# Case Series

# Correspondence Simon Lévesque simon.levesque@inspq.qc.ca

Received 16 July 2015
Revised 22 December 2015
Accepted 26 January 2016

# Enteroccocus pallens as a potential novel human pathogen: three cases of spontaneous bacterial peritonitis

Simon Lévesque,<sup>1</sup> Yves Longtin,<sup>2</sup> Marc-Christian Domingo,<sup>1</sup> Cynthia Massé,<sup>1</sup> Harold Bernatchez,<sup>3</sup> Christiane Gaudreau<sup>4</sup> and Cécile Tremblay<sup>1,5</sup>

**Introduction:** *Enterococcus pallens* is one of the four yellow-pigmented members of the genus *Enterococcus*. To date, a single report of *E. pallens* isolated from a human sample has been published.

**Case presentation:** We report three cases of *E. pallens* spontaneous bacterial peritonitis in patients with liver cirrhosis that all occurred in Quebec, Canada. Ascitic fluid analysis revealed the presence of *E. pallens* in culture. Identification was made by classical biochemical testing and MALDI-TOF MS, as well as 16S rRNA and elongation factor (*tut*) gene sequencing. Two of the three patients recovered after antimicrobial treatment.

**Conclusion:** This report identifies *E. pallens* as a novel human pathogen that appears to possess particular but as-yet unidentified virulence factors that favour the development of peritoneal fluid infections, as previously reported for other *Enterococcus* species. Clinical microbiologist should be aware of this micro-organism which can be identified by phenotypic and molecular methods.

Keywords: Enteroccocus pallens; liver cirrhosis; peritonitis; Quebec.

#### Introduction

Enterococcus pallens is one of the four yellow-pigmented Enterococcus. To date, a single report of E. pallens isolated from a human sample has been published (Tyrrell et al., 2002). The type strain (ATCC BAA-351) was obtained from a peritoneal dialysate of a patient from Quebec, Canada, who developed peritonitis secondary to intestinal perforation. Here, we report the first case series of human infections caused by this pathogen.

Abbreviations: LSPQ, Laboratoire de santé publique du Québec; NASH, non-alcoholic steatohepatitis; SBP, spontaneous bacterial peritonitis.

The GenBank/EMBL/DDBJ accession numbers for the 16S rRNA and *tuf* gene sequences of the isolates LSPQ-04155, LSPQ-04156, LSPQ-04157 and LSPQ-04217 are KP793151 and KP793155, KP793152 and KP793156, KP793153 and KP793158, and KP793154 and KP793157, respectively.

## **Case Reports**

Patient 1 was a 72-year-old woman known for cirrhosis secondary to non-alcoholic steatohepatitis (NASH) who was admitted in March 2004 to a hospital in Montreal, Canada, because of a 4 month history of increasing abdominal girth, progressive dyspnoea and exacerbation of leg oedema. Her medical history also included coronary artery disease, hypertension, type 2 diabetes mellitus and nephrolithiasis. On examination, the patient was icteric. Her vital signs were normal. She was alert and oriented. The abdomen was markedly distended but non-tender. Laboratory tests revealed a white cell count of  $6.2 \times 10^9$ cells l<sup>-1</sup>. Ultrasonography of the abdomen revealed an echogenic liver, a patent portal vein and moderate ascites. A diagnosis of hydrops secondary to NASH-related liver failure was made, and a 1.5 l day<sup>-1</sup> fluid restriction and a diuretic therapy were initiated. On day 14 of hospitalization, the patient complained of mild abdominal pain.

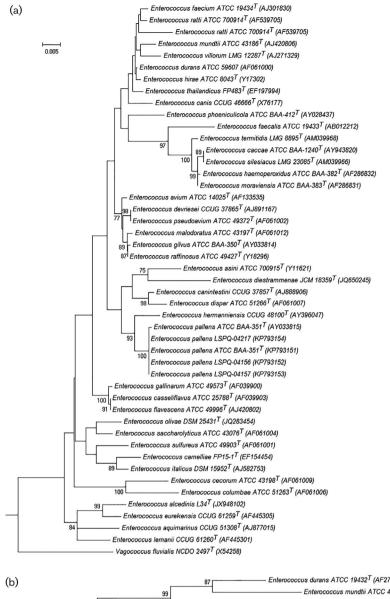
<sup>&</sup>lt;sup>1</sup>Laboratoire de santé publique du Québec/Institut national de santé publique du Québec, Sainte-Anne-de-Bellevue, Québec, Canada

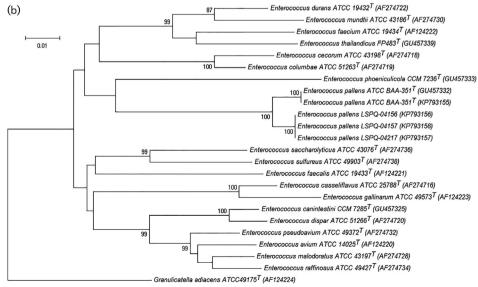
<sup>&</sup>lt;sup>2</sup>SMBD - Jewish General Hospital, Québec, Canada

<sup>&</sup>lt;sup>3</sup>Centre hospitalier régional Rimouski-Neigette, Québec, Canada

<sup>&</sup>lt;sup>4</sup>Centre Hospitalier de l'Université de Montréal, Québec, Canada

<sup>&</sup>lt;sup>5</sup>Centre de recherche de l'Université de Montréal, Québec, Canada





2 JMM Case Reports

**Fig. 1.** Neighbour-joining phylogenetic tree showing the relationships of the three patient's isolates to related *Enterococcus* species. Bootstrap values were calculated from 1000 replications. Names and GenBank accession numbers are given. Relatedness between (a) partial sequences (1181 bp) of the 16S rRNA gene and (b) partial sequences (599 bp) of the *tuf* gene. GenBank accession numbers for 16S rRNA gene sequences: LSPQ-04155 (KP793151), LSPQ-04156 (KP793152), LSPQ-04157 (KP793153) and LSPQ-04217 (KP793154). GenBank accession numbers for *tuf* gene sequences: LSPQ-04155 (KP793155), LSPQ-04156 (KP793156), LSPQ-04157 (KP793158) and LSPQ-04217 (KP793157). Bar, estimated number of substitutions per site.

Abdominal paracentesis was performed to improve her symptoms; 3 l of straw-coloured fluid were drained. Ascitic fluid analysis revealed a red blood cell count of  $103 \times 10^6$ cells  $l^{-1}$ , a leukocyte count of  $62 \times 10^6$  cells  $l^{-1}$  and an absence of malignant cells. A differential leukocyte count was not performed. No organism was seen on Gram staining of the fluid, but E. pallens was isolated on culture. The patient developed an episode of fever (rectal temperature, 39.1 °C) 1 h after the procedure for which antibiotic therapy consisting of intravenous ticarcillin and clavulanic acid was initiated. Blood cultures were negative. On day 16 of hospitalization, the patient complained of shortness of breath and became hypotensive (systolic pressure 95 mmHg). Abdominal paracentesis was repeated; 2.61 of cloudy yellowish fluid was removed. The ascitic fluid leukocyte count was  $600 \times 10^6$  cells  $1^{-1}$  with >70 %neutrophils. Culture of peritoneal fluid once again yielded E. pallens. The patient fully recovered after a 7 day course of intravenous ticarcillin and clavulanic acid.

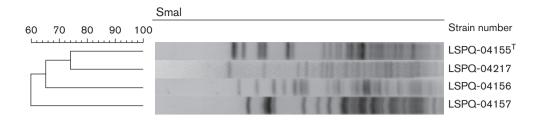
Patient 2 was a 62-year-old male known for liver cirrhosis and refractory ascites requiring drainage every other week. He was admitted in May 2013 to a hospital in Rimouski, Canada, due to abdominal pain, increasing abdominal girth and fever. A presumptive diagnosis of spontaneous bacterial peritonitis (SBP) was made and antibiotic therapy consisting of intravenous ceftriaxone was initiated. His medical history also included chronic partial thrombosis of the portal vein, thrombocytopenia, coagulopathy, chronic kidney disease and dilated cardiomyopathy. On the second day of hospitalization, an abdominal paracentesis was performed. Ascitic fluid analysis revealed a white blood cell count of  $1611 \times 10^6$  cells  $1^{-1}$  with 91 % neutrophils. Direct examination of the ascitic fluid showed no micro-organisms, but E. pallens was isolated on culture. Antimicrobial treatment was switched to imipenem and the patient fully recovered.

Patient 3 was a 62-year-old man with alcoholic liver cirrhosis and portal hypertension who was admitted in July 2013 to a hospital in Montreal, Canada, because of hepatic encephalopathy. He was known for refractory ascites requiring large-volume paracentesis twice monthly. His medical history also included a chronic excoriated umbilical hernia, previous hepatic encephalopathy and lower leg lymphoedema. Ascitic fluid analysis revealed a white blood cell count of  $447 \times 10^6$  cells l<sup>-1</sup> with 62 % neutrophils. A diagnosis of SBP was made and antibiotic therapy consisting of intravenous piperacillin/tazobactam was initiated. *E. pallens* was isolated from the ascitic fluid

culture, which prompted the addition of intravenous vancomycin on days 4 and 5 of hospitalization. Blood and urine cultures were negative. Chest radiography showed no consolidation or infiltrate. Repeat abdominal paracentesis was performed on day 3 of hospitalization which showed  $<\!250\times10^6$  neutrophils  $l^{-1}$  and a negative ascitic fluid culture. A third abdominal paracentesis was performed on day 5, which this time showed a very high neutrophil count  $(2725\times10^6~{\rm cells~l^{-1}})$  and yielded *Escherichia coli*. The patient died on day 8 due to global deterioration.

# **Investigations**

The three isolates were sent to the Laboratoire de santé publique du Québec (LSPQ) for identification. Classical biochemical testing, as well as 16S rRNA (Bekal et al., 2006) and elongation factor (tuf) gene sequencing were performed. For tuf gene analysis, new primers were designed in order to amplify a 619 bp fragment of the gene from all the species of the genus Enterococcus (forward primer EntA1 5'-ATCTTAGTAGTTTCTGCTGCTGA, reverse primer EntA2 5'-GTAGAATTCAGGACGGTAGT-TAG). The same primers were used for sequencing. Fig. 1 shows the phylogenetic trees of 16S rRNA and tuf gene sequences reconstructed with MEGA5 software (Tamura et al., 2011). Multiple sequence alignment using the neighbour-joining method showed that isolates LSPO-04155 (i.e. the type strain ATCC BAA-351), LSPO-04156, LSPQ-04157 and LSPQ-04217 were closely related to each other, displaying 100 and 98 % similarity for 16S rRNA and tuf, respectively. Interestingly, the biochemical characteristics of the three isolates were identical to those of the type strain, with the exception of methyl-α-glucopyranoside which was positive, whilst the type strain was negative. The three isolates could not be identified using VITEK MS MALDI-TOF. The species E. pallens was not included in the VITEK MS IVD version 2.0 (bioMérieux), a clinically relevant species database, and SARAMIS version 4.1, the Research Use Only database. New spectra entries were added to the SARAMIS database using cultures from different times of incubation for each of the three isolates and also from the reference strain LSPO-04155 (ATCC BAA-351). Based on this newly updated library, the SARAMIS database was able to identify the three isolates correctly to species level. The Bruker Biotyper database (Bruker Daltonics) contains the species *E. pallens*. The three strains were correctly identified with the Bruker Biotyper system. PFGE (Gagnon et al., 2011) was performed and showed that all isolates had different PFGE



**Fig. 2.** Smal PGFE patterns of three *E. pallens* isolates compared with the reference strain LSPQ-04155 (ATCC BAA-351), identified with the letter 'T'. The scale represents the percentage similarity between the isolates.

patterns (Fig. 2). Susceptibility profiles to seven antimicrobials were obtained by broth microdilution according to Clinical and Laboratory Standards Institute guidelines (CLSI, 2014). All isolates were susceptible to penicillin, ampicillin, vancomycin, teicoplanin, quinupristin/dalfopristin, daptomycin and linezolid.

#### Discussion

SBP is a common infection that affects patients with liver cirrhosis and ascites. The infection occurs following the translocation of gut microflora into the ascites and subsequent proliferation due to impaired host defence (Bennett et al., 2015). Unsurprisingly, Enterobacteriaceae, in particular E. coli and Klebsiella pneumoniae, account for the majority of SBP. Enterococcus species, Streptococcus pneumoniae and other Streptococcus spp. are also common causative agents (Bennett et al., 2015). The microbiology of SBP may be evolving in some regions. Reports of increasing prevalence of Gram-positive SBP in Turkey (Yakar et al., 2010) and of the emergence of Enterococcus species SBP in Germany and in Greece have been published (Alexopoulou et al., 2013; Reuken et al., 2012). Here, we report the first series of patients who developed SBP due to E. pallens. In humans, this micro-organism had previously been isolated only once from the peritoneal dialysate of a patient with secondary peritonitis in Quebec (Tyrrell et al., 2002). The fact that all four human infections described so far were peritonitis is noteworthy. Hence, E. pallens, as well as other Enterococcus species, appears to possess particular but as-yet unidentified virulence factors that favour the development of peritoneal fluid infections. Whether E. pallens preferentially colonizes the digestive tract of patients with cirrhosis or whether it colonizes humans regardless of underlying conditions but mainly causes infections in patients with cirrhosis remains to be determined. Also, the fact that all four E. pallens human infections reported to date in the literature originate from the province of Quebec raises the question of the geographical distribution of this micro-organism. However, as the four isolates analysed are not genetically related, the hypothesis of a single regional clone is unlikely. Further studies are needed to determine the prevalence of E. pallens and to better

understand its virulence factors. Clinical microbiologist should be aware of this micro-organism which can be identified by phenotypic and molecular methods. As *E. pallens* and other *Enterococcus* species are intrinsically resistant to cephalosporins, clinicians should suspect these pathogens in the context of failure of first-line empirical agents.

In conclusion, we describe the first case series of human infections due to *E. pallens*. Together with the only previous reported human case, these suggest that *E. pallens* has a propensity to cause peritoneal fluid infections.

## **Acknowledgements**

We thank Agata Klebucki and Nancy Cloutier from LSPQ for technical assistance. We thank Maureen Hastie for English revision. We also thank bioMérieux Canada for access to the VITEK MS MALDI-TOF system and Kathryn Bernard from the National Microbiology Laboratory at Winnipeg, Canada for Bruker Biotyper identification. This study was conducted with LSPQ internal funding. S. L. has a research agreement with bioMérieux Canada.

### References

Alexopoulou, A., Papadopoulos, N., Eliopoulos, D. G., Alexaki, A., Tsiriga, A., Toutouza, M. & Pectasides, D. (2013). Increasing frequency of gram-positive cocci and gram-negative multidrug-resistant bacteria in spontaneous bacterial peritonitis. *Liver Int* 33, 975–981.

Bekal, S., Gaudreau, C., Laurence, R. A., Simoneau, E. & Raynal, L. (2006). *Streptococcus pseudoporcinus* sp. nov., a novel species isolated from the genitourinary tract of women. *J Clin Microbiol* 44, 2584–2586.

Bennett, J. E., Dolin, R. & Blaser, M. J. (2015). Mandell, Douglas, and Bennett's Principles and Practice of Infectious Diseases, 8th edn. Philadelphia, PA: Churchill Livingstone Elsevier.

**CLSI (2014).** Performance Standards for Antimicrobial Susceptibility Testing; 24th Informational Supplement M100-S24. Wayne, PA: Clinical and Laboratory Standards Institute.

Gagnon, S., Lévesque, S., Lefebvre, B., Bourgault, A. M., Labbé, A. C. & Roger, M. (2011). *vanA*-containing *Enterococcus faecium* susceptible to vancomycin and teicoplanin because of major nucleotide deletions in Tn*1546*. *J Antimicrob Chemother* **66**, 2758–2762.

Reuken, P. A., Pletz, M. W., Baier, M., Pfister, W., Stallmach, A. & Bruns, T. (2012). Emergence of spontaneous bacterial peritonitis due to enterococci – risk factors and outcome in a 12-year retrospective study. *Aliment Pharmacol Ther* 35, 1199–1208.

4 JMM Case Reports

Tamura, K., Peterson, D., Peterson, N., Stecher, G., Nei, M. & Kumar, S. (2011). MEGA5: molecular evolutionary genetics analysis using maximum likelihood, evolutionary distance, and maximum parsimony methods. *Mol Biol Evol* 28, 2731–2739.

Tyrrell, G. J., Turnbull, L., Teixeira, L. M., Lefebvre, J., Carvalho, M. G., Facklam, R. R. & Lovgren, M. (2002). *Enterococcus gilvus* sp. nov. and

Enterococcus pallens sp. nov. isolated from human clinical specimens. *J Clin Microbiol* **40**, 1140–1145.

Yakar, T., Güçlü, M., Serin, E. & Alişkan, H (2010). A recent evaluation of empirical cephalosporin treatment and antibiotic resistance of changing bacterial profiles in spontaneous bacterial peritonitis. *Dig Dis Sci* 55, 1149–1154.