A community-acquired lung abscess attributable to odontogenic flora

This article was published in the following Dove Press journal:
Infection and Drug Resistance

Wenjia Guo¹

Bo Gao¹

Li Li¹

Wei Gai²

Jianghui Yang³

Yan Zhang²

Lijun Wang⁴

¹Department of Pulmonary and Critical Care Medicine, Beijing Tsinghua Changgung Hospital, School of Clinical Medicine, Tsinghua University, Beijing, People's Republic of China; ²National Engineering Research Center for Beijing Biochip Technology, Beijing, People's Republic of China; ³Department of Pathology, Beijing Tsinghua Changgung Hospital, School of Clinical Medicine, Tsinghua University, Beijing, People's Republic of China; ⁴Clinical Research Center, Beijing Tsinghua Changgung Hospital, School of Clinical Medicine, Tsinghua University, Beijing, People's Republic of China

Abstract: A lung abscess is an infectious pulmonary disease characterized by pus-filled cavity formation and often an air-fluid level. In this article, we described an indolent community-acquired lung abscess suspected as a tumor previously. A 56-year-old male presented with cough and expectoration for 2 months and hemoptysis for 2 weeks. His physical examinations, whole blood count and C-reactive protein level were normal. The chest computed tomography (CT) scan showed a 40×38×39 mm high-density mass in the right upper pulmonary lobe, with irregular borders. The pathology of a CT-guided percutaneous needle aspiration biopsy showed numerous inflammatory cells and bacteria infiltration without tumor lesions. Bacteriological detection of lung tissue revealed the cause was odontogenic flora. A next-generation sequencing demonstrated the etiologic correlation between lung abscess and periodontitis. After a 2-month pathogen-directed oral antibiotics therapy combined with chlorhexidine gargle oral care, this patient showed a remarkable improvement. Periodontitis can be a cause of a lung abscess, which would be taken into account in the treatment regimes preventing infectious recurrence.

Keywords: community-acquired lung abscess, periodontitis, odontogenic flora

Introduction

A lung abscess is an infectious pulmonary disease characterized by pus-filled cavity formation and often an air-fluid level. Primary lung abscesses are often polymicrobial and contain multiple anaerobic species.¹ The bacteriological characteristics, however, have gradually changed. *Klebsiella pneumoniae*^{2,3} was reported as the most common organism in lung abscess cases in Asia, moreover, in alcoholics with poor oral hygiene, the spectrum of pathogens contains *Staphylococcus aureus*, *Streptococcus pyogenes* and *Actinomyces*. Thus, etiology diagnosis cannot be ignored. On the other hand, the most common predisposing factor was an unhygienic oral cavity.² Nevertheless, the impact of dental diseases on lung abscess is often underestimated in routine clinical practice. Here, we describe a case of indolent community-acquired lung abscess (CALA) due to odontogenic flora.

Case presentation

A 56-year-old male complained of cough and expectoration for 2 months and hemoptysis for 2 weeks. He reported no fever, dyspnea, chest pain or distress. On physical examination, his temperature was 36.5°C, heart rate 72 bpm, blood pressure 127/72 mmHg and oxygen saturation 98% when he was breathing

Correspondence: Lijun Wang
Clinical Research Center, Beijing Tsinghua Changgung Hospital, School of Clinical Medicine, Tsinghua University, No. 168 Litang Road, Changping District, Beijing 102218, People's Republic of China
Tel +86 1 381 037 0450
Email wlj7878@163.com

ambient air. His breath sounds were clear without rales. The laboratory tests revealed white blood cell count was 8.48×10^9 cell/L, containing 64.3% neutrophils, 26.8% lymphocytes and 7.5% monocytes. The C-reactive protein was 6 mg/L within a normal range. The chest computed tomography (CT) scan (Figure 1) showed a $40 \times 38 \times 39$ mm high-density mass in the right upper pulmonary lobe, with irregular borders. As a result, lung cancer was suspected. Then, we performed a CT-guided percutaneous fine-needle aspiration biopsy in order to

Figure 1 A chest computed tomography (CT) scan on admission demonstrated a high-density $40 \times 38 \times 39$ mm mass on the right upper lung lobe, with irregular borders.

identify either tumor or infection. The patient suffered from fever and the serum procalcitonin increased to 3.42 ng/mL after aspiration. The patient had underlying diseases, including diabetes mellitus and hyperlipemia. He had about 30-year history of smoking or alcohol consumption, and he quit smoking 7 years ago.

Bacteriological detection of expectorated sputum samples yielded no pathogens with routine aerobic culture. However, Gram stain of the lung biopsy suggested a variety of bacteria (Figure 2C); then, tissue culture detected moderate *Fusobacterium nucleatum* and *Parvimonas micra*. In addition, the pathology showed significant inflammatory cells and bacteria infiltration without tumor lesions (Figure 2A and B). Thus, the diagnosis of a community-acquired pulmonary abscess was concluded.

Since the isolated pathogens are oral cavity microbiome constitute, we further performed the oral examination and dental orthopantomography (Figure 3). The patient had severe chronic periodontitis.

The antibiotics regimes are metronidazole (0.4 g, tid) and oral amoxicillin/clavulanate potassium (1 mg/0.5 mg, tid) for 2 months. At the same time, oral care with chlorhexidine gargle was advocated every day. After therapy, the mass decreased remarkably (Figure 4). Now, the patient was recovered.

Figure 2 Pathology findings with inflammatory cells infiltration (A) and bacteria cluster (B) (HE, $\times 200$). Gram staining of lung tissues showed a variety of bacteria (C). Black arrows indicate bacteria evidence.

Figure 3 Several odontolith with high-density on dental orthopantomography.

Figure 4 Chest CT showed a significantly decreased lung mass lesion after 2-month therapy.

Abbreviation: CT, computed tomography.

Next-generation sequencing (NGS)

A NGS of both the lung tissue and odontolith was performed by National Engineering Research Center for Beijing Biochip Technology, in order to explore the correlation between periodontitis and lung abscess.

Multiple anaerobic bacteria genomes were found in the lung tissue and listed in [Table 1](#), comprising of *Porphyromonas gingivalis*, *P. micra*, *F. nucleatum*, *Tannerella forsythia* and *Treponema* species. Our bacteriological culture coincides with NGS results. Moreover, the NGS results from odontolith were generally

Table 1 The next-generation sequencing (NGS) from the lung tissue

Name	Gene length
<i>Porphyromonas gingivalis</i> _SJD	2,508,424
<i>Parvimonas micra</i>	1,132,442
<i>Fusobacterium nucleatum</i> _subs	1,076,720
<i>Tannerella forsythia</i> _3313	1,029,601
<i>Treponema denticola</i> _ASLM	54,082

Table 2 The next-generation sequencing (NGS) from the odontolith

Name	Gene length
<i>Propionibacterium acidifaciens</i> _F0233	2,100,174
<i>Tannerella forsythia</i>	1,505,894
<i>Desulfobulbus</i> _sp._oral_taxon_041_str._Dsb1-5	438,407
<i>Treponema socranskii</i> _subsp._ _socranskii_VPI_DR56BR1116_=_ATCC_35536	290,118
<i>Selenomonas noxia</i> _F0398	167,091
<i>Corynebacterium matruchotii</i> _ATCC_33806	138,896
<i>Actinomyces massiliensis</i> _4401292	131,469
<i>Capnocytophaga granulosa</i> _ATCC_51502	122,080
<i>Porphyromonas gingivalis</i> _SJD12	60,964
<i>Neisseria sicca</i> _ATCC_29256	49,590
<i>Veillonella parvula</i> _HSIVPI	48,251

consistent with lung tissue ([Table 2](#)), including *T. forsythia*, *P. gingivalis* and *Treponema* species. The NGS confirmed the etiologic correlation between lung abscess and periodontitis.

Discussion

Classically, lung abscess could be classified into aspiration lung abscess, secondary lung abscess and hematogenous lung abscess, while aspiration pneumonia could be the most common cause.⁴ Actually, anaerobes comprise 60–80% of lung abscess etiologic pathogens.¹ Dental diseases such as gingivitis usually provide the inoculums in which large quantities of anaerobic bacteria are colonized and then spread into lung to develop aspiration lung abscess. Takayanagi⁵ investigated the etiologic pathogens of CALA in Japan. It indicated that periodontal disease was highly present in 61% of these patients with lung abscess. Meanwhile, alcohol ingestion was the most important risk factor.² In our case, the patient not only had a history of alcohol consumption and a severe periodontitis, but also most of the pathogens in odontolith can be found in the abscess lesion. Thus, chronic periodontitis is likely to be a cause for lung abscess in this patient.

Previously, mainly bacterial cultures of sputum and bronchoalveolar lavage fluid had been used to identify the etiology of lung abscess. Due to the contamination of oral flora, the etiology could not be clarified in many cases.¹ Percutaneous aspiration biopsy is a promising method at a time when majority clinicians have ignored performing any bacteriological tests in cases of pulmonary infection. The differential diagnosis in this patient just benefits from the aspiration biopsy. Since lung abscesses are often mixed infection, conventional bacterial culture is

challenging to recover all the pathogens. Until the last decade, a new class of sequencing platforms defined “Next Generation Sequencing” (NGS) technologies has been developed and applied for the identification of pathogens in clinical material.⁶ Recently a blinded, prospective cohort study⁷ was published to evaluate the value of NGS in patients with suspected infection. As a result, NGS provided a high negative predictive value with 96±5%. Moreover, the detection rates of pathogens with NGS were significantly higher than that with conventional methods (36% vs 11%). To further determine the origin of pathogens, NGS technology with both biopsy tissue and odontolith were utilized and odontogenic flora were confirmed as the pathogens. To our knowledge, it is the first report that a patient with odontogenic CALA was confirmed by NGS. The sequencing could make potentially contributions to mixed infection, especially exploring the source of the infection to make effective strategy.⁸

In case of etiology of lung abscess is clear, targeting therapy would be cost-effective. In general, all anaerobes isolated from the respiratory samples are susceptible to beta-lactam/beta-lactamase combinations such as amoxicillin-clavulanate. After 2-month oral antibiotics, amoxicillin-clavulanate and metronidazole, combined with chlorhexidine gargle oral care, this patient showed a remarkable improvement.

Conclusion

Dental diseases can be a cause of a lung abscess, which is often easily overlooked. Then, dental treatment is as significant as antibiotics in the therapy of lung abscess with chronic periodontitis. Finally, NGS will be a promising method for microbiological diagnosis and tracking trace in patients with infection.

Abbreviations

CALA, community-acquired lung abscess; NGS, next-generation sequencing technology.

Ethics approval and consent to participate

The ethics committee of Beijing Tsinghua Changgung Hospital approved the study procedure. The reference number of the approval is 19117-0-01. The patient provided a written informed consent for the case details to be published.

Availability of data and material

All the related data are presented in the manuscript.

Disclosure

The authors report no conflicts of interest in this work.

References

1. Bartlett JG. The role of anaerobic bacteria in lung abscess. *Clin Infect Dis*. 2005;40:923–925. doi:10.1086/428586
2. Mohapatra MM, Rajaram M, Mallick A. Clinical, radiological and bacteriological profile of lung abscess – an observational hospital based study. *Open Access Maced J Med Sci*. 2018;6:1642–1646. doi:10.3889/oamjms.2018.374
3. Wang JL, Chen KY, Fang CT, Hsueh PR, Yang PC, Chang SC. Changing bacteriology of adult community-acquired lung abscess in Taiwan: *Klebsiella pneumoniae* versus anaerobes. *Clin Infect Dis*. 2005;40:915–922. doi:10.1086/428574
4. Yazbeck MF, Dahdel M, Kalra A, Browne AS, Pratter MR. Lung abscess: update on microbiology and management. *Am J Ther*. 2014;21:217–221. doi:10.1097/MJT.0b013e3182383c9b
5. Takayanagi N, Kagiya N, Ishiguro T, Tokunaga D, Sugita Y. Etiology and outcome of community-acquired lung abscess. *Respiration*. 2010;80:98–105. doi:10.1159/000312404
6. Deurenberg RH, Bathoorn E, Chlebowicz MA, et al. Application of next generation sequencing in clinical microbiology and infection prevention. *J Biotechnol*. 2017;243:16–24. doi:10.1016/j.jbiotec.2016.12.022
7. Parize P, Muth E, Richaud C, et al. Untargeted next-generation sequencing-based first-line diagnosis of infection in immunocompromised adults: a multicentre, blinded, prospective study. *Clin Microbiol Infect*. 2017;23:574 e1–e6. doi:10.1016/j.cmi.2017.02.006
8. Besser J, Carleton HA, Gerner-Smith P, Lindsey RL, Trees E. Next-generation sequencing technologies and their application to the study and control of bacterial infections. *Clin Microbiol Infect*. 2018;24:335–341. doi:10.1016/j.cmi.2017.10.013

Infection and Drug Resistance

Publish your work in this journal

Infection and Drug Resistance is an international, peer-reviewed open-access journal that focuses on the optimal treatment of infection (bacterial, fungal and viral) and the development and institution of preventive strategies to minimize the development and spread of resistance. The journal is specifically concerned with the epidemiology of

antibiotic resistance and the mechanisms of resistance development and diffusion in both hospitals and the community. The manuscript management system is completely online and includes a very quick and fair peer-review system, which is all easy to use. Visit <http://www.dovepress.com/testimonials.php> to read real quotes from published authors.

Submit your manuscript here: <https://www.dovepress.com/infection-and-drug-resistance-journal>

Dovepress