulmonary disease

Opinion statement

Treatment of non-tuberculous mycobacterial lung disease (NTM-LD) is challenging for several reasons including the relative resistance of NTM to currently available drugs and the difficulty in tolerating prolonged treatment with multiple drugs. Yet-to-be-done, large, multicenter, prospective randomized studies to establish the best regimens will also be arduous because multiple NTM species are known to cause human lung disease, differences in virulence and response to treatment between different species and strains within a species will make unbias randomization difficult, the need to distinguish relapse from a new infection, and the difficulty in adhering to the prescribed treatment due to intolerance, toxicity, and/or drug-drug interactions, often necessitating modification of therapeutic regimens. Furthermore, the out-of-state resident status of many patients seen at the relatively few centers that care for large number of NTM-LD patients pose logistical issues in monitoring response to treatment. Thus, current treatment regimens for NTM-LD are largely based on small case series, retrospective analyses, and guidelines based on expert opinions. It has been nearly 10 years since the publication of a consensus quideline for the treatment of NTM-LD. This review is a summary of the available evidence on the treatment of the major NTM-LD until more definitive studies and guidelines become available.

Introduction

Treatment of non-tuberculous mycobacterial lung disease (NTM-LD) is challenging because different NTM with varying virulence and drug susceptibility are known to be causative agents, the relative resistance of NTM to available antibiotics, requirement of multidrug regimens for an extended period of time, frequent intolerance of the prescribed regimens, and the relatively high frequency of relapse and/or reinfection. Furthermore, expertrecommended antibiotic regimens are largely based on case series, small randomized studies, expert opinions, and anecdotal clinical experience. Herein, we review the current recommendations on treatment of NTM-LD based on the available evidence.

The prevalence of NTM lung disease and potential sources of infections

Well accepted among the clinical and scientific community is the rising rates of NTM-LD. Because of the chronicity of NTM-LD, the best measures of disease burden in a population are prevalence rates [1]. In several countries including Australia, Canada, Germany, and Taiwan, the prevalence has increased by 1.5-to 6-fold over the past decade [2–5]. The Hawaiian Islands were found to have the highest period prevalence of NTM-LD (396 cases/100,000 persons for a 10-year time period) in a sampling of 2.3 million Medicare Part B beneficiaries enrolled from 1997 to 2007 [6••]. These numbers are likely underestimated because of a lack of mandatory reporting in the USA.

There are potentially multiple reasons for the increased number of NTM-LD cases. One possibility is that the true number of cases is not increasing, but rather there are simply more cases being diagnosed due

to greater utilization of chest CT scan (prompting sputum cultures in those with abnormalities), more reliance on more sensitive molecular diagnostic techniques, and greater awareness of NTM-LD by clinicians. However, increased exposure to NTM is also a factor as evinced by a study showing that skin test reactivity to purified protein derivative-B (a mixture of antigens from Mycobacterium intracellulare) in the 1999–2000 period (~7400 subjects) was significantly greater than those tested in the 1971-1972 period (~1500 subjects) (17 vs. 11 %, respectively) [7]. Greater exposure may be related to enhanced biofilm development on polyvinyl chloride, steel, and polycarbonate surfaces—where there is greater propensity for NTM biofilm formation compared to copper and glass surfaces [8]-and subsequent inhalation of fractured biofilm carried in fine water aerosols such as that seen with high-efficiency showerheads, water misters, and hot tub bubble generators [9]. In particular, NTM glycopeptidolipids have been linked to enhanced formation of biofilms [10]. The production of extracellular polymeric substances (i.e., lipids, polysaccharides, and nucleic acids) in a three-dimensional matrix during the production of biofilms is a successful strategy used by NTM for survival in the environment [11]. Compared to planktonic bacteria, those residing in biofilms exhibit 10- to 1000-fold greater resistance to antimicrobial agents and decontaminants [8, 12]. The close proximity of NTM to each other in biofilms can also promote horizontal gene transfer (e.g., acquisition of drug resistance genes) and switch to slower growth rates that inhibit antibiotic efficiency. With increasing use of chlorination, it is possible that NTM—which are relatively resistant to chlorine except for Mycobacterium scrofulaceum which has virtually disappeared with widespread chlorination-have increased in number because of less competition from chlorine-sensitive organisms. We have also speculated that climate change with global warming and natural disasters may also contribute to increasing infection and prevalence of NTM-LD [13].

While a link has been made to water and associated biofilms as potential sources of NTM lung infections [14], a recent study found that *M. intracellulare*—a member of the *Mycobacterium avium* complex (MAC) group of organisms and historically the most common cause of NTM-LD in the USA—was not commonly found in water biofilms; instead, Mycobacterium chimaera (another MAC organism) was found in household water and biofilm samples [15•]. However, since traditional nonsequencing methods of NTM identification cannot speciate M. chimaera with any reliability, it is likely that a significant number of isolates previously classified as M. avium or M. intracellulare were in fact M. chimaera or another species under the MAC umbrella [16]. Indeed, a report from Germany showed that 143/166 (85 %) of M. intracellulare isolates initially identified by 16S rRNA gene-based methods were actually misidentified and were reclassified as M. chimaera after multilocus sequence analyses including a combination of 16S rRNA gene and 16S-23S internal transcribed spacer (ITS) region sequencing [17]. Furthermore, rpoB gene and 16S-23S ITS multilocus sequence analyses used

to speciate 448 clinical MAC isolates in the USA revealed the dominant MAC to be *M. avium* (54 %), *M. chimaera* (28 %), and *M. intracellulare* (18 %) [18••]. Thus, since a significant number of previously identified *M. intracellulare* clinical isolates may actually be another MAC species, the infrequent finding of *M. intracellulare* in water biofilms should not diminish the importance of water and their associated biofilms as sources for NTM infections.

Brief synopsis of NTM identification and diagnosis of NTM lung disease

Determining the precise NTM species and even subspecies is critical since the antibiotic regimen and prognosis varies with the responsible NTM. Newer molecular techniques that allow more precise identification include line probe hybridization and PCR. For the most precise identification of NTM species and subspecies, sequencing of *hsp65* and *rpoB* genes, as well as the 16S-23S ITS is recommended [19-23]. Furthermore, the infrequent use of NTM genotyping in the clinical setting makes it difficult to ascertain whether recurrent disease is due to a relapse or a new infection acquired from the environment.

There is no reliable biomarker that distinguishes NTM colonization vs. NTM-LD. Thus, after identification of NTM from the respiratory tract, other clinical factors must be considered to determine if the isolated NTM is clinically relevant. A guideline published jointly by the American Thoracic Society/Infectious Diseases Society of America (ATS/IDSA) espoused that, at a minimum, NTM-LD can be diagnosed when NTM-positive patients are symptomatic (e.g., cough, sputum, fatigue) and have radiographic features that are consistent with NTM-LD (e.g., nodules, bronchiectasis, tree-in-bud opacities, cavities). The NTM species can also help predict the clinical relevance; e.g., at one end of the spectrum, *M. gordonae* almost always is a laboratory contaminant and not a culprit organism whereas the presence of *Mycobacterium kansasii* is a better predictor of NTM-LD (Table 1).

Even if one has established the presence of NTM-LD, patient factors such as age, comorbid conditions, life expectancy, and goals of care (e.g., curative vs. symptom control) are important to consider since

Organisms recovered	Relative likelihood of disease
M. kansasii	High
MAC, M. abscessus complex, M. chelonae, M. malmoense, M. szulgai, M. xenopi	Intermediate depending on imaging, symptoms, and repeated isolation; <i>M. szulgai</i> in certain regions is considered to be more pathogenic
M. simiae, M. fortuitum, M. terrae	Low
M. gordonae	Very low

Table 1. Relative likelihood that NTM isolated from respiratory tract indicate disease

treatment for NTM-LD entails multiple antibiotics for an extended period of time (months to several years).

Treatment of NTM-LD is based on small clinical studies and expert opinion

Although treatment trials have been performed on patients with MAC-LD, participant numbers are generally small and often single-centered [24-28]. Large, multicenter, prospective trials reminiscent of the studies performed on tuberculosis in Hong Kong, East Africa, the USA, and several other countries have not been done with NTM-LD. Challenges to conduct such studies include the relatively few NTM-LD patients at most centers, out-of-state resident status of many patients at the few centers that care for large number of NTM-LD patients, the difficulty of distinguishing relapse and reinfection with standard laboratory techniques, and the relatively high frequency of intolerance or toxicity to various antibiotic regimens in an older population, often necessitating temporary or permanent discontinuation of certain drugs or modification of regimens based on comorbid conditions or drug-drug interactions. It has been nearly 10 years since the publication of the ATS/IDSA guidelines in the diagnosis and treatment of NTM-LD [29]. Yet, Admejian et al. recently showed that following the treatment guidelines is poor among clinicians who treat such patients [30]. We discuss below the treatment of NTM-LD caused by MAC, M. kansasii, Mycobacterium malmoense, Mycobacterium xenopi, Mycobacterium szulgai, Mycobacterium simiae, and Mycobacterium abscessus complex.

Treatment for MAC

MAC has traditionally been composed of several well-known species, including *M. avium* and *M. intracellulare* but more recently, also include a third group known as MAC "X," composed of at least eight less common MAC species [15•]. In the USA, standard laboratory testing is completed by partial sequencing and subsequent phylogenetic analysis of the *16S rRNA* gene, and thus, the delineation of different strains of MAC is frequently unavailable. Currently, all species within MAC are similarly treated according to standard ATS/IDSA guidelines [29]. But clearly, different "MAC strains" are associated with differential virulence [31]; if future analyses find the virulence of MAC species significantly differs, distinguishing these MAC species may ultimately prove useful for prognosticating and perhaps guiding therapy [32, 33].

In contrast to treatment for tuberculosis, where in vitro susceptibility testing is a useful guide for effective in vivo response to antibiotics [29], the Clinical Laboratory Standards Institute and ATS/IDSA guidelines recommend testing MAC susceptibility only to clarithromycin (or azithromycin) [29]. This recommendation is based on studies showing a relationship between clinical efficacy and minimum inhibitory concentration (MIC) for clarithromycin, but not for rifampicin, ethambutol, or streptomycin [34–38]. No studies have established superiority of one macrolide over another [28]. Combined rifampin and ethambutol drug susceptibility testing has shown mutual lowering of the individual drug MIC, but the clinical importance of this in vitro synergy is controversial [39, 40]. Susceptibility to amikacin ($\leq 16 \ \mu g/mL$ for susceptible, 32 $\mu g/mL$ for intermediate, and $\geq 64 \ \mu g/mL$ for resistant) may also help predict treatment success [41].

Two major radiographic phenotypes of NTM-LD are seen—nodularbronchiectasis often involving the right middle lobe and lingula segment and the fibrocavitary form often involving the upper lobes—although patients may have features of both. For patients with nodularbronchiectasis MAC-LD without cavitation, it is acceptable to treat with a macrolide, rifamycin, and ethambutol—the standard three-drug regimen—thrice weekly (Table 2) [29, 42–44]. In a study that compared such dosing in 180 MAC-LD patients with nodular-bronchiectasis who

Regimen	Drug	Dose and schedule ^a	Known adverse effects
Fibrocavitary disease: daily three-drug regimen	Azithromycin <i>or</i> clarithromycin <i>and</i>	250 mg QD 1000 mg QD	Diarrhea, hearing loss, metallic taste, QT prolongation
± thrice weekly aminogly- coside for at least 12 months of negative sputum cultures.	Rifampin <i>or</i> rifabutin ^b and	10 mg/kg QD (maximum 600 mg) <i>or</i> 150–300 mg QD	Hepatitis, drug-drug interactions, fever, chills, arthralgias, thrombocytopenia, leukopenia, acute hemolytic anemia, uveitis, decreases clarithromycin levels
	Ethambutol	15 mg/kg QD (max 2.4 g)	Optic neuritis (color blindness, scotoma, decreased visual acuity and/or visual defect), hepatitis
	± Amikacin or streptomycin	10–25 mg/kg ^c IV or IM TIW for the first 2–3 months	Renal failure, cranial VII (vestibular and cochlear) toxicity, hypomagnesemia
	Clofazamine (for intolerance to rifamycins)	100-300 mg QD	GI intolerance (abdominal pain, nausea, vomiting, diarrhea – occurring in 40-50 %), skin discoloration, QT prolongation
Nodular-bronchiectasis: thrice weekly for at least	Azithromycin <i>or</i> clarithromycin <i>and</i>	500 mg TIW 1000 mg TIW	See above
12 months of negative sputum cultures.	Rifampin <i>or</i> rifabutin ^a and	600 mg TIW 150–300 mg TIW	See above
	Ethambutol	25 mg/kg TIW	See above

Table 2. Treatment of Mycobacterium avium complex (MAC)

Key references: [29, 45••, 93]

IV intravenous, IM intramuscular, QD once daily, TIW thrice weekly

^aUnless indicated, all dosages are given orally

^bReserve rifabutin for advanced or previously treated disease. Rifabutin levels may be increased by macrolides. Hematologic complications is more common with rifabutin than rifampin

^cSome centers favor lower doses for better tolerability. Monitor peak aminoglycoside level weekly

completed >12 months of multidrug therapy, thrice weekly dosing was as good as daily dosing with combined treatment success rate of 84 % (sputum conversion with no evidence of microbiologic relapse) [28]. Interestingly, most of the recurrences (75 %) were due to new infections rather than a true relapse (25 %) [28]. Overall, the standard three-drug regimen of a macrolide, rifamycin, and ethambutol gives a durable culture conversion rate of ~60–80 % [25, 28, 34, 35, 37, 45••].

For those with fibrocavitary disease, daily dosing rather than intermittent therapy with the standard three-drug regimen is recommended (Table 2) [43]. With cavitary or severe nodular-bronchiectatic disease, the addition of an aminoglycoside during the first 2–3 months of therapy is recommended based on a study showing that addition of streptomycin to the standard three-drug regimen significantly improved the sputum conversion rates, albeit long-term outcome was not significantly better [27]. There are small case series describing inhaled amikacin to treat MAC-LD with varying success [46–48]. An ongoing multicenter study is recruiting patients to determine if inhaled liposomal amikacin helps for MAC-LD recalcitrant to the standard three-drug regimen (http://www.insmed.com/clinical-trials/).

Commensurate with the importance of a macrolide in the treatment of MAC-LD, the development of macrolide resistance in a MAC isolate is strongly associated with treatment failure and increased mortality [49]. Factors associated with emergence of macrolide resistance include macrolide monotherapy and dual therapy with a macrolide and fluoroquinolone [49]. Thus, avoiding these practices is a critical element in the management of patients with MAC-LD. However, once macrolide resistance develops, aggressive therapy, usually including use of an injectable aminoglycoside and possibly lung resection should be considered [49].

Second-line agents for MAC are generally reserved for those with disease recalcitrant to first-line treatment, intolerance, or unacceptable adverse effects to one or more of the first-line agents, and/or macrolide resistance. For patients intolerant to rifamycins, substitution with clofazimine is a viable option with prolonged sputum conversion rates for clofazimine, macrolide, and ethambutol or minocycline to be ~ 65 %, similar to that seen with rifampin-based three-drug therapy [25, 45••, 50, 51•, 52, 53]. Clofazimine synergizes with clarithromycin or amikacin against MAC [50, 52]. The use of linezolid was assessed in a retrospective multicenter study across six NTM centers in over 100 patients, 33 % of whom had MAC-LD [54]. Many of these patients had stable or improved disease on treatment, but significant side effects occurred in about half. Bedaquiline, approved by the FDA in 2013 for the treatment of multidrug-resistant tuberculosis, appears to be welltolerated in patients with refractory MAC-LD [55]. A UK study compared 24 months of treatment for MAC-LD with either ciprofloxacinethambutol-rifamycin or macrolide-ethambutol-rifamycin and found similar and very low cure rates for both arms (23-24 %), but this may be due to the fact that two thirds had cavitary disease [26], a sign of more severe disease. While this finding suggests that ciprofloxacin is equivalent to clarithromycin in efficacy, experts in North America and several other regions still consider clarithromycin or azithromycin the

most important drug available for MAC. Moxifloxacin has been used with some success in MAC-LD patients who failed clarithromycin-based therapy [56]. Thus, for refractory or resistant MAC-LD, there is little published data for treatment options apart from ATS/IDSA guidelines.

Treatment for M. kansasii

M. kansasii is one of the more common causes of NTM-LD in the Western hemisphere but is less common in Asia [57]. M. kansasii is phylogenetically the most closely related NTM species to M. tuberculosis [58]. It is considered the most virulent NTM and classically causes upper lobe fibrocavitary lung disease similar to tuberculosis. Historically, isolation of M. kansasii has been considered to almost always predict true NTM-LD. However, a recent review of 19 papers published over the course of 60 years found that 1008 of 2672 patients (38%) evaluated after isolation of M. kansasii isolation did not meet clinical criteria for disease [59].

There are no randomized trials comparing treatment regimens for M. kansasii. Fortunately, M. kansasii is among the most antibiotic-responsive NTM species with a low rate of treatment failure or relapse (<1 %) among patients who completed the ATS/IDSA recommended treatment of daily isoniazid, rifampin, and ethambutol until cultures are negative for at least 12 months (Table 3) [29]. Rifampin is the most important agent in this regimen as cure was uncommon in the pre-rifampin era [60].

M. kansasii is also typically susceptible to macrolides, fluoroquinolones, and the aminoglycosides but is intrinsically resistant to pyrazinamide. Susceptibility testing should be performed for rifampin and clarithromycin [61]. Because isoniazid activity is lower for M. kansasii than for M. tuberculosis and macrolides have strong in vitro activity, an alternative regimen is replacing isoniazid with a macrolide. Indeed, two studies used a regimen of clarithromycin, rifampin, and ethambutol daily or thrice weekly for at least 12 months of culture negativity while on treatment and identified no relapses after a long period of follow-up [62, 63]. Patients receiving shorter courses (9 to 12 months) with different

Regimen	Drug	Dose and schedule	Known adverse effects
Daily for 12 months of negative cultures	Isoniazid ^a and	300 mg QD	Hepatotoxicity, peripheral neuropathy drug-induced lupus, agranulocytosis
	Rifampin and	600 mg QD	See above
	Ethambutol	15 mg/kg QD (max 2.4 g)	See above
	Azithromycin or clarithromycin may be substituted for isoniazid	See above	See above

^aConsider adding pyridoxime 20 mg QD to lower risk of INH-induced peripheral neuropathy

regimens have experienced unacceptable rates of relapse (6-10 %) [64, 65]. Earlier regimens also included streptomycin in the first 3 months [60, 66], but since high cure rates are typically achieved with oral regimens, aminoglycosides are rarely necessary. For rifampin-resistant M. kansasii, ATS/IDSA recommends a combination of clarithromycin, moxifloxacin, and a third agent with in vitro susceptibility such as ethambutol or sulfamethoxazole [29].

Treatment for *M. malmoense*

M. malmoense is an uncommon pathogen in the USA but is a more common cause of NTM-LD in Europe. It grows more slowly than MAC in liquid media and historically, it is a difficult species to treat. Similar to MAC, many clinicians cannot correlate in vitro susceptibilities with an in vivo response. A prospective study of 106 patients with M. malmoense-LD was performed over a 5-year period by the British Thoracic Society (BTS) [67]. The results of 2 years of treatment with rifampin plus ethambutol were equivalent to rifampin, ethambutol plus isoniazid, although only 53 % of patients were alive at 5 years and 44 of the original 106 patients (42 %) were cured of the infection [67]. In a follow-up study, the BTS randomly assigned 167 patients with M. malmoense-LD to clarithromycin, rifampin, and ethambutol, or ciprofloxacin, rifampin, and ethambutol. Overall response rates were low, but the group receiving clarithromycin had slightly better clinical response and lower mortality [26]. There have been several other retrospective studies with varying success to rifampicin and ethambutol with or without a macrolide [4500, 68, 69]. Overall, the optimum antibiotic regimen to treat M. malmoense remains unknown but based on available evidence, standard MAC therapy is a reasonable place to start (Table 4).

Treatment of M. szulgai

M. szulgai, an organism closely related to M. malmoense [70], is one of a few NTM that possess the ESAT-6 and CFP-10—small secretory proteins produced by M. tuberculosis but only by a few NTM-indicating that it may contain

Regimen	Drug	Dose and schedule	Known adverse effects
Daily therapy for at least 12 months of negative sputum cultures	Clarithromycin ^a or azithromycin <i>and</i>	1000 mg QD or 250 mg QD	See above
	Rifampin <i>and</i>	600 mg QD	See above
	Ethambutol	15 mg/kg QD	See above
	Moxifloxacin in place of the macrolide	400 mg QD	Tendonitis, QT prolongation photosensitivity

^aFluoroquinolones may be used as an alternative in those intolerant to macrolides

virulence genes similar to *M. tuberculosis* [71]. Consistent with this hypothesis is that isolation of *M. szulgai* from a patient generally indicates actual disease [72]. *M. szulgai* is known to cause chronic lung disease and skin and soft tissue infections. Treatment regimens that consist of clarithromycin, rifampin, and ethambutol (± ciprofloxacin) have been used successfully (Table 5) [72, 73].

Treatment of M. xenopi

Treatment of Muschasterium and and

. . . .

M. xenopi is named for its isolation from the toad *Xenopus laevis* [74]. It is mainly isolated from water sources [29] and clinically more relevant in certain parts of the world including Europe [75•, 76, 77].

M. xenopi is often recalcitrant to treatment, with correspondingly lower sputum conversion rates, reduced long-term cure rates, and associated with higher mortality [24, 26, 45••, 77, 78]. There is resistance to typical anti-tuberculosis drugs such as rifampin, isoniazid, and ethambutol [79]; however, the rifabutin MIC_{50} for a small number of strains tested was <0.5 [80]. In time-kill kinetic studies, moxifloxacin and clarithromycin are equally effective [81]. In vitro activity with ciprofloxacin and amikacin was limited [82], whereas no in vitro synergy between rifampin and ethambutol was seen [39]. Andrejak and co-workers performed a comprehensive study of antimicrobials using drugs in vitro, sera of treated mice to assess antimicrobial activities ex vivo, and in vivo murine models to test drug activities [83••]. They found that in vitro, two-drug combinations of ethambutol plus either rifamycin or moxifloxacin, and of clarithromycin plus moxifloxacin showed the best bactericidal activities; ex vivo, a three-drug combination of ethambutol plus a rifamycin and either clarithromycin or moxifloxacin was best [83••]. Interestingly, for the in vivo studies, amikacin-containing regimen had the greatest bactericidal activity with no difference in regimens containing clarithromycin or moxifloxacin [83••]. Based on a few trials, isoniazid is not effective and may in fact be associated with worse outcomes [24, 84]. Short of more robust studies, the most efficacious drugs are the rifamycins, macrolides, and the fluoroquinolones [26, 29, 45., 77, 85]. Thus, based on available evidence, it is recommended that M. xenopi-LD be treated with a combination of a macrolide, rifamycin, ethambutol \pm a fluoroquinolone (Table 6) [26, 45••].

able 5. Treatment of <i>Mycobacterium szulga</i>			
Regimen	Drug	Dose and schedule	Known adverse effects
No proven regimen. Daily therapy	Clarithromycin and	1000 mg QD	See above
for >12 months of negative cultures.	Rifampin and	10 mg/kg QD (max 600 mg)	See above
	Ethambutol and	15 mg/kg QD (max 2.4 g)	See above
	± Ciprofloxacin	250-750 mg BID	See moxifloxacin above
Key references: [72, 73]			

Regimen	Drug	Dose and schedule	Known adverse effects
No proven regimen. Daily therapy for >12 months of negative cultures	Clarithromycin <i>and</i> Rifampin <i>and</i> Ethambutol ± Moxifloxacin ± Amikacin ^a	1000 mg QD 10 mg/kg QD (max 600 mg) 15 mg/kg QD (max 2.4 g) 400 mg QD 10–25 mg/kg IV or IM TIW for the first 2–3 months	See above See above See above See above See above
Key references: [26, 45••] ^a Based on an in vivo murine study [83••]			

Table 6. Treatment of Mycobacterium xenopi

Treatment of *M. simiae*

M. simiae is isolated from water sources in the environment [29, 70] and is capable of, but rarely causes disease, i.e., a positive culture is a low predictor of having actual NTM-LD [86, 87]. When implicated in pulmonary disease, its clinical presentation has similarities to that seen with MAC-LD [88]. This organism is resistant to many of the standard tuberculosis drugs such as isoniazid, rifamycin, ethambutol, and para-aminosalicylic acid [80, 89]. In one report, 27 of 86 isolates (31 %) were susceptible to streptomycin [90]. Agents that have in vitro activity include ofloxacin, amikacin, clarithromycin, ethionamide, cycloserine, and clofazimine [91]. Like M. xenopi, no in vitro synergy was seen between rifampin and ethambutol [39]; however, synergy could be demonstrated between amikacin and clofazimine [52]. Jeong et al. recently reported the first confirmed South Korean patient with M. simiae-LD who, despite receiving 12 month treatment with azithromycin, rifampin, ethambutol, and moxifloxacin, failed to achieve culture conversion [92]. ATS/IDSA guidelines primarily suggest a clarithromycin-based, multiple drug regimen, citing that drugs such as moxifloxacin, sulfamethoxazole/trimethoprim, or linezolid have activity against M. simiae (Table 7) [29]. For cavitary disease, extrapolation for MAC treatment suggests amikacin for a period of time could be helpful. Overall, the results of therapy are often disappointing and treatment for *M. simiae* remains problematic [93].

Treatment for M. abscessus complex and M. chelonae

Among the rapidly growing mycobacteria (RGM)—defined historically as observable growth from a subculture on solid medium in <7 days—those belonging to the *M. abscessus* complex are the most clinically relevant to humans. Whole genome sequencing support categorizing *M. abscessus* complex into three distinct subspecies: *M. abscessus* sensu stricto, *M. bolletii*, and *M. massiliense* [94].

Regimen	Drug	Dose and schedule	Known adverse effects
No proven regimen. Relationship of in vitro susceptibility and clinical outcome is not clear.	Clarithromycin and Rifampin and Ethambutol ± Moxifloxacin ± Trimethoprim- sulfamethoxazole	1000 mg QD 10 mg/kg QD (max 600 mg) 15 mg/kg QD (max 2.4 g) 400 mg QD One double-strength tablet BID	See above See above See above See above Hypersensitivity, rash, myelosuppression, interstitial nephritis, increased liver function tests
Key reference: [29]			

In the clinical laboratory, distinguishing these three *M. abscessus* complex subspecies can be difficult and often requires sequencing of *hsp65*, *rpoB*, *secA*, erythromycin ribosomal methylase 41 (*erm41*), and/or *16S–23S rRNA ITS*.

Pang and coworkers performed in vitro susceptibility testing of 40 international reference RGM to 20 antimicrobial agents and found that amikacin, tigecycline, and linezolid had potent activities and that the fluoroquinolones, cefoxitin, and meropenem had good activities [95]. Although the Clinical and Laboratory Standards Institute recommends drug susceptibility testing to ten antimicrobial agents (amikacin, cefoxitin, clarithromycin, ciprofloxacin, doxycycline or minocycline, imipenem, linezolid, moxifloxacin, trimethoprim-sulfamothoxazole, and tobramycin) for M. abscessus complex isolates [96, 97] and recommended by some experts [94, 98], others cite insufficient evidence that in vitro susceptibility correspond with in vivo response with the exception of the macrolides [99]. The method of drug susceptibility testing (e.g., disk diffusion vs. broth dilution methods with the latter method preferred) may account for differences in drug susceptibility and hence, differences in reliability of the results in predicting outcome [96]. One study from South Korea noted sputum conversion and maintenance of negative sputum cultures for more than 12 months were significantly lower in patients whose isolates were resistant to clarithromycin (2/12, 17 %) compared with those whose isolates were susceptible or intermediate to clarithromycin (21/33, 64 %) [100].

A distinguishing feature of *M. abscessus* sensu stricto and *M. bolletii* from *M. massiliense* is that the first two RGM have poor treatment outcomes. The reason for this is the presence of the functional *erm41* gene in both *M. abscessus* sensu stricto and *M. bolletii*, which, upon exposure to macrolides, modifies the binding site for macrolides and induces resistance [101]. Clarithromycin may have greater propensity to induce greater erm41 expression resulting in greater macrolide resistance than azithromycin in *M. abscessus* sensu stricto infection [102] although others have not found a difference in the two macrolides to induce such resistance [103].

Additionally, *M. abscessus* sensu stricto possesses other enzymes and efflux pumps that may confer in vivo antibiotic resistance [36].

Koh et al. summarized the four studies (three from South Korea and one from Denver, Colorado) examining the treatment outcome for *M. abscessus*-LD [94]. In the three studies that did not distinguish the subspecies of *M. abscessus* complex, the sputum conversion rate without evidence of relapse was 50–70 %. In contrast, in the one study from South Korea where *M. massiliense* was distinguished from *M. abscessus* sensu stricto, the sputum conversion rate was significantly more favorable for *M. massiliense* (88 %) than for *M. abscessus* sensu stricto (25 %) [104••]. This finding indicates that the presence of the *erm41* gene may be the reason for the generally poor response of *M. abscessus* sensu stricto to medical therapy as oppose to the more favorable outcomes in the treatment of lung disease due to *M. massiliense* [104••, 105, 106]. Thus, this is a good example where distinguishing the subspecies has important prognostic implications for patients.

The only oral drugs with reliable activity against *M. abscessus* complex organisms are the macrolides and clofazimine, although others that have been used in treatment include ciprofloxacin, moxifloxacin, linezolid, or doxycycline [107]. Similar to that for MAC, clofazimine was found to synergize with clarithromycin or amikacin against *M. abscessus* [50]. Parenteral drugs used in the treatment of *M. abscessus* complex include cefoxitin, imipenem, tigecycline, and amikacin.

Current guidelines recommend an initial intensive phase of daily clarithromycin or azithromycin and two parenteral drugs for 2–4 months (e.g., amikacin thrice weekly plus daily imipenem or cefoxitin in divided doses) followed by macrolide therapy with at least one other oral agent (fluoroquinolone, linezolid, clofazimine) or inhaled amikacin [45••, 108, 109]. If the isolate is identified as *M. abscessus* sensu stricto or *M. bolletii*, then the macrolide should not be used and inhaled amikacin plus clofazimine should be considered in the continuation phase (Table 8). The total duration of therapy is based on culture conversion, typically treating for ≥ 12 months of negative sputum cultures. On the other hand, if the goal is symptom control and not intent to cure, then the continuation phase can be shortened.

In contrast to *M. abscessus* sensu stricto, *M. chelonae*—a much less common cause of lung disease—does not have an active *erm41* gene and thus should not have inducible macrolide resistance.

Role of resectional lung surgery

Surgical resection can play an important adjunctive role in the management of NTM-LD [29, 110]. However, there is a paucity of robust evidence to guide the calculation of risks and benefits of selecting NTM-LD patients for lung resection. A number of single-center, retrospective case series have reported long-term microbiological success after lung resection for NTM-LD [111–119]. Case series of *M. abscessus* patients have suggested improved outcomes with surgical resection relative to antibiotic treatment alone [100, 108, 120]. Nonetheless, several caveats are critical in interpretation of these data. First, these reports come from

Regimen	Drug	Dose and schedule	Known adverse effects
M. abscessus sensu stricto or M. a	<i>bscessus</i> subsp. <i>bolletii</i>		
A reasonable regimen includes clofazimine, cefoxitin, and amikacin initially, then clofazimine, inhaled amikacin, and ± linezolid for >12 months of negative culture. If the goal	Clofazimine and	100-300 mg QD	See above
	Cefoxitin or	2 g IV BID or TID	Thrombophlebitis, induration, hypotension, and rash
	Imipenem and	500–1000 mg QID IV (BID also used)	Seizures, bone marrow suppression, increased liver function tests
is symptom control, length of treatment can be shortened.	Amikacin	15–25 mg/kg IV TIW for the first 2–3 months	See above. <i>M. abscessus</i> can acquire resistance due to mutation of the 16S rRNA gene
	± Tigecycline	50 mg IV QD or BID	Hepatotoxicity, pancreatitis, photosensitivity, tooth discoloration, and anorexia
	± Linezolid	600 mg PO BID	Serotonin syndrome, myelosuppression, neuropathy, metabolic acidosis
M. abscessus subsp. massiliense			
Daily regimen of macrolide + cefoxitin or imipenem ±	Azithromycin or clarithromycin <i>and</i>	250 mg QD 1000 mg QD	See above
thrice weekly amikacin for at least 12 months of negative	Cefoxitin or	2 g BID or TID IV for ≥2 months	See above
culture.	Imipenem	500–1000 mg QID IV (BID also used)	See above
	Amikacin	15–25 mg/kg IV TIW for the first 2–3 months	See above
	± Moxifloxacin	400 mg QD	See above

Table 8. Treatment of Mycobacterium abscessus complex

centers with extensive experience with surgical techniques specific to infectious lung disease. Results should not be extrapolated to centers with less experience in these highly specialized techniques. Second, patients seen in these centers are highly selected and may not be representative of NTM patients generally.

Expert opinion emphasizes consideration of surgical resection under the following circumstances: (i) for pathogens that are less amenable to medical therapy such as *M. abscessus* or macrolide-resistant MAC organisms, (ii) for severely involved but localized disease (e.g., focal bronchiectasis or cavitation), and/or (iii) for patients who have responded poorly to initial medical treatment and for life-threatening complications such as hemoptysis [29, 110]. Patients are typically treated with intensive antibiotic regimens for 2–3 months prior to surgery to maximally reduce bacterial burden [110]. ATS/IDSA guidelines conclude that—in light of the lack of widely-accepted criteria for patient selection and the high potential morbidity—decision-making regarding surgical resection should optimally be made in conjunction with experienced NTM treatment centers [29].

Adjunctive treatment

In addition to antimicrobial therapy, avoidance of potential sources of exposure (e.g., aerosolized water, soil, and biofilms) is necessary to prevent reinfection. In patients who are on inhaled corticosteroids, a trial to limit their use may potentially help with treatment as inhaled corticosteroids have been shown to be a risk factor for acquiring NTM-LD and by inference, presumably may hinder response to therapy [121]. Moreover, measures to improve airway clearance in the bronchiectatic airways are essential adjunct to drug treatment—e.g., airway clearance devices (Acapella®, Aerobika®, TheraVest®), agents to enhance expectoration of mucous (e.g., inhaled hypertonic saline or mannitol), and optimization of nutrition. Other measures to maximize favorable treatment outcome are listed in Table 9.

Traditional and more novel predictors of treatment outcome

Traditional factors that may contribute to suboptimal outcome include innate and acquired resistance including to the endogenous antimicrobial peptide cathelicidin [36, 122], under-dosing of antimicrobials [123], the presence of other pathogens in the respiratory tract (NTM or non-NTM bacteria or fungi), and newly acquired NTM infections during treatment or after completion.

In 72 Japanese patients with MAC-LD on triple drug regimen, in whom 51 % experienced treatment success-defined as sputum

Measures	Key references
Test NTM for resistance to macrolide and possibly the aminoglycosides	[27, 34, 35, 41]
Use at least a three-drug regimen of azithromycin, rifampin, and ethambutol for MAC-LD, thrice weekly for non-cavitary minimal disease and daily for cavitary and more severe disease	[29, 42–44]
If rifampin is used with macrolides, the dose of macrolides may need to be increased and preferably, macrolide levels checked	[34, 45●●, 123]
Consider including intravenous or inhaled amikacin for those with more severe or cavitary disease	[27, 47, 48]
Take measures to avoid re-exposure such as avoidance of aerosolized soil and water	[124]
Consider increasing hot water heater temperature to 130 °F	[127, 128]
Airway clearance mechanisms for bronchiectasis (e.g., Acapella® valve, Aerobika®, TheraVest®, hypertonic saline)	[129]
Surgical lung resection for severe but localized disease in selected patients	[29, 110]

Table 9. Measures to maximize favorable treatment outcome

conversion without relapse—low soil exposure (defined as ≤ 2 h of soilrelated activities per week) was associated with significantly greater sputum conversion, lower rates of relapse, and higher treatment success rates than those with high soil exposure (>2 h per week) [124]. This finding is consistent with the long-standing observation with now experimental proof that a substantial number of recurrences are due to a new infection rather than a true relapse [28].

While treatment outcome is most often discussed in the context of the antibiotic regimen used and perhaps drug susceptibility, some investigators have examined the association between outcome and genotype of the organisms. Kikuchi and coworkers categorized patients with M. avium-LD as having a therapeutic responsiveness (defined as microbiologic and radiologic improvement) or unresponsiveness and then retrospectively determined whether there was an association between the *M. avium* genotype of the variable number tandem repeats (VNTR) at 16 minisatellite loci and responsiveness to therapy [125]. Performing principal component analysis of the raw VNTR data, they were able to identify genetic features that were associated with therapeutic response to clarithromycin-containing regimen. Subsequently, they constructed a multivariate model to predict therapeutic responsiveness using VNTR data from only four minisatellite loci [125]. This study would lend credence to the notion that different strains of NTM-even within the same species—may have differential virulence. In contrast, Kim S-Y et al. found no association between the clinical characteristics, drug susceptibility, disease progression, and M. intracellulare cluster based on VNTR genotyping [126].

Conclusions and future developments

In conclusion, we summarized the treatment of NTM-LD based on available data with the caveat that many of the recommendations are based on clinical experience, small clinical studies, and expert opinions. Systematic, multicenter studies are needed to provide more robust evidence-based recommendations for treatment and outcome analysis of NTM-LD.

Compliance with ethical standards

Conflict of interest

Dr. Julie V. Philley, Dr. Jennifer R. Honda, Dr. Michael M. Chan, Dr. Shannon Kasperbauer, Dr. Nicholas D. Walter, and Dr. Edward D. Chan declare that they have no conflict of interest. Dr. Mary Ann DeGroote is a co-PI for the preclinical evaluation of new therapeutic entities for NTM therapeutics with Crestonepharma Inc. This is an SBIR phase II grant. There is no overlap with antimicrobial agents described in this review.

Human and animal rights and informed consent

This article does not contain any studies with human or animal subjects performed by any of the authors.

References and Recommended Reading

Papers of particular interest, published recently, have been highlighted as:

- Of importance
- •• Of major importance
- 1. Prevots DR, Marras TK. Epidemiology of human pulmonary infection with nontuberculous mycobacteria: a review. Clin Chest Med. 2015;36:13.
- 2. Lai CC, Tan CK, Chou CH, Hsu HL, Liao CH, Huang YT, et al. Increasing incidence of nontuberculous mycobacteria, Taiwan, 2000-2008. Emerg Infect Dis. 2010;16:294.
- Marras TK, Mendelson D, Marchand-Austin A, May K, Jamieson FB. Pulmonary nontuberculous mycobacterial disease, Ontario, Canada, 1998-2010. Emerg Infect Dis 2013; 19: 1889.
- Ringshausen FC, Wagner D, de Roux A, Diel R, Hohmann D, Hickstein L, et al. Prevalence of nontuberculous mycobacterial pulmonary disease, Germany, 2009-2014. Emerg Infect Dis. 2016;22:1102.
- Thomson RM. NTM working group at Queensland TB Control Centre and Queensland Mycobacterial Reference Laboratory. Changing epidemiology of pulmonary nontuberculous mycobacteria infections. Emerg Infect Dis. 2010;16:1576.
- 6.•• Adjemian J, Olivier KN, Seitz AE, Holland SM, Prevots DR. Prevalence of nontuberculous mycobacterial lung disease in U.S. Medicare beneficiaries. Am J Respir Crit Care Med. 2012;185:881.

A comprehensive study examining the prevalence of nontuberculous mycobacterial (NTM) lung disease in Medicare Part B Beneficiaries in the United States. It found that in this elderly population, the prevalence increased by 8.2% per year from 1997 to 2007, with significant racial/ethnic and geographic differences. Hawaii had the highest period prevalence among the 50 states at 396 cases of NTM lung disease per 100,000.

- Khan K, Wang J, Marras TK. Nontuberculous mycobacterial sensitization in the United States: national trends over three decades. Am J Respir Crit Care Dis. 2007;176:306.
- Williams MM, Yakrus MA, Arduino MJ, Cooksey RC, Crane CB, Banerjee SN, et al. Structural analysis of biofilm formation by rapidly and slowly growing nontuberculous mycobacteria. Appl Environ Microbiol. 2009;75:2091.
- Mullis SN, Falkinham JO 3rd. Adherence and biofilm formation of *Mycobacterium avium*, *Mycobacterium intracellulare* and *Mycobacterium abscessus* to household plumbing materials. J Appl Microbiol. 2013;115:908.
- Recht J, Kolter R. Glycopeptidolipid acetylation affects sliding motility and biofilm formation in *Mycobacterium smegmatis*. J Bacteriol. 2001;183:5718.
- Sousa S, Bandeira M, Carvalho PA, Duarte A, Jordao L. Nontuberculous mycobacteria pathogenesis and biofilm assembly. Int J Mycobacteriol. 2015;4:36.

- Simões M. Antimicrobial strategies effective against infectious bacterial biofilms. Curr Med Chem. 2011;18:2129.
- 13. Honda JR, Bernhard JN, Chan ED. Natural disasters and nontuberculous mycobacteria: a recipe for increased disease? Chest. 2015;147:304.
- 14. Falkinham JO 3rd, Iseman MD, de Haas P, van Soolingen D. *Mycobacterium avium* in a shower linked to pulmonary disease. J Water Health. 2008;6:209.
- 15.• Wallace RJ Jr, Iakhiaeva E, Williams MD, Brown-Elliott BA, Vasireddy S, Vasireddy R, et al. Absence of *Myco-bacterium intracellulare* and presence of *Mycobacterium chimaera* in household water and biofilm samples of patients in the United States with *Mycobacterium avium* complex respiratory disease. J Clin Microbiol. 2013;51:1747.

This study sequenced specific genes of NTM isolated from water sources in households of patients with *Mycobacterium avium* complex (MAC) lung disease. While most of the clinical respiratory isolates (91 %) were *M. intracellulare*, unexpectedly most of the isolates (70 %) from household water samples were *M. chimaera;* in fact, no *M. intracellulare* were identified from the water sources. While this study does not necessarily imply that water or water-associated biofilms are not important sources of NTM infection, it does demonstrate that proper NTM species identification is important and that different MAC species occupydifferent environmental niches.

- Tortoli E, Rindi L, Garcia MJ, Chiaradonna P, Dei R, Garzelli C, et al. Proposal to elevate the genetic variant MAC-A, included in the *Mycobacterium avium* Complex, to species rank as *Mycobacterium chimaera* sp. nov. Int J Syst Evol Microbiol. 2004;54(Pt 4):1277.
- 17. Schweickert B, Goldenberg O, Richter E, Göbel UB, Petrich A, Buchholz P, et al. Occurrence and clinical relevance of *Mycobacterium chimaera* sp. nov., Germany. Emerg Infect Dis. 2008;14:1443.
- 18. •• Boyle DP, Zembower TR, Reddy S, Qi C. Comparison of clinical features, virulence, and relapse among *Mycobacterium avium* complex species. Am J Respir Crit Care Med. 2015;191:1310.

With availability of gene sequencing, increasing number of distinct species are now identified under the "*Mycobacterium avium* complex (MAC)" umbrella. Using a clinical consensus definition of the presence of NTM lung disease, this study suggests (but does not necessarily prove) that the three major MAC species(*M. avium, M. intracellulare,* and *M. chimaera*) are likely to have differences in virulence.

19. Frothingham R, Wilson KH. Sequence-based differentiation of strains in the Mycobacterium avium complex. J Bacteriol. 1993;175:2818.

- 20. Ben Salah I, Adékambi T, Raoult D, Drancourt M. rpoB sequence-based identification of *Mycobacterium avium* complex species. Microbiology. 2008;154:3715.
- 21. Zelazny AM, Root JM, Shea YR, Colombo RE, Shamputa IC, Stock F, et al. Cohort study of molecular identification and typing of *Mycobacterium abscessus*, *Mycobacterium massiliense*, and *Mycobacterium bolletii*. J Clin Microbiol 2009; 47: 1985.
- 22. Macheras E, Roux AL, Bastian S, Leão SC, Palaci M, Sivadon-Tardy V, et al. Multilocus sequence analysis and rpoB sequencing of *Mycobacterium abscessus* (sensu lato) strains. J Clin Microbiol. 2011;49:491.
- 23. Jang MA, Koh WJ, Huh HJ, Kim SY, Jeon K, Ki CS, et al. Distribution of nontuberculous mycobacteria by multigene sequence-based typing and clinical significance of isolated strains. J Clin Microbiol. 2014;52:1207.
- 24. Research Committee of the British Thoracic Society. First randomised trial for pulmonary disease caused by *M. avium intracellulare, M. malmoense* and *M. xenopi* in HIV negative patients: rifampicin, ethambutol and isoniazid versus rifampicin and ethambutol. Thorax. 2001;56:167.
- 25. Field SK, Cowie RL. Treatment of *Mycobacterium aviumintracellulare* Complex lung disease with a macrolide, ethambutol, and clofazimine. Chest. 2003;124:1482.
- Jenkins PA, Campbell IA, Banks J, Gelder CM, Prescott RJ, Smith AP. Clarithromycin vs ciprofloxacin as adjuncts to rifampicin and ethambutol in treating opportunist mycobacterial lung diseases and an assessment of *Mycobacterium vaccae* immunotherapy. Thorax. 2008;63:627.
- Kobashi Y, Matsushima T, Oka MA. Double-blind randomized study of aminoglycoside infusion with combined therapy for pulmonary *Mycobacterium avium* complex disease. Respir Med. 2007;101:130.
- Wallace RJ Jr, Brown-Elliott BA, McNulty S, Philley JV, Killingley J, Wilson RW, et al. Macrolide/Azalide therapy for nodular/bronchiectatic mycobacterium avium complex lung disease. Chest. 2014;146:276.
- Griffith DE, Aksamit T, Brown-Elliott BA, Catanzaro A, Daley C, Gordin F, et al. An official ATS/IDSA statement: diagnosis, treatment, and prevention of nontuberculous mycobacterial diseases. Am J Respir Crit Care Med. 2007;175:367.
- Adjemian J, Prevots DR, Gallagher J, Heap K, Gupta R, Griffith D. Lack of adherence to evidence-based treatment guidelines for nontuberculous mycobacterial lung disease. Ann Am Thorac Soc. 2014;11:9.
- 31. Tateishi Y, Hirayama Y, Ozeki Y, Nishiuchi Y, Yoshimura M, Kang J, et al. Virulence of *Mycobacterium avium* complex strains isolated from immunocompetent patients. Microb Pathog. 2009;46:6.
- 32. Stout JE, Hopkins GW, McDonald JR, Quinn A, Hamilton CD, Reller LB, et al. Association between 16S–23S internal transcribed spacer sequence groups of *Mycobacterium avium* complex and pulmonary disease. J Clin Microbiol. 2008; 46.
- 33. Han XY, Tarrand JJ, Infante R, Jacobson KL, Truong M. Clinical significance and epidemiologic analyses of

Mycobacterium avium and *Mycobacterium intracellulare* among patients without AIDS. J Clin Microbiol. 2005;43:4407.

- Kobashi Y, Abe M, Mouri K, Obase Y, Kato S, Oka M. Relationship between clinical efficacy for pulmonary MAC and drug-sensitivity test for isolated MAC in a recent 6-year period. J Infect Chemother. 2012;18:436.
- Kobashi Y, Yoshida K, Miyashita N, Niki Y, Oka M. Relationship between clinical efficacy of treatment of pulmonary *Mycobacterium avium* Complex disease and drug-sensitivity testing of *Mycobacterium avium* Complex isolates. J Infect Chemother. 2006;12:195.
- 36. van Ingen J, Boeree MJ, van Soolingen D, Mouton JW. Resistance mechanisms and drug susceptibility testing of nontuberculous mycobacteria. Drug Resist Updat. 2012;15:149.
- Wallace RJ Jr, Brown BA, Griffith DE, Girard WM, Murphy DT. Clarithromycin regimens for pulmonary *Mycobacterium avium* complex. The first 50 patients. Am J Respir Crit Care Med. 1996;153:1766.
- Tanaka E, Kimoto T, Tsuyuguchi K, Watanabe I, Matsumoto H, Niimi A, et al. Effect of clarithromycin regimen for *Mycobacterium avium* Complex pulmonary disease. Am J Respir Crit Care Med. 1999;160:866.
- 39. van Ingen J, Hoefsloot W, Mouton JW, Boeree MJ, van Soolingen D. Synergistic activity of rifampicin and ethambutol against slow-growing nontuberculous mycobacteria is currently of questionable clinical significance. Int J Antimicrob Agents. 2013;42:80.
- Hoffner SE, Heurlin N, Petrini B, Svenson SB, Källenius G. *Mycobacterium avium* complex develop resistance to synergistically active drug combinations during infection. Eur Respir J. 1994;7:247.
- 41. Brown-Elliott BA, Iakhiaeva E, Griffith DE, Woods GL, Stout JE, Wolfe CR, et al. *vitro* activity of amikacin against isolates of *Mycobacterium avium* complex with proposed MIC breakpoints and finding of a 16S rRNA gene mutation in treated isolates. J Clin Microbiol. 2013;51:3389.
- 42. Griffith DE, Brown BA, Murphy DT, Girard WM, Couch L, Wallace RJ Jr. Initial (6-month) results of three-times-weekly azithromycin in treatment regimens for *Mycobacterium avium* complex lung disease in human immunodeficiency virus-negative patients. J Infect Dis. 1998;178:121.
- 43. Lam PK, Griffith DE, Aksamit TR, Ruoss SJ, Garay SM, Daley CL, et al. Factors related to response to intermittent treatment of *Mycobacterium avium* complex lung disease. Am J Respir Crit Care Med. 2006;173:1283.
- 44. Griffith DE, Brown BA, Cegielski P, Murphy DT, Wallace RJ Jr. Early results (at 6 months) with intermittent clarithromycin-including regimens for lung disease due to *Mycobacterium avium* complex. Clin Infect Dis. 2000;30:288.
- 45.•• van Ingen J, Ferro BE, Hoefsloot W, Boeree MJ, van Soolingen D. Drug treatment of pulmonary nontuberculous mycobacterial disease in HIV-negative patients: the evidence. Expert Rev Anti-Infect Ther. 2013;11:1065.

A comprehensive review of drug treatment of nontuberculous mycobacterial lung disease by pioneering experts in the field.

- Olivier KN, Shaw PA, Glaser TS, Bhattacharyya D, Fleshner M, Brewer CC, et al. Inhaled amikacin for treatment of refractory pulmonary nontuberculous mycobacterial disease. Ann Am Thorac Soc. 2014;11:30.
- 47. Davis KK, Kao PN, Jacobs SS, Ruoss SJ. Aerosolized amikacin for treatment of pulmonary *Mycobacterium avium* infections: an observational case series. BMC Pulm Med. 2007;7:2.
- Safdar A. Aerosolized amikacin in patients with difficult-to-treat pulmonary nontuberculous mycobacteriosis. Eur J Clin Microbiol Infect Dis 2012; 31: 1883.
- Griffith DE, Brown-Elliott BA, Langsjoen B, Zhang Y, Pan X, Girard W, et al. Clinical and molecular analysis of macrolide resistance in *Mycobacterium avium* complex lung disease. Am J Respir Crit Care Med. 2006;174:928.
- 50. Ferro BE, Meletiadis J, Wattenberg M, de Jong A, van Soolingen D, Mouton JW, et al. Clofazimine prevents the regrowth of *Mycobacterium abscessus* and *Mycobacterium avium* type strains exposed to amikacin and clarithromycin. Antimicrob Agents Chemother. 2015;60:1097.
- 51.• Jarand J, Davis JP, Cowie RL, Field SK, Fisher DA. Longterm follow-up of *Mycobacterium avium* complex lung disease in patients treated with regimens including clofazimine and/or rifampin. Chest. 2016;149:1285.

In 107 patients with *Mycobacterium avium* complex lung disease followed for at least 6 months after completion of treatment, patients who received a macrolide, ethambutol, and clofazamine had significantly greater sputum conversion than those that received macrolide, ethambutol, and rifampin although there was no difference in microbiologic relapse or retreatment rates (occurring in about one-third) between the twotreatment groups. These findings lend credence to the increasing evidence that clofazamine should be considered in the regimen against nontuberculous mycobacteria (NTM). The study also suggests that a significant number of those with recurrent disease may in fact have a new NTM infection from the environment rather than a true relapse.

- 52. van Ingen J, Totten SE, Helstrom NK, Heifets LB, Boeree MJ, Daley CL. *In vitro* synergy between clofazimine and amikacin in treatment of nontuberculous mycobacterial disease. Antimicrob Agents Chemother. 2012;56:6324.
- 53. Roussel G, Igual J. Clarithromycin with minocycline and clofazimine for Mycobacterium avium intracellulare complex lung disease in patients without the acquired immune deficiency syndrome. GETIM. Groupe d'Etude et de Traitement des Infections à Mycobactéries. Int J Tuberc Lung Dis. 1998;2:462.
- 54. Winthrop KL, Ku JH, Marras TK, Griffith DE, Daley CL, Olivier KN, et al. The tolerability of linezolid in the treatment of nontuberculous mycobacterial disease. Eur Respir J. 2015;45:1177.

- Philley JV, Wallace RJ Jr, Benwill JL, Taskar V, Brown-Elliott BA, Thakkar F, et al. Preliminary results of bedaquiline as salvage therapy for patients with nontuberculous mycobacterial lung disease. Chest. 2015;148:499.
- Koh WJ, Hong G, Kim SY, Jeong BH, Park HY, Jeon K, et al. Treatment of refractory *Mycobacterium avium* complex lung disease with a moxifloxacin-containing regimen. Antimicrob Agents Chemother. 2013;57:2281.
- 57. Bittner MJ, Horowitz EA, Safranek TJ, Preheim LC. Emergence of *Mycobacterium kansasii* as the leading mycobacterial pathogen isolated over a 20-year period at a midwestern Veterans Affairs hospital. Clin Infect Dis. 1996;22:1109.
- Wang J, McIntosh F, Radomski N, Dewar K, Simeone R, Enninga J, et al. Insights on the emergence of *Mycobacterium tuberculosis* from the analysis of *Mycobacterium kansasii*. Genome Biol Evolution. 2015;7:856.
- Moon SM, Park HY, Jeon K, Kim S-Y, Chung MJ, Huh HJ, et al. Clinical significance of *Mycobacterium kansasii* isolates from respiratory specimens. PLoS One. 2015;10:e0139621.
- 60. Santin M, Dorca J, Alcaide F, Gonzalez L, Casas S, Lopez M, et al. Long-term relapses after 12-month treatment for *Mycobacterium kansasii* lung disease. Eur Respir J. 2009;33:148.
- 61. Woods GL, Brown-Elliott BA, Conville PS, Desmond EP, Hall GS, Lin G, et al. Susceptibility testing of mycobacteria, nocardiae, and other aerobic actinomtcetes: Approved Standard. In: Institute CaLS, editor. CSLI document M24-A2, 2nd ed. Wayne, Pennsylvania; 2011.
- 62. Shitrit D, Baum GL, Priess R, Lavy A, Shitrit AB, Raz M, et al. Pulmonary *Mycobacterium kansasii* infection in Israel, 1999-2004: clinical features, drug susceptibility, and outcome. Chest. 2006;129:771.
- 63. Griffith DE, Brown-Elliott BA, Wallace RJ Jr. Thriceweekly clarithromycin-containing regimen for treatment of *Mycobacterium kansasii* lung disease: results of a preliminary study. Clin Infect Dis. 2003;37:1178.
- 64. Kaustova J, Chmelik M, Ettlova D, Hudec V, Lazarova H, Richtrova S. Disease due to *Mycobacterium kansasii* in the Czech Republic: 1984-89. Tuber Lung Dis. 1995;76:205.
- 65. *Mycobacterium kansasii* pulmonary infection: a prospective study of the results of nine months of treatment with rifampicin and ethambutol. Research Committee, British Thoracic Society. Thorax. 1994; 49: 442.
- Ahn CH, Lowell JŘ, Ahn SS, Ahn SI, Hurst GA. Shortcourse chemotherapy for pulmonary disease caused by *Mycobacterium kansasii*. Am Rev Respir Dis. 1983;128:1048.
- 67. Research Committee of the British Thoracic Society. Pulmonary disease caused by *M. malmoense* in HIV negative patients: 5-yr follow-up of patients receiving standardised treatment. Eur Respir J. 2003;21:478.
- 68. Murray MP, Laurenson IF, Hill AT. Outcomes of a standardized triple-drug regimen for the treatment of

nontuberculous mycobacterial pulmonary infection. Clin Infect Dis. 2008;47:222.

- 69. Hoefsloot W, van Ingen J, de Lange WC, Dekhuijzen PN, Boeree MJ, van Soolingen D. Clinical relevance of *Mycobacterium malmoense* isolation in The Netherlands. Eur Respir J. 2009;34:926.
- Wayne LG, Sramek HA. Agents of newly recognized or infrequently encountered mycobacterial diseases. Clin Microbiol Rev. 1992;5:1.
- 71. van Ingen J, de Zwaan R, Dekhuijzen R, Boeree M, van Soolingen D. Region of difference 1 in nontuberculous mycobacterium species adds a phylogenetic and taxonomical character. J Bacteriol. 2009;191:5865.
- 72. van Ingen J, Boeree MJ, de Lange WC, de Haas PE, Dekhuijzen PN, van Soolingen D. Clinical relevance of *Mycobacterium szulgai* in The Netherlands. Clin Infect Dis. 2008;46:1200.
- 73. van Ingen J, Boeree M, Janssen M, Ullmann E, de Lange W, de Haas P, et al. Pulmonary *Mycobacterium szulgai* infection and treatment in a patient receiving antitumor necrosis factor therapy. Nat Clin Pract Rheumatol. 2007;3:414.
- 74. Marks J, Schwabacher H. Infection due to *Mycobacterium xenopi*. Br Med J. 1965;1:32.
- 75. Hoefsloot W, van Ingen J, Andrejak C, Angeby K, Bauriaud R, Bemer P, et al. Nontuberculous Mycobacteria Network European Trials Group. The geographic diversity of nontuberculous mycobacteria isolated from pulmonary samples: an NTM-NET collaborative study. Eur Respir J. 2013;42:1604.

This study tabulated the major NTM species isolated from respiratory samples and reported in 2008 to the NTM-Network European Trials Group. In data from over 20,000 patients from 30 countries across sixcontinents, *Mycobacterium avium* complex (MAC) organisms predominated in most countries. Importantly, there were significant differences among the different countries and continents in the geographical distribution of MAC species and other more common NTM.

- 76. Martín-Casabona N, Bahrmand AR, Bennedsen J, Thomsen VO, Curcio M, Fauville-Dufaux M, et al. Spanish Group for Non-Tuberculosis Mycobacteria. Non-tuberculous mycobacteria: patterns of isolation. A multi-country retrospective survey. Int J Tuberc Lung Dis. 2004;8:1186.
- 77. van Ingen J, Boeree MJ, de Lange WC, Hoefsloot W, Bendien SA, Magis-Escurra C, et al. *Mycobacterium xenopi* clinical relevance and determinants, the Netherlands. Emerg Infect Dis. 2008;14:385.
- Andréjak C, Lescure FX, Pukenyte E, Douadi Y, Yazdanpanah Y, Laurans G, et al. *Mycobacterium xenopi* pulmonary infections: a multicentric retrospective study of 136 cases in north-east France. Thorax. 2009;64:291.
- Banks J, Jenkins PA. Combined versus single antituberculosis drugs on the *in vitro* sensitivity patterns of non-tuberculous mycobacteria. Thorax. 1987;42:838.

- 80. Watt B. *In-vitro* sensitivities and treatment of less common mycobacteria. J Antimicrob Chemother. 1997;39:567.
- Ferro BE, van Ingen J, Wattenberg M, van Soolingen D, Mouton JW. Time-kill kinetics of slowly growing mycobacteria common in pulmonary disease. J Antimicrob Chemother. 2015;70:2838.
- van Ingen J, van der Laan T, Dekhuijzen R, Boeree M, van Soolingen D. *In vitro* drug susceptibility of 2275 clinical non-tuberculous mycobacterium isolates of 49 species in The Netherlands. Int J Antimicrob Agents. 2010;35:169.
- 83.•• Andréjak C, Almeida DV, Tyagi S, Converse PJ, Ammerman NC, Grosset JH. Improving existing tools for *Mycobacterium xenopi* treatment: assessment of drug combinations and characterization of mouse models of infection and chemotherapy. J Antimicrob Chemother. 2013;68:659.

Lung disease due to *Mycobacterium xenopi* is often recalcitrant to treatment. This comprehensive study examined potential drugs and combinations of drugs against this organism using in vitro (drugs against bugs), ex vivo (sera of treated mice to assess antimicrobial activities), and in vivo (murine) models. The investigators found that in vitro, 1) ethambutol plus either rifamycin or moxifloxacin and 2) clarithromycin plus moxifloxacin showed the best bactericidal activities; ex vivo, a three-drug combination of ethambutolplus a rifamycin and either clarithromycin or moxifloxacin was best. Interestingly, for the in vivo mouse studies, amikacin-containing regimen had the greatest bactericidal activity with no difference in regimens containing clarithromycin or moxifloxacin. This study is important because it provides a valuable foundation for future clinical trials in humans.

- Varadi RG, Marras TK. Pulmonary Mycobacterium xenopi infection in non-HIV-infected patients: a systematic review. Int J Tuberc Lung Dis. 2009;13:1210.
- 85. Watt B, Rayner A, Harris G. Comparative activity of azithromycin against clinical isolates of mycobacteria. J Antimicrob Chemother. 1996;38:539.
- van Ingen J, Boeree MJ, Dekhuijzen PN, van Soolingen D. Clinical relevance of *Mycobacterium simiae* in pulmonary samples. Eur Respir J. 2008;31:106.
- Krasnow I, Gross W. *Mycobacterium simiae* infection in the United States. A case report and discussion of the organism. Am Rev Respir Dis. 1975;111:357.
- Boisvert H, Truffot C. Relationships between "Mycobacterium simiae" and the "M. avium-intracellularescrofulaceum" complex (author's transl. Ann Microbiol (Paris). 1979;130B:457.
- Bell RC, Higuchi JH, Donovan WN, Krasnow I, Johanson WG Jr. *Mycobacterium simiae*. Clinical features and follow-up of twenty-four patients. Am Rev Respir Dis. 1983;127:35.
- Valero G, Peters J, Jorgensen JH, Graybill JR. Clinical isolates of *Mycobacterium simiae* in San Antonio, Texas. An 11-yr review. Am J Respir Crit Care Med. 1995;152:1555.
- 91. Valero G, Moreno F, Graybill JR. Activities of clarithromycin, ofloxacin, and clarithromycin plus

ethambutol against *Mycobacterium simiae* in murine model of disseminated infection. Antimicrob Agents Chemother. 1994;38:2676.

- 92. Jeong SH, Kim SY, Lee H, Ham JS, Hwang KB, Hwang S, et al. Nontuberculous mycobacterial lung disease caused by *Mycobacterium simiae*: the first reported case in South Korea. Tuberc Respir Dis (Seoul). 2015;78:432.
- 93. Philley JV, Griffith DE. Treatment of slowly growing mycobacteria. Clin Chest Med. 2015;36:79.
- 94. Koh WJ, Stout JE, Yew WW. Advances in the management of pulmonary disease due to Mycobacterium abscessus complex. Int J Tuberc Lung Dis. 2014;18:1141.
- 95. Pang H, Li G, Wan L, Jiang Y, Liu H, Zhao X, et al. *In vitro* drug susceptibility of 40 international reference rapidly growing mycobacteria to 20 antimicrobial agents. Int J Clin Exp Med. 2015;8:15423.
- 96. Broda A, Jebbari H, Beaton K, Mitchell S, Drobniewski F. Comparative drug resistance of *Mycobacterium abscessus* and *M. chelonae* isolates from patients with and without cystic fibrosis in the United Kingdom. J Clin Microbiol. 2013;51:217.
- 97. Document M100-S11. Clinical and Laboratory Standards Institute (CLSI). Performance standards for antimicrobial susceptibility testing. Eleventh informational supplement. Document no. M100-S11. Wayne, PA: CLSI; 2001.
- Ryu YJ, Koh WJ, Daley CL. Diagnosis and treatment of nontuberculous mycobacterial lung disease: Clinicians' perspectives. Tuberc Respir Dis (Seoul). 2016;79:74.
- Philley JV, Griffith DE. Management of nontuberculous mycobacterial (NTM) lung disease. Semin Respir Crit Care Med. 2013;34:135.
- 100. Jeon K, Kwon OJ, Lee NY, Kim BJ, Kook YH, Lee SH, et al. Antibiotic treatment of *Mycobacterium abscessus* lung disease: a retrospective analysis of 65 patients. Am J Respir Crit Care Med. 2009;180:896.
- 101. Nash KA, Brown-Elliott BA, Wallace RJ Jr. A novel gene, erm(41), confers inducible macrolide resistance to clinical isolates of *Mycobacterium abscessus* but is absent from *Mycobacterium chelonae*. Antimicrob Agents Chemother. 2009;53:1367.
- 102. Choi GE, Shin SJ, Won CJ, Min KN, Oh T, Hahn MY, et al. Macrolide treatment for *Mycobacterium abscessus* and *Mycobacterium massiliense* infection and inducible resistance. Am J Respir Crit Care Med. 2012;186:917.
- 103. Maurer FP, Castelberg C, Quiblier C, Böttger EC, Somoskövi A. Erm(41)-dependent inducible resistance to azithromycin and clarithromycin in clinical isolates of *Mycobacterium abscessus*. J Antimicrob Chemother. 2014;69:1559.
- 104. •• Koh WJ, Jeon K, Lee NY, Kim BJ, Kook YH, Lee SH, et al. Clinical significance of differentiation of *Mycobacterium massiliense* from *Mycobacterium abscessus*. Am J Respir Crit Care Med. 2011;183:405.

This study sheds new light on why some patients with *M. abscessus* lung disease respond poorly to treatment whereas others do better. Previous work showed that with more robust

speciation, *M. abscessus* organisms are comprised of at least three distinct species within the so-called *M. abscessus* complex: *M.abscessus* sensu stricto, *M. bolletii*, and *M. massiliense*. In patients infected with *M. abscessus* sensu stricto, response to treatment is significantly poorer than those with *M. massiliense*; this is likely due, in large part, to the presence of an inducible gene in *M. abscessus* sensu stricto that confers resistance to a macrolide in the presence, interestingly, of a macrolide.

- 105. Kim HS, Lee KS, Koh WJ, Jeon K, Lee EJ, Kang H, et al. Serial CT findings of Mycobacterium massiliense pulmonary disease compared with *Mycobacterium abscessus* disease after treatment with antibiotic therapy. Radiology. 2012;263:260.
- 106. Lyu J, Kim BJ, Kim BJ, Song JW, Choi CM, Oh YM, et al. A shorter treatment duration may be sufficient for patients with *Mycobacterium massiliense* lung disease than with *Mycobacterium abscessus* lung disease. Respir Med. 2014;108:1706.
- Wallace RJ Jr, Brown-Elliott BA, Ward SC, Crist CJ, Mann LB, Wilson RW. Activities of linezolid against rapidly growing mycobacteria. Antimicrob Agents Chemother. 2001;45:764.
- 108. Jarand J, Levin A, Zhang L, Huitt G, Mitchell JD, Daley CL. Clinical and microbiologic outcomes in patients receiving treatment for *Mycobacterium abscessus* pulmonary disease. Clin Infect Dis. 2011;52:565.
- 109. Kasperbauer SH, De Groote MA. The treatment of rapidly growing mycobacterial infections. Clin Chest Med. 2015;36:67.
- Mitchell JD. Surgical approach to pulmonary nontuberculous mycobacterial infections. Clin Chest Med. 2015;36:117.
- Nelson KG, Griffith DE, Brown BA, Wallace RJ Jr. Results of operation in *Mycobacterium avium-intracellulare* lung disease. Ann Thorac Surg. 1998;66:325.
- 112. Shiraishi Y, Nakajima Y, Takasuna K, Hanaoka T, Katsuragi N, Konno H. Surgery for *Mycobacterium avium* complex lung disease in the clarithromycin era. Eur J Cardiothorac Surg. 2002;21:314.
- 113. Yu JA, Pomerantz M, Bishop A, Weyant MJ, Mitchell JD. Lady Windermere revisited: treatment with thoracoscopic lobectomy/segmentectomy for right middle lobe and lingular bronchiectasis associated with non-tuberculous mycobacterial disease. Eur J Cardiothorac Surg. 2011;40:671.
- 114. Mitchell JD, Bishop A, Cafaro A, Weyant MJ, Pomerantz M. Anatomic lung resection for nontuberculous mycobacterial disease. Ann Thorac Surg. 2008;85:1887.
- 115. Watanabe M, Hasegawa N, Ishizaka A, Asakura K, Izumi Y, Eguchi K, et al. Early pulmonary resection for Mycobacterium avium complex lung disease treated with macrolides and quinolones. Ann Thorac Surg. 2006;81:2026.
- 116. Corpe RF. Surgical management of pulmonary disease due to *Mycobacterium avium-intracellulare*. Rev Infect Dis. 1981;3:1064.

- 117. Moran JF, Alexander LG, Staub EW, Young WG Jr, Sealy WC. Long-term results of pulmonary resection for atypical mycobacterial disease. Ann Thorac Surg. 1983;35:597.
- 118. Shiraishi Y, Katsuragi N, Kita H, Hyogotani A, Saito MH, Shimoda K. Adjuvant surgical treatment of nontuberculous mycobacterial lung disease. Ann Thorac Surg. 2013;96:287.
- 119. Shiraishi Y, Nakajima Y, Katsuragi N, Kurai M, Takahashi N. Pneumonectomy for nontuberculous mycobacterial infections. Ann Thorac Surg. 2004;78:399.
- 120. Griffith DE, Girard WM, Wallace RJ Jr. Clinical features of pulmonary disease caused by rapidly growing mycobacteria. An analysis of 154 patients. Am Rev Respir Dis. 1993;147:1271.
- 121. Andréjak C, Nielsen R, Thomsen VO, Duhaut P, Sørensen HT, Thomsen RW. Chronic respiratory disease, inhaled corticosteroids and risk of nontuberculous mycobacteriosis. Thorax. 2013;68:256.
- 122. Honda JR, Hess T, Malcolm KC, Ovrutsky AR, Bai X, Irani VR, et al. Pathogenic nontuberculous mycobacteria resist and inactivate cathelicidin: implication of a novel role for polar mycobacterial lipids. PLoS One. 2015;10:e0126994.
- 123. van Ingen J, Egelund EF, Levin A, Totten SE, Boeree MJ, Mouton JW, et al. The pharmacokinetics and pharmacodynamics of pulmonary *Mycobacterium*

avium complex disease treatment. Am J Respir Crit Care Med. 2012;186:559.

- 124. Ito Y, Hirai T, Fujita K, Kubo T, Maekawa K, Ichiyama S, et al. The influence of environmental exposure on the response to antimicrobial treatment in pulmonary *Mycobacterial avium* complex disease. BMC Infect Dis. 2014;14:522.
- 125. Kikuchi T, Kobashi Y, Hirano T, Tode N, Santoso A, Tamada T, et al. *Mycobacterium avium* genotype is associated with the therapeutic response to lung infection. Clin Microbiol Infect. 2014;20:256.
- 126. Kim SY, Lee ST, Jeong BH, Park HY, Jeon K, Kim JW, et al. Genotyping of *Mycobacterium intracellulare* isolates and clinical characteristics of lung disease. Int J Tuberc Lung Dis. 2013;17:669.
- 127. Schulze-Röbbecke R, Buchholtz K. Heat susceptibility of aquatic mycobacteria. Appl Environ Microbiol 1992; 58: 1869.
- 128. Falkinham JO 3rd. Nontuberculous mycobacteria from household plumbing of patients with nontuberculous mycobacteria disease. Emerg Infect Dis. 2011;17:419.
- 129. Chan ED, Iseman, MD. Bronchiectasis. In: Broaddus C EJ, King TE, Lazarus SC, Mason R, Murray J, Nadel J, Slutsky AS (editors). Textbook of Respiratory Medicine, 6th ed. in press: Elsevier Press; 2016.