

First detection of insertion sequence element ISPa1328 in the *oprD* porin gene of an imipenem-resistant *Pseudomonas aeruginosa* isolate from an idiopathic pulmonary fibrosis patient in Marseille, France

C. Al-Bayssari¹, C. Valentini¹, C. Gomez²,
M. Reynaud-Gaubert² and J.-M. Rolain¹

1) Aix-Marseille Université, Unité de Recherche sur les Maladies Infectieuses et Tropicales Emergentes (URMITE), UM63, CNRS 7278, IRD 198, Inserm 1095, IHU Méditerranée Infection, Faculté de Médecine et de Pharmacie and 2) Service de Pneumologie, Centre de Soins de la Mucoviscidose (CRCM) et en Transplantation Pulmonaire Adulte, Hôpital Nord, Marseille, France

Abstract

We report here the first case of a carbapenem-resistant *Pseudomonas aeruginosa* clinical isolate harboring the insertion sequence (IS) element ISPa1328 in the *oprD* gene in an idiopathic pulmonary fibrosis patient in France previously treated with imipenem.

New Microbes and New Infections © 2015 The Authors. Published by Elsevier Ltd on behalf of European Society of Clinical Microbiology and Infectious Diseases.

Keywords: Carbapenem resistance, insertion sequence, *oprD* gene, *Pseudomonas aeruginosa*, pulmonary fibrosis

Original Submission: 24 February 2015; **Accepted:** 6 May 2015
Available online 14 May 2015

Corresponding author: J.-M. Rolain, Aix-Marseille Université, Unité de Recherche sur les Maladies Infectieuses et Tropicales Emergentes (URMITE), UM63, CNRS 7278, IRD 198, Inserm 1095, IHU Méditerranée Infection, Faculté de Médecine et de Pharmacie, 27 Bd Jean Moulin 13005 Marseille, France
E-mail: jean-marc.rolain@univ-amu.fr
The first two authors contributed equally to this article, and both should be considered first author

Carbapenems are frequently used as last drug choice for the treatment of multidrug-resistant *Pseudomonas aeruginosa* infections [1], but the emergence of carbapenem resistance is increasingly reported. Carbapenem resistance in *P. aeruginosa* may be due to low permeability, multidrug efflux pumps or the production of class B β -lactamases (metallo- β -lactamases, MBLs) [2]. However, the main mechanism of carbapenem resistance in *P. aeruginosa* remains the loss or the alteration of the outer membrane porin (*oprD*) through mutations, deletions or insertions in the *oprD* gene [3]. Clinical isolate of *P. aeruginosa* exhibiting high-level resistance to carbapenems was isolated from a sputum sample of a 69-year-old man with idiopathic pulmonary fibrosis treated by the association of either ciprofloxacin or ceftazidime or tazocillin with tobramycin or colimycin and later by imipenem. Strain (PA 461) was cultured on trypticase soy agar plate at 37°C for 24 hours, and identification was confirmed by matrix-assisted desorption ionization–time of flight mass spectrometry (MALDI-TOF MS) (Microflex; Bruker Daltonics, Bremen, Germany) with FlexControl software, as previously described [4]. Antibiotic susceptibility testing performed on Mueller-Hinton agar by standard disk diffusion method as recommended by the European Committee on Antimicrobial Susceptibility Testing (EUCAST; <http://www.eucast.org/>), showed that the isolate was resistant to almost all antibiotics, including β -lactams, as well as fluoroquinolones and rifampicin, but remained susceptible to aminoglycosides and colistin. Phenotypic detection of carbapenemase by modified Hodge test and imipenem–ethylene diamine tetra-acetic acid test, performed as previously described [5], were negative. The presence of MBLs genes, investigated by PCR as previously described [6], confirmed that this isolate did not produce a carbapenemase. Amplification of the *oprD* gene using previously described primers [3,6] resulted in PCR product of 2.6 kb instead of 1332 bp. Sequence analysis of the 2668 bp product revealed insertion of a sequence of 1336 bp at position 610 in *oprD*. Within the sequence, an open reading frame of 1227 bp was found and corresponded to ISPa1328 from *P. aeruginosa* (GenBank accession AY539833). This insertion sequence (IS) was bordered by two terminal imperfect repeats and flanked on both sides by direct repeat sequences of 7 bp (CCAAGAG) (Fig. 1). Multilocus sequence typing, performed as previously described (<http://pubmlst.org/paeruginosa>), showed a novel sequence type (ST1797) and thus a novel clone of *P. aeruginosa*.

Random transposition of IS elements is known to be a form of adaptation of bacteria to environmental changes. To date, the presence of IS elements disrupting the *oprD* gene has been reported in South Africa (ISPa26) [7], Spain (ISPa133) [8], China (ISPa1328, ISPre2) [9], the United States (ISPa8 [1] and ISPa1328 [10]) and France (ISPa46) [6]. The occurrence of multidrug-

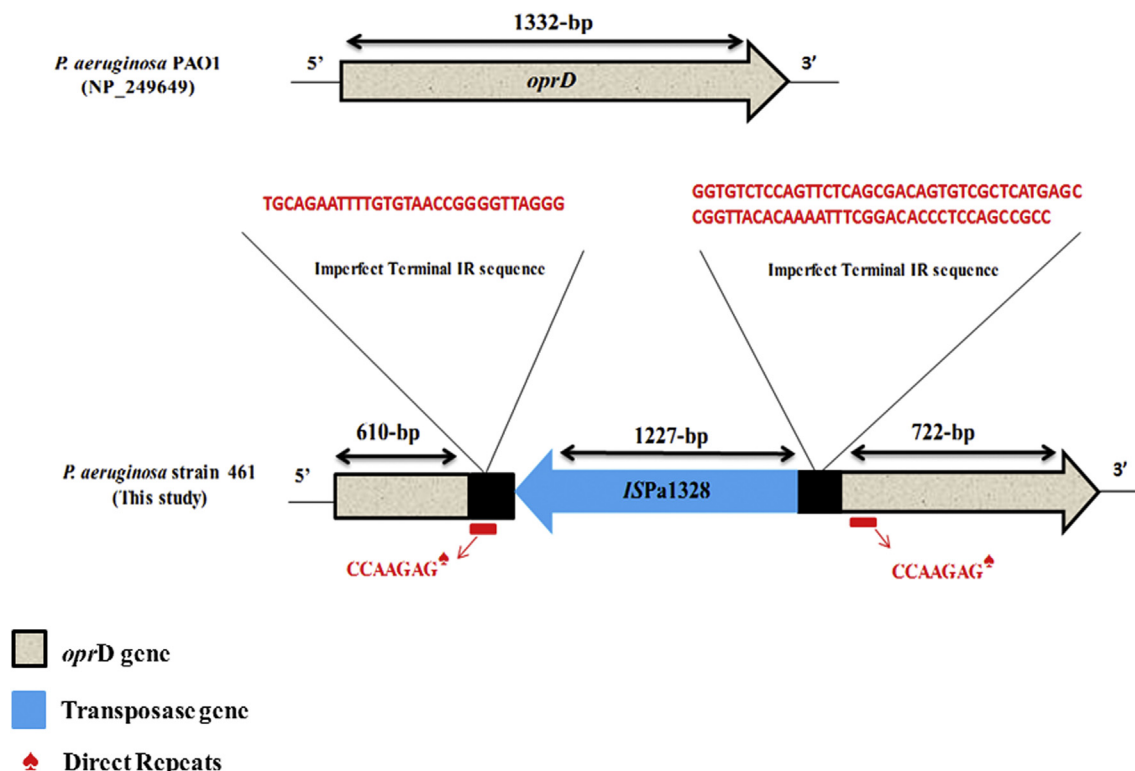


FIG. 1. Schematic representation of *oprD* gene of PA461 disrupted by ISPa1328 compared to reference strain PAO1.

resistant bacteria is associated with the extensive use of broad-spectrum antimicrobial drugs in treating humans. This could explain the emergence of imipenem-resistant *P. aeruginosa* in this patient, the result of direct and specific antibiotic selective pressure created by the use of imipenem.

In conclusion, we report for the first time in France the emergence of ISPa1328 in a patient with idiopathic pulmonary fibrosis associated with carbapenem resistance.

Conflict of interest

None declared.

Acknowledgement

The authors thank L. Hadjadj for technical assistance.

References

[1] Fowler RC, Hanson ND. Emergence of carbapenem resistance due to the novel insertion sequence ISPa8 in *Pseudomonas aeruginosa*. *PLoS One* 2014;9:e91299.

[2] Walsh TR, Toleman MA, Poirel L, Nordmann P. Metallo-beta-lactamases: the quiet before the storm? *Clin Microbiol Rev* 2005;18:306–25.
 [3] Al Bayssari C, Diene SM, Loucif L, Gupta SK, Dabboussi F, Mallat H, et al. Emergence of VIM-2 and IMP-15 carbapenemases and inactivation of *oprD* gene in carbapenem-resistant *Pseudomonas aeruginosa* clinical isolates from Lebanon. *Antimicrob Agents Chemother* 2014;58:4966–70.
 [4] Seng P, Rolain JM, Fournier PE, La SB, Drancourt M, Raoult D. MALDI-TOF-mass spectrometry applications in clinical microbiology. *Future Microbiol* 2010;5:1733–54.
 [5] Lee K, Lim YS, Yong D, Yum JH, Chong Y. Evaluation of the Hodge test and the imipenem-EDTA double-disk synergy test for differentiating metallo-beta-lactamase-producing isolates of *Pseudomonas* spp. and *Acinetobacter* spp. *J Clin Microbiol* 2003;41:4623–9.
 [6] Diene SM, L'homme T, Bellulo S, Stremmer N, Dubus JC, Mely L, et al. ISPa46, a novel insertion sequence in the *oprD* porin gene of an imipenem-resistant *Pseudomonas aeruginosa* isolate from a cystic fibrosis patient in Marseille, France. *Int J Antimicrob Agents* 2013 Sep;42(3):268–71.
 [7] Evans JC, Segal H. A novel insertion sequence, ISPA26, in *oprD* of *Pseudomonas aeruginosa* is associated with carbapenem resistance. *Antimicrob Agents Chemother* 2007;51:3776–7.
 [8] Ruiz-Martinez L, Lopez-Jimenez L, d'Ostuni V, Fuste E, Vinuesa T, Vinas M. A mechanism of carbapenem resistance due to a new insertion element (ISPa133) in *Pseudomonas aeruginosa*. *Int Microbiol* 2011;14:51–8.
 [9] Wang J, Zhou JY, Qu TT, Shen P, Wei ZQ, Yu YS, et al. Molecular epidemiology and mechanisms of carbapenem resistance in *Pseudomonas aeruginosa* isolates from Chinese hospitals. *Int J Antimicrob Agents* 2010;35:486–91.
 [10] Wolter DJ, Acquazzino D, Goering RV, Sammut P, Khalaf N, Hanson ND. Emergence of carbapenem resistance in *Pseudomonas aeruginosa* isolates from a patient with cystic fibrosis in the absence of carbapenem therapy. *Clin Infect Dis* 2008;46:e137–41.