



A Case of Localized Fungal Pneumonia Caused by *Rhodotorula mucilaginosa* in an Immunocompetent Patient

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Dear Editor,

Rhodotorula species, belonging to the family *Sporidiobolaceae*, are yeasts that are present in the environment and can cause opportunistic infections in immunocompromised patients [1, 2]. Among *Rhodotorula* species, only *R. mucilaginosa*, *R. glutinis*, and *R. minuta* have been reported to cause human infections [1, 2]. *Rhodotorula* sp. are mostly isolated from the blood of immunocompromised patients, where they cause fungemia or catheter-related infections. Only one case of pneumonia caused by *R. mucilaginosa* has been reported to date in an immunocompromised patient [3]. We report the first case of fungal pneumonia caused by *R. mucilaginosa* in an immunocompetent patient. The infection was confirmed by fungal ribosomal RNA (rRNA) sequencing using wedge-resected lung tissue.

This case report was approved by the Institutional Review Board of Hallym University Dongtan Sacred Heart Hospital, Hwaseong, Korea (IRB No. 2020-03-017). A waiver of consent was granted due to the retrospective nature of this study. A 52-year-old man visited our hospital in July 2019 because of a pulmonary nodule found incidentally during a health check-up at a local clinic in May 2019. The patient did not have any underlying disease and his complete blood count was within the normal range. A lobu-

lated nodular lesion measuring 3.5×1.6 cm was observed in the left lower lobe on radiological examination (computed tomography (CT); Fig. 1A). Serological screening tests for parasites (*Clonorchis*, *Paragonimus*, *Cysticercus*, *Sparganum*, and *Toxocara*) and the *Aspergillus* antigen (Galactomannan) test yielded negative results. Histopathological evaluation of a percutaneous needle biopsy revealed scattered, brown, oval, yeast spores (5–7 μm) and chronic granulomatous inflammation, leading us to suspect fungal infection (Fig. 1B) [4]. Fungal culture using needle biopsy tissue was not performed at that time.

Empirical antifungal therapy with fluconazole 400 mg/day was initiated and continued for three months (Aug 14–Nov 23, 2019), because cryptococcal infection was suspected based on the histopathological findings. During antifungal treatment, CT revealed a slight decrease in nodule size (2.6×1.6 cm), which increased (to 3.1×1.9 cm) three months after discontinuation of the antifungal therapy. Wedge resection of the lung lesion was performed for definite diagnosis and treatment on March 5, 2020, and the fresh specimen was partitioned and sent for histopathological evaluation, fungal culture, and fungal rRNA sequencing. The histopathological features of the wedge-resected lung tissue were similar to those of the previous needle biopsy specimen.

Received: March 26, 2020

Revision received: April 21, 2020

Accepted: July 20, 2020

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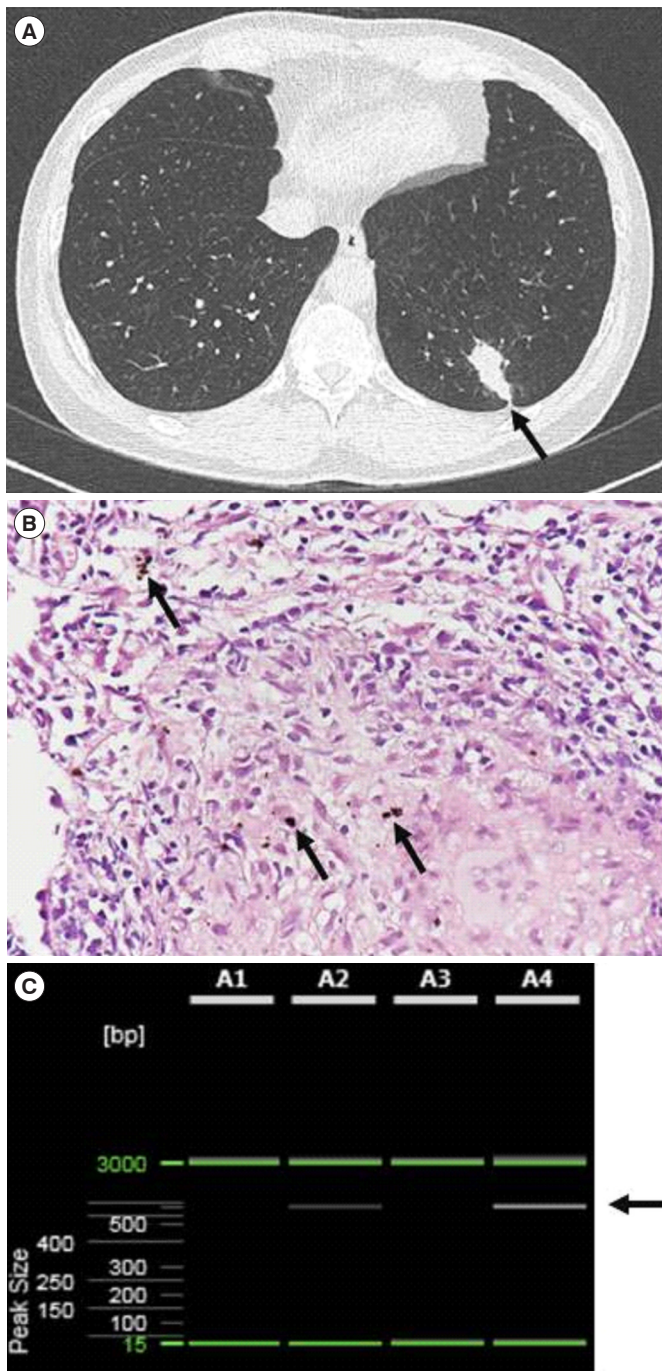


Fig. 1. Radiologic, histopathological, and PCR findings regarding *Rhodotorula* pneumonia for the case presented. (A) Chest CT scan shows a nodular lesion (arrow) in the left lower lobe. (B) The lung biopsy section shows scattered, brown, oval, yeast spores (arrows) and chronic granulomatous inflammation (Hematoxylin-eosin stain, $\times 400$). (C) Capillary electrophoresis of PCR products shows no amplicon for the fungal ITS1 region (lanes A1 and A3) and distinct amplicons for the fungal D1/D2 region of the large subunit rRNA (lanes A2 and A4).

Abbreviations: CT, computed tomography; rRNA, ribosomal RNA; ITS1, internal transcribed spacer 1.

Fungal culture of the resected lung tissue showed no growth, probably due to the antifungal treatment.

DNA extracted from the wedge-resected lung tissue was used for PCR amplification and sequencing of the fungal internal transcribed spacer 1 (ITS1) region and the D1/D2 region of the large subunit rRNA gene (Fig. 1C) [5]. The ITS1 region was not amplified; however, the D1/D2 region of the large subunit rRNA gene showed 100% (560 bps /560 bps) sequence identity with the corresponding region in *R. mucilaginosa* based on a basic local alignment search tool using the sequence obtained (<https://blast.ncbi.nlm.nih.gov/>).

As *R. mucilaginosa* pneumonia has never been reported in immunocompetent patients, it was extremely difficult to draw the differential diagnosis of pulmonary nodules. We thought that cryptococcal infection was the most likely type of fungal infection and began antifungal treatment. *R. mucilaginosa* pneumonia was diagnosed based on the correlation between histopathological findings and fungal rRNA sequencing data. *Rhodotorula* contamination from the environment could be ruled out, as similar histopathological findings regarding fungal morphology in granulomatous regions were observed in two biopsy sections taken six months apart. This patient was discharged after wedge biopsy, and the solitary pulmonary nodule completely disappeared on follow-up chest X-ray, one month later.

This case is unique compared with the previously-reported *Rhodotorula* infection cases from Korea, which involved either immunocompromised patients or patients undergoing invasive procedures (i.e., central venous catheterization and continuous peritoneal dialysis) [6–8], in that our patient was immunocompetent, without ongoing invasive procedures.

In summary, we report the first case of *R. mucilaginosa* pneumonia in an immunocompetent patient. The diagnosis was made based on combined histopathological and fungal rRNA sequencing findings in wedge-resected lung tissue. *R. mucilaginosa* pneumonia infection should be considered when conducting differential diagnosis for pulmonary nodules in immunocompetent patients.

ACKNOWLEDGEMENTS

We are grateful to Ju Un Park for the excellent technical assistance.

AUTHOR CONTRIBUTIONS

HSK designed and performed the study, obtained ethical ap-

proval from IRB, and wrote the manuscript; HK collected the data and wrote the manuscript; CHK and HL diagnosed and treated the patient, provided clinical information, and edited the manuscript; YB, SB, HK, and HSK contributed to the diagnosis of *Rhodotorula* infection. All authors read and approved the final manuscript.

CONFLICTS OF INTEREST

None declared.

RESEARCH FUNDING

None declared.

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