## Brief Original Article

# Extensively drug-resistant *Acinetobacter baumannii* isolated in a university hospital: Role of inter-hospital transmission

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#### Abstract

Introduction: *Acinetobacter baumannii* causes severe infections that primarily affect intensive care unit (ICU) patients. It has a high prevalence of multidrug resistance, including carbapenems, and a high potential for intra-hospital and inter-hospital transmission. The aim of this study was to determine the origin of extensively drug-resistant (XDR) *A. baumannii* isolates in our hospital during 2009.

Methodology: This was an observational retrospective study. Isolates of *A. baumannii* were obtained from patients hospitalized during 2009. XDR isolates were defined using criteria published by Magiorakos *et al.*. The isolates were classified as community acquired, hospital acquired, and inter-hospital transmission.

Results: A total of 48 isolates of *A. baumannii* were isolated during 2009, corresponding to 34 patients. Of these, 18 (53%) were susceptible, 6 (18%) were multidrug resistant (MDR), and 10 (29%) were XDR. Of the 10 XDR isolates, 9 were isolated from patients transferred from other hospitals. The median time of hospitalization in origin hospitals was 17 days, while the median time of hospitalization in the study hospital, previous to isolation of *A. baumannii*, was 1 day. A total of 6 out of 10 patients had a positive culture taken on the day of admission. None of the patients shared a clinical ward or time during hospitalization. Genotypic characterization demonstrated the existence of two clones (A and B) which were geographically associated with patients transferred from two different regions of the country.

Conclusions: During 2009, all XDR A. baumannii isolates were recovered from patients coming from other hospitals, indicative of interhospital transmission.

Key words: Acinetobacter baumannii; carbapenem resistance; inter-hospital transmission.

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#### Introduction

Acinetobacter baumannii is a major cause of nosocomial infections worldwide, especially in the intensive care unit (ICU) setting [1,2]. In Chile, carbapenem resistance in the ICU is over 70% [3], leaving few therapeutic options (polymyxins and tigecycline) for treatment of severe cases [4]. Both clonal transmission within hospitals [5] and interhospital transmission of A. baumannii have been reported [6-9]. However, the mechanism of A. baumannii transmission requires clarification to allow for better evaluation of current infection control measures with the aim of preventing hospital outbreaks and decreasing morbidity and mortality caused by clinical infections, especially in critically ill patients [10]. The aim of this study was to determine the origin of extensively drug-resistant (XDR) A. baumannii in our hospital during 2009 and to evaluate the importance of inter-hospital transmission.

### Methodology

This was a retrospective observational study performed in Hospital Clínico Universidad Católica de Chile, a tertiary care university teaching hospital. Cases were included if *A. baumannii* was isolated from any clinical sample during 2009 and met the definition of an XDR isolate. Cases were detected using records available in the clinical microbiology laboratory. Only the first isolate recovered from a patient was analyzed. Epidemiological information was collected from clinical records.

*A. baumannii* isolates were identified by conventional methods [11] and by an automatized system with GNI+ card (Vitek 2 Compact, bioMerieux, Lyon, France). Determination of antimicrobial susceptibility was carried out by agar dilution using 2009 Clinical and Laboratory Standards Institute (CLSI) criteria. [12] *A. baumanii* was classified as susceptible, multidrug resistant (MDR), and XDR using criteria published by Magiorakos *et al.* [13]. XDR isolates were genotyped by pulsed-field gel electrophoresis (PFGE) using the *Smal* restriction enzyme. Banding patterns were interpreted using the criteria specified by Tenover *et al.* [14].

The *A. baumannii* isolates were classified into three categories: a) community acquired: isolates recovered from patients within 48 hours of admission with no hospitalization or other (*e.g.*, long-term) healthcare treatment in the previous six months; b) hospital acquired: isolates recovered from patients 48 hours after admission, with epidemiological link with patients with the same strain hospitalized in the same service, at the same time; or c) inter-hospital transmission: isolates from patients transferred from or hospitalized in other centers in the previous six months, or from patients receiving healthcare (dialysis, chemotherapy, nursing home) in ambulatory centers in the previous six months and with no epidemiological link with patients hospitalized with the same strain.

## Results

During 2009, 48 isolates of A. baumannii were obtained from 34 patients. Of the 34 isolates analyzed, 18 were susceptible, 6 were MDR, and 10 were XDR. These 10 XDR A. baumannii were resistant to most classes of antibiotics, such as penicillins (including combinations with inhibitors), all classes of cephalosporins, carbapenems, fluoroquinolones, aminoglicosides, and others. Thev remained susceptible only to polimixin B (100%) and tigecycline (100%).

Of the 10 XDR isolates, 9 were recovered from patients transferred from other hospitals, and one was

from a patient with a chronic ulcer in the leg who was being treated at an outpatient clinic outside of the health network. Thus, all of these isolates met the criteria for inter-hospital transmission, as outlined above. The median time from admission to isolation was 1 day (range, 1 to 9 days), and all patients had prolonged hospitalization in their origin hospitals, with a median time of 18 days (range, 1 to 60 days). A total of 6 out of 10 patients had positive culture taken on the admission day. None of the patients shared clinical wards or time during hospitalization in the hospital (Table 1).

Genotypic characterization was possible for 8 out of 10 XDR isolates, revealing the existence of two clones. Five isolates were indistinguishable and designated Clone A. These isolates were recovered from patients transferred from four hospitals in the central area of Chile. Two additional isolates were closely related and designated Clone B. These isolates were recovered from patients coming from two hospitals in the south of Chile. The last isolate was unique and was cultured from a patient treated at an outpatient clinic (Table 1).

All 10 patients were under contact precaution during the total length of their hospitalization. Room cleaning had been implemented with no changes until now.

## Discussion

*A. baumannii* is the fifth most common cause of infections affecting patients admitted to ICUs worldwide, reaching 9% of all infections in this group. It is one of the main causes of ventilator-associated pneumonia in Latin American countries [2]. In Chile,

Table 1. Epidemiological data of 10 patients infected/colonized with extensively drug-resistant A. baumannii.

| Isolate No. | Culture sample       | Length of<br>hospitalization (α)<br>(days) | Previous<br>isolation (β) | Time from admission to culture date (days) | Clonal<br>relationship (γ) | Geographical<br>origin (µ) |
|-------------|----------------------|--|---------------------------|--|----------------------------|----------------------------|
| 3           | CSF                  | 17   | Yes                       | 1  | Clone A                    | South                      |
| 6           | ETA                  | 32   | Yes                       | 1  | Clone A                    | Center                     |
| 7           | ETA                  | 19   | No                        | 9  | Clone A                    | Center                     |
| 8           | Tissue               | 1  | No                        | 1  | Clone A                    | Center                     |
| 10          | ETA                  | 6  | No                        | 6  | Clone A                    | Center                     |
| 2           | Abdominal collection | 28   | No                        | 1  | Clone B                    | South                      |
| 9           | Blood                | 4  | No                        | 1  | Clone B                    | South                      |
| 4           | Tissue               | Outpatient                                 | No                        | 1  | Unrelated                  | Center                     |
| 1           | ETA                  | 60   | No                        | 5  | NA                         | Center                     |
| 5           | Blood                | 1  | No                        | 9  | NA                         | Center                     |

( $\alpha$ ) Length of hospitalization in previous center; ( $\beta$ ) previous isolation: patient already known to be colonized or infected at admission; ( $\gamma$ ) clonal relationship based in the pulsed-field gel electrophoresis pattern; ( $\mu$ ) geographical origin in Chile; CSF: cerebrospinal fluid; ETA: endo-traqueal aspirate.

the resistance to imipenem and meropenem in ICU patients was 76% and 80.2%, respectively, during 2012. Carbapenem resistance is mainly caused by carbapenemases belong to the OXA family [15]. Several outbreaks of clonal transmission within hospitals associated with contaminated environment and patient to patient transmission have been reported [16]. Also, inter-hospital transmission of MDR isolates has been frequently reported [6-9,17]. Epidemic MDR strains are usually introduced to the hospital by a colonized or infected patient, from whom it can spread to other patients and contaminate the environment.

In this study, we demonstrated that during 2009 in our institution, all XDR A. baumannii strains were isolated from patients coming from other hospitals or healthcare centers, indicative of inter-hospital transmission. We believe that the patients with XDR baumannii did not acquire А. the infection/colonization in our hospital because XDR A. baumannii has very low incidence in our hospital (during 2009, only 10 patients in our hospital were infected or colonized with XDR A. baumannii out of more than 20,000 discharges). A total of 9 out of 10 patients with XDR A. baumannii in our hospital were transferred from other hospitals, where A. baumannii has been endemic for many years and has a high incidence, especially in ICUs. In addition, 6 of these patients had positive culture on admission. The other 4 patients had had prolonged hospitalization in the previous hospital, had positive cultures on days 5 and 6, and 2 had positive cultures on day 9 after admission. Finally, none of the 10 patients shared a room or time with patients with XDR A. baumannii infection.

These data support our belief that implementing contact isolation precautions for transferred patients, both at admission and any time when *A. baumannii* is isolated, is essential for successfully preventing nosocomial transmission in hospitals. In addition, we also believe that environmental cleaning is important to prevent the spread of these strains. PFGE genotyping demonstrated geographic-specific XDR clones in patients (*i.e.*, related to a particular region of Chile), suggesting inter-hospital transmission within these regions.

This study has several limitations. It was a retrospective, one-year, single-hospital study. However, the data allowed us to reach important conclusions regarding appropriate infection control measures for transfer patients potentially colonized with *A. baumannii*, which is especially important given the XDR nature of these organisms in our country.

## Conclusions

During 2009 in our institution, all XDR *A. baumannii* strains were isolated from patients coming from other hospitals, indicative of inter-hospital transmission. Future studies are needed to confirm and expand these findings to better inform infection control practices aimed at limiting the transmission of *A. baumannii* in developing countries.

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#### References

- 1. Garnacho-Montero J, Amaya-Villar R (2010) Multiresistant *Acinetobacter baumannii* infections: epidemiology and management. Curr Opin Infect Dis 23: 332-339.
- Vincent JL, Rello J, Marshall J, Silva E, Anzueto A, Martin CD, Moreno R, Lipman J, Gomersall C, Sakr Y, Reinhart K; EPIC II Group of Investigators (2009) International study of the prevalence and outcomes of infection in intensive care units. JAMA 302: 2323-2329.
- Cifuentes DM, Silva F, García P, Bello H, Briceño I, Calvo AM, Labarca J; Grupo Colaborativo de Resistencia Bacteriana de Chile (2014) Antimicrobial susceptibility in Chile 2012. Rev Chil Infectol 31: 123-130.
- 4. Maragakis L, Perl T (2008) *Acinetobacter baumannii*: epidemiology, antimicrobial resistance, and treatment options. Clin Infect Dis 46: 1254-1263.
- Barbolla RE, Centrón D, Maimone S, Rospide F, Salgueira C, Altclas J, Catalano M (2008) Molecular epidemiology of *Acinetobacter baumannii* spread in an adult intensive care unit under an endemic setting. Am J Infect Control 36: 444-452.
- Mezzatesta ML, D'Andrea MM, Migliavacca R, Giani T, Gona F, Nucleo E, Fugazza G, Pagani L, Rossolini GM, Stefani S (2012) Epidemiological characterization and distribution of carbapenem resistant *Acinetobacter baumannii* clinical isolates in Italy. Clin Microbiol Infect 18: 160-166.
- Wang H, Guo P, Sun H, Wang H, Yang Q, Chen M, Xu Y, Zhu Y (2007) Molecular epidemiology of clinical isolates of carbapenem-resistant *Acinetobacter* spp. from Chinese hospitals. Antimicrob Agents Chemother 51: 4022-4028.
- van den Broek PJ, Arends J, Bernards AT, De Brauwer E, Mascini EM, van der Reijden TJ, Spanjaard L, Thewessen EA, van der Zee A, van Zeijl JH, Dijkshoorn L (2006) Epidemiology of multiple *Acinetobacter* outbreaks in The Netherlands during the period 1999-2001. Clin Microbiol Infect 12: 837-843.
- Marais E, De Jong G, Ferraz V, Maloba B, Dusé AG (2004) Interhospital transfer of pan-resistant *Acinetobacter* strains in Johannesburg, South Africa. Am J Infect Control 32: 278-281.
- Lautenbach E, Synnestvedt M, Weiner MG, Bilker WB, Vo L, Schein J, Kim M (2009) Epidemiology and impact of imipenem resistance in *Acinetobacter baumannii*. Infect Control Hosp Epidemiol 30: 1186-1192.
- Schreckenberger PC, Daneshvar MI, Weyant RS, Hollis DG (2005) Acinetobacter, Achromobacter, Chryseobacterium, Moraxella and other nonfermentative gram-negative rods. In Murray PR, Baron EJ, Jorgensen JH, editors. Manual of

Clinical Microbiology, 8th edition. Washington: American Society for Microbiology. 749-779.

- Clinical and Laboratory Standards Institute (2009) Performance Standards for Antimicrobial Susceptibility Testing: Nineteenth Informational Supplement M100-S19. Wayne, USA: CLSI.
- Magiorakos AP, Srinivasan A, Carey RB, Carmeli Y, Falagas ME, Giske CG, Harbarth S, Hindler JF, Kahlmeter G, Olsson-Liljequist B, Paterson DL, Rice LB, Stelling J, Struelens MJ, Vatopoulos A, Weber JT, Monnet DL (2012) Multidrugresistant, extensively drug-resistant and pandrug-resistant bacteria: an international expert proposal for interim standard definitions for acquired resistance. Clin Microbiol Infect 18: 268-281.
- Tenover FC, Arbeit RD, Goering RV, Mickelsen PA, Murray BE, Persing DH, Swaminathan B (1995) Interpreting chromosomal DNA restriction patterns produced by pulsed field gel electrophoresis: criteria for bacterial strain typing. J Clin Microbiol 33: 2233-2239.
- 15. Labarca JA, Salles MJ, Seas C, Guzmán-Blanco M (2014) Carbapenem resistance in Pseudomonas aeruginosa and

Acinetobacter baumannii in the nosocomial setting in Latin America. Crit Rev Microbiol 27: 1-17.

- van den Broek PJ, van der Reijden YKJ, van Strijen E, Helmig-Schurter AV, Bernards AT, Dijkshoorn L (2009) Endemic and epidemic *Acinetobacter* species in a university hospital: an 8-year survey. J Clin Microbiol 47: 3593-3599.
- 17. Sader HS, Mendes CF, Pignatari AC, Pfaller MA (1996) Use of macrorestriction analysis to demonstrate interhospital spread of multiresistant *Acinetobacter baumannii* in São Paulo, Brazil. Clin Infect Dis 23: 631-634.

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