Triple Trouble: Disseminated Penicilliosis in a Cancer patient with COVID-19

Sujeet Kamtalwar1,2  Sumeet Mirgh2,3  Ashwini More1,2  Anant Gokarn2,3  Sachin Dhumal2,4  Palak Sharma2,5  Sujata Lall2,6  Nikhil Patkar2,7  Sweta Rajpal2,9  Vivek Bhat2,6  Navin Khattry2,3  Sudeep Gupta2,3

1 Department of General Medicine, Advanced Centre for Treatment, Research and Education in Cancer, ACTREC, Kharghar, Maharashtra, India
2 Homi Bhabha National Institute, Mumbai, India
3 Department of Medical Oncology, Advanced Centre for Treatment, Research and Education in Cancer, ACTREC, Kharghar, Maharashtra, India
4 Department of Radiation Oncology, Advanced Centre for Treatment, Research and Education in Cancer, ACTREC, Kharghar, Maharashtra, India
5 Department of Medical Administration, Advanced Centre for Treatment, Research and Education in Cancer, ACTREC, Kharghar, Maharashtra, India
6 Department of Microbiology, Advanced Centre for Treatment, Research and Education in cancer, ACTREC, Kharghar, Maharashtra, India
7 Department of Hematolymphoid, Advanced Centre for Treatment, Research and Education in cancer, ACTREC, Kharghar, Maharashtra, India
8 Department of Radiodiagnosis, Advanced Centre for Treatment, Research and Education in cancer, ACTREC, Kharghar, Maharashtra, India
9 Department of Hematopathology Laboratory, Advanced Centre for Treatment, Research and Education in Cancer, ACTREC, Kharghar, Maharashtra, India

Address for correspondence Sumeet Mirgh, MD, DM, Adult Hematolymphoid and BMT, Department of Medical Oncology, Advanced Centre for Treatment, Research and Education in Cancer (ACTREC), Kharghar, Maharashtra, India and Homi Bhabha National Institute, Mumbai, India (e-mail: drsumeetmirgh@gmail.com).

Abstract
Penicilliosis is a fungal infection caused by the fungus Penicillium marneffei or Talaromyces marneffei. Penicilliosis is commonly seen in immunocompromised patients such as in HIV (AIDS). Herein, we present a case of penicilliosis in an oral cavity cancer patient who was admitted for the management of SARS-CoV-2 infection at our hospital. A 50-year-old male patient operated on for squamous cell carcinoma of the oral cavity who completed his adjuvant chemoradiation 2 months ago, presented to our hospital with dry cough for more than 3 weeks. His nasopharyngeal swab was positive for the severe acute respiratory distress syndrome (SARS-CoV-2). During his hospital stay for SARS-CoV-2 infection, he was diagnosed with disseminated penicilliosis. The patient was treated with intravenous antifungals caspofungin and voriconazole. However, he succumbed to disseminated fungal sepsis. This case highlights the need to consider penicilliosis as a possible opportunistic pathogen, especially in immunocompromised patients such as cancer.

Keywords
► penicilliosis
► cancer
► oral cavity
► SARS-CoV-2
► COVID-19

Introduction
Penicilliosis infection is commonly seen in immunocompromised patients such as acquired immunodeficiency syndrome (AIDS), cancer and persons with autoimmune diseases.1-3 Disseminated penicilliosis is transmitted in humans from the inhalation of spores from the environment.4 There are published reports in the literature showing penicilliosis infections in lung cancer patients.5 In 1959, G. Segretain, a mycologist from the Pasteur Institute at Paris was the first person to discover penicilliosis. Symptomatology varies as per the system involved. The diagnosis of penicilliosis is suspected or done through examination of either cytology or biopsy samples.6 Herein, we report the first case...
of disseminated penicilliosis infection in a cancer patient with concomitant SARS-CoV-2 infection.

Case

A 50-year male patient, premorbidly healthy, was diagnosed with carcinoma of buccal mucosa (non-keratinizing squamous cell carcinoma) T4-N3b-M0. He received two cycles of neo-adjuvant chemotherapy (paclitaxel + carboplatin + cetuximab) followed by right buccal mucosa composite resection + left neck dissection followed by pectoralis major myocutaneous flap. Post-surgery, he received concurrent radiation (60Gy/30# of RT over 43 days) along with weekly cisplatin. Two months after completion of concurrent chemoradiation, he presented with a 3-week history of dry cough. Nasopharyngeal swab was positive for SARS-CoV-2 infection using standard real-time polymerase chain reaction (RT-PCR test). He was hospitalized with a diagnosis of severe COVID-19 as per ordinal scale. On admission, the patient was tachypneic, and hypoxic with bilateral crepitations. X-ray chest (Fig. 1A) showed bilateral infiltrates. In view of severe COVID-19, and in anticipation of a difficult airway considering a recent history of major facial surgery, an elective tracheostomy was performed on the day of hospital admission. After tracheostomy, he was supported with invasive mechanical ventilation with volume control mode (FiO₂ 50%). Meanwhile, he was treated with intravenous dexamethasone (6 mg once a day as per the RECOVERY trial) for severe COVID-19, along with broad-spectrum antibiotic (meropenem) and prophylactic enoxaparin. Non-directed broncho-alveolar lavage (ND-BAL) was sent for bacterial culture that showed a sensitive strain of Pseudomonas aeruginosa. Over the next 96 hours, he improved with respect to defervescence, oxygen requirement, and lung infiltrates. On day 10, he was on minimal oxygen support (2 L/min) without any tachypnea. Repeat ND-BAL cultures were negative for any bacterial growth. However, on day 12, we noticed an increase in respiratory rate, with a rise in C-reactive protein (CRP) and radiological evidence of increased lung infiltrates (Fig. 1B). In view of normal procalcitonin, high CRP, and repeatedly negative blood and tracheal cultures, we suspected an impending recurrent cytokine storm due to SARS-CoV-2 infection. On account of better lung penetration of methylprednisolone in comparison to dexamethasone and based on evidence for utility of bevacizumab at the time of patient management, he was treated with a single dose of intravenous bevacizumab 400 mg, intravenous methylprednisolone 20 mg 12 hourly, along with broad-spectrum antibiotics (colistin + ceftazidime-avibactam + aztreonam). However, his lung infiltrates continued to worsen. In view of rising CRP with worsening infiltrates, recent steroid use, and recent use of broad-spectrum antibiotics, the possibility of an opportunistic fungal infection was considered. His blood sugars and neutrophil counts were normal. An ND-BAL fungal culture was repeated. After 24 hours on Sabouraud dextrose agar (SDA) at 25°C; it showed a filamentous grayish–white growth in 24 hours. The growth later became yellowish green in the center surrounded by white periphery and had radial folds. On Lacto phenol Cotton Blue Mount (LPCB), penicillillar morphology was seen. Hyaline short hyphae that were septate and branched bearing chains of elliptical or lemon shaped phialoconidia were seen. Dimorphism showing yeast to mycelial phase conversion could not be demonstrated. Hence, it was probably diagnosed as Penicillium non-marneffei spp. Voriconazole plus caspofungin has been shown to work well for patients with penicilliosis non-marneffei infection. Hence, he was started on injection caspofungin pre-emptively pending further identification. Computerized tomogram (CT) of the chest and abdomen showed multiple nodular lesions in bilateral lung fields with cavitations, ground glass opacities, along with well-defined hypodense lesions in segment VII, VIII and multiple satellite foci consistent with a fungal etiology (Fig. 1C). An aspiration of liver abscess was not performed in view of poor general condition of the patient. There were no signs of active COVID-19 disease on CT scan. Patient was continued on antibacterials and injection caspofungin. Over the next 96 hours, ND-BAL culture growth was reported as Penicillium species (non-marneffei) (Fig. 2). ND-BAL galactomannan levels were

Fig. 1 (A) X-ray chest showing a few nodular opacities in both lung fields, maximum in right lower zone suggestive of infective etiology. (B) X-ray chest showing increase in nodular opacities as compared to a prior X ray, features suggestive of worsening of pulmonary infection. (C) Multinodular lesions in bilateral lung fields with cavitations (red arrow) and surrounding ground glass opacities (green arrow) [in top two panels and left middle panel] suggestive of infective etiology more likely to be fungal. Multiple hypodense (blue arrow) hepatic lesions in segments VII and VIII, suggestive of multiple abscesses. [right middle panel and both lower panels].
positive (Index 3.601), though serum galactomannan levels were negative. The day wise trend of CBC, CRP, and procalcitonin is provided in Table 1. After 1 week, voriconazole was added as he continued to worsen with respect to respiratory parameters, rising CRP, worsening in infiltrates, and negative bacterial cultures. Subsequently, he had worsening hypoxia necessitating mechanical ventilation. Despite maintaining therapeutic voriconazole levels, he progressed to septic shock. On day 27 of hospital admission, he succumbed to disseminated penicilliosis.

Discussion
Penicillium species (especially *P. marneffei*) have been proven to cause fatal invasive infections in immunocompromised cases. Pulmonary fungal infections with *Penicillium non-marneffei* species have been reported in immunocompromised hosts. Disseminated disease with hepatosplenic involvement and pericardial involvement are reported. Invasive fungal infection can be attributed to some of the rare *Penicillium* species such as *P. commune*, *P. purpurogenum*, *P. oxalicum*, *P. notatum*, *P. citrinum*, and *P. brevicompactum*. Definitive diagnosis of *Penicillium* infection is made by culture of biological specimens with a turn-around time of approximately a week or more. Although no serological tests are developed for diagnosing penicilliosis, galactomannan assay can show cross-reactivity with *Penicillium* species and can pre-empt diagnosis of invasive penicilliosis before microbiological cultures.

In retrospect, looking back to the SARS epidemic in 2003, fungal infection was the most common cause of mortality accounting for 25 to 73% of all deaths. However, in contrast, data of fungal co-infection in COVID-19 are scarce. A meta-analysis by Peng et al reported an overall pooled proportion of COVID-19 with fungal co-infection of 0.12%, with an overall mortality of 0.17%. They described a higher proportion of patients with co-infection from Asia, in comparison to Europe. However, the majority of their fungal co-infections were caused by *Aspergillus*. Possible risk factors for the

**Table 1** Day-wise trend of hematological and biochemistry report of the patient

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<th>Day of COVID-19 positivity</th>
<th>Hb (g/dL)</th>
<th>TLC $\times 10^9$ (per L)</th>
<th>ANC $\times 10^9$ (per L)</th>
<th>ALC $\times 10^9$ (per L)</th>
<th>Platelets $\times 10^9$ (per L)</th>
<th>CRP (mg%)</th>
<th>Procalcitonin (ng/mL)</th>
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Fig. 2 Lactophenol cotton blue stain mount from the growth of the *Penicillium* spp showing conidiophores having secondary branches called metulae. On metulae are arranged flask-shaped phialides bearing conidia. The entire structure is forming the characteristic ‘penicillus’ or ’brush’ appearance.
development of this invasive fungal infection in our patient were the presence of an underlying malignancy, use of steroids and broad-spectrum antibiotics. Similar to literature, we treated our patient with voriconazole, albeit without success. As per the literature on talaromycosis in cancer patients, cases have been reported in lung cancer patients with favorable and poor outcomes. However, in the former case, there was a relapse of the fungal infection, 5 months after initial treatment. We also found one case report of nodular Hodgkin’s lymphoma patient and an ovarian cancer patient infected with talaromycosis. In the latter case, Penicillium spp. was resistant to all agents, except amphotericin-B. It is important to note that routine anti-fungal susceptibility, except for Candida is not routinely available in India. While we administered voriconazole to our patient, he succumbed despite maintaining therapeutic drug levels. While both COVID-19 and penicilliosis predominantly affect the respiratory system, the mortality of penicilliosis is much higher than the case fatality in COVID-19.

**Conclusion**

To our knowledge, this is the first reported case of *Penicillium* co-infection with COVID-19 in the oral cavity cancer patient. While worsening respiratory symptoms in a COVID-19 patient are commonly attributed to SARS-CoV-2 or a bacterial superinfection, physicians should be cognizant for the possibility of a concomitant fungal infection which can be fatal.

**Declaration of Patient Consent**

The authors certify that they have obtained all appropriate patient consent forms. In the form, the patient has given his consent for his images and other clinical information to be reported in the journal. The patient understands that his name and initials will not be published and due efforts will be made to conceal his identity, but anonymity cannot be guaranteed.

**Funding**

None.

**Conflict of Interest**

None declared.

**References**


