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The First Case of Preauricular Fistular Abscess Caused by *Peptoniphilus grossensis*

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Dear Editor,

The genus *Peptoniphilus* comprises obligatory anaerobic grampositive cocci. The name *Peptoniphilus* means "affinity to peptones," which are the major energy source of this genus [1]. More than 28 *Peptoniphilus* species have been isolated from the gut microbiome, and most of them are commensal organisms; *Peptoniphilus grossensis* was first isolated from human feces [2]. There are no reports on human diseases caused by *P. grossensis*.

A preauricular fistula or sinus is a common congenital abnormality, with a prevalence of approximately 2% in the Korean population and 0.1%–10% in several other countries [3]. In some cases, it can get infected. We report the first case of a preauricular fistular abscess caused by *P. grossensis* in a human. This study was approved by The Institutional Review Board of Gyeongsang National University Hospital, Jinju, Korea approved the study (approval number 202108013) and waived the need for informed consent.

A 23-year-old woman was admitted for preauricular pain, swelling, and a febrile sensation that started 10 days earlier. Two years prior, she had suffered an *Actinomyces odontolyticus*-infected preauricular fistula. The remainder of her history was unremarkable. She was suspected of suffering recurrent preauric

ular fistula infections. She had an elevated C-reactive protein level (21.5 mg/L; reference range <5 mg/L) and erythrocyte sedimentation rate (57 mm/hr; \leq 20 mg/L). Two closed pus specimens were obtained through incision and drainage. Microscopic examination using Gram's stain revealed monomorphic gram-positive cocci, numerous white blood cells (WBCs), and 1–5 epithelial cells per high-power field (Fig. 1). Three days later, anaerobic gram-positive cocci were isolated from a thioglycollate broth culture. They were subcultured on Brucella agar plates and incubated in an anaerobic chamber. The bacteria grew slowly, and a pinpoint colony formed after two days (Fig. 1). The colony was non-hemolytic, grayish, and circular. The bacteria were small gram-positive cocci.

The bacteria could not be identified by matrix-associated laser desorption ionization-time of flight mass spectrometry (bioMérieux, Marcy l'Étoile, France). Biochemical testing using the VI-TEK 2 system (bioMérieux) identified the bacteria as *P. asac-charolyticus*, with a 95% probability. 16S rRNA sequencing was performed to identify the bacterial species. According to the Ez-Taxon database (v2.1; http://www.ezbiocloud.net), the bacteria were identical to *P. grossensis* (100%), followed by *P. senega-lensis* (99.42%), *P. lacydonensis*, and *P. rhinitidis* (99.14%). As 16S rRNA sequencing did not provide resolution at the species

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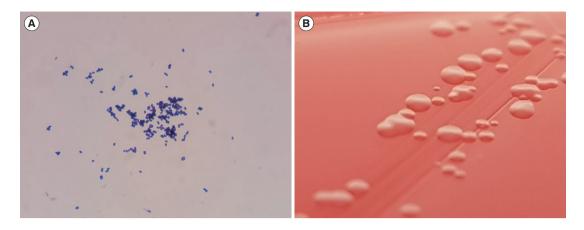


Fig. 1. Microscopic image and colony morphology of *Peptoniphilus grossensis*. (A) Gram-positive cocci (Gram stain, ×1,000). (B) Pinpoint and grayish colonies (×100) on a Brucella agar plate after 48 hours incubation.

level, whole-genome sequencing using the Illumina MiSeq platform (Illumina, San Diego, CA, USA) was performed. The genome size of the pathogen was 2,038,961 bp, and the data were analyzed using the TrueBac ID-Genome system (www.truebacid.com; ChunLab, Seoul, Korea) [4]. Based on the average nucleotide identity, the bacteria were classified under the genus *Peptoniphilus*. The bacteria were identified as *P. grossensis* (96. 25%), *P. senegalensis* (88.60%), *P. lacydonensis* (85.74%), and *P. rhinitidis* (85.83%). Beta-lactamase screening was negative.

Intravenous ampicillin/sulbactam was administered for four days after initial incision and drainage. The patient was discharged as no bleeding or purulent discharge from the wound was observed. An oral cefditoren, a third-generation cephalosporin, was prescribed for seven days for preauricular pain at the time of discharge.

Gram-positive anaerobic cocci are mostly commensal and opportunistic pathogens [5]. Infections from these bacteria are often polymicrobial [6]. Therefore, the clinical significance of individual anaerobic cocci is difficult to determine in clinical practice. Herein, monomorphic gram-positive cocci and numerous WBCs were observed on a direct smear of closed pus obtained aseptically. A single species was recovered, indicating that this bacterium caused the infection and highly likely is pathogenic.

Peptoniphilus are fastidious bacteria and identifying them clinically is time-consuming. *P. grossensis* was isolated but not identified within a clinically significant timeframe. Antibiotic susceptibility testing of anaerobic bacteria is not feasible in most clinical settings, because it requires manual agar dilution or broth microdilution with supplementation of hemin, vitamin K, and laked sheep or horse blood [7]. We identified three antibiotic-resistance genes in the isolate genome: *erm (A)*, which confers

resistance to erythromycin, and *aph(3')-IIIa and ant(6)-Ia*, which confer resistance to aminoglycosides. We did not assess phenotypic antibiotic susceptibility, because the clinical breakpoints for macrolides and aminoglycosides are not provided in the CLSI or European Committee on Antimicrobial Susceptibility Testing guidelines [7, 8]. Anaerobic gram-positive cocci, including *Pept-oniphilus* spp., do not exhibit high-level resistance against other antibiotics [6, 9]. After incision, drainage, and beta-lactam combination drug treatment as empirical therapy, the prognosis of the patient was good.

This report highlights the potential pathogenicity of *P. grossensis*, even in immunocompetent young subjects. Technological advances can help detect and identify more bacteria in human specimens. Further investigation of the pathogenesis and clinical significance of rare bacteria such as *P. grossensis* is needed [5, 10].

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AUTHOR CONTRIBUTIONS

Conceptualization: Byun JH. Data curation: Kim M, Shin H, Lee DH, Koh EH. Writing-review and editing: Byun JH, Kim M. Final approval of the manuscript: all authors.

CONFLICTS OF INTEREST

The authors have no conflicts of interest to declare.

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