

## ABSTRACT

**Objective:** To evaluate the high prevalence of *Acinetobacter* spp. (ASP) displaying elevated tigecycline MICs (>2 mg/L) in Latin American (LATAM) hospitals surveyed by the SENTRY Program. We recently noted a significant difference in the percentage of ASP with tigecycline MICs >1 mg/L in LATAM compared to other geographic regions of the world (12.5% vs. 3.9-6.6%).

**Methods:** 1,950 ASP were received from LATAM during 2005-2011-period (9 hospitals from 2005 to 2010 and 20 in 2011). Isolates were susceptibility tested according to CLSI guidelines. *A. baumannii* (ACB) isolates from 2011 displaying tigecycline MIC values >2 mg/L were molecularly typed by PFGE. Expression of *adeA* and *adeF* encoding the efflux pumps AdeABC and AdeFGH was determined for 18 unique isolates from 2011 using high quality DNA-free RNA preparations and measured by quantitative RT-PCR, normalized using *rpoB* and compared to ACB ATCC 19606.

**Results:** ASP displaying tigecycline MIC values >1 mg/L varied from 6.9 to 32.2% in the study period and showed an increase from 14.6% in 2010 to 32.2% in 2011 ( $p < 0.0001$ ; OR=0.16[0.09-0.28]). Isolates with confirmed tigecycline MIC values >2 mg/L were 49 *A. baumannii* and 1 *A. pittii* (formerly genomic species 3; by MALDI-TOF). Isolates were mainly from Sao Paulo, Brazil (SP; 29 isolates) and Guadalajara, Mexico (14), but also from Durango, Mexico (3), Florianopolis, Brazil (1), Panama City, Panama and Santiago, Chile (1). PFGE showed that 15/29 isolates from SP belonged to a single clone. Three other clusters were noted in the same hospital (5, 3 and 2 isolates). Ten strains from Guadalajara belonged to a major clone and the remaining 4 strains belonged to two other PFGE types. All three strains from Durango were genetically related. Expression results of AdeABC and AdeFGH tested for 18 strains with MIC values ranging to 4 to 8 mg/L showed that only two isolates had significantly greater expression of AdeFGH (>10-fold difference from the control ATCC strain) both from clonal groups from SP and displaying tigecycline MIC values of 4 mg/L. All strains had AdeABC expression similar to the control strain.

**Conclusions:** We documented the recent increase of ASP displaying elevated tigecycline resistance in LATAM hospitals, dominantly due to the clonal expansion of isolates in Brazil and Mexico. Control of tigecycline usage in those countries and more strict infection control practices in the involved centres will be needed to contain these ACB outbreaks.

## INTRODUCTION

*Acinetobacter baumannii* is a nosocomial pathogen that can cause various types of opportunistic infections in patients suffering of underlying conditions that have been hospitalized for extended periods. These organisms that are commonly isolated worldwide have the ability to acquire resistance to several antimicrobial agents and *A. baumannii* populations display high resistance rates to virtually all antimicrobial agents that are clinically available. Mechanisms that lead to multidrug resistance are particularly important for this species; and it may occur due to horizontal acquisition of genetic elements carrying several resistance genes or overexpression of chromosomally encoded efflux systems that can lead to the extrusion of compounds from several antimicrobial classes.

Tigecycline resistance in *A. baumannii* has been related to the overexpression of two resistance-nodulation-cell division (RND) multidrug transporters, AdeABC and AdeFGH. AdeABC overexpression contributes to resistance to various antimicrobial classes, including  $\beta$ -lactams, aminoglycosides, quinolones and tigecycline. Additionally, this efflux system is regulated by *adeRS* operon that is a two-component regulation system and mutations on the components *adeS* and *adeR* have also been associated with tigecycline resistance in single-step mutants. AdeFGH overexpression confers high-level resistance to quinolones, chloramphenicol, trimethoprim and clindamycin as well as decreased susceptibility to tetracycline, tigecycline and sulfamethoxazole without affecting  $\beta$ -lactams and aminoglycosides.

In this study, we evaluated a total of 1,950 *A. baumannii* isolates collected in Latin American hospitals as part of the SENTRY Antimicrobial Surveillance Program to document an increase in tigecycline resistance in the most recent years. Isolates from 2011 were evaluated for clonality and 18 unique strains were submitted to further assays to determine the expression of AdeABC and AdeFGH.

## MATERIALS AND METHODS

**Bacterial isolates.** A total of 1,950 *Acinetobacter* spp. isolates collected in nine (2005-2010) to twenty (2011) Latin American hospitals were analysed. Only one isolate per patient from documented bloodstream infections were included in the study. Species identification was confirmed for all isolates displaying tigecycline MIC values >2 mg/L from 2011 by Matrix-Associated Laser Desorption Ionization-Time Of Flight mass spectrometry (MALDI-TOF MS) using the Bruker Daltonik MALDI Biotyper (Billerica, Massachusetts, USA) by following the instructions of the manufacturer.

**Antimicrobial susceptibility testing.** All isolates were susceptibility tested using the broth microdilution method as described by the Clinical and Laboratory Standards Institute (CLSI, M07-A9) with freshly made Mueller-Hinton broth. Quality control (QC) was performed using *Escherichia coli* ATCC 25922 and *Pseudomonas aeruginosa* ATCC 27853. All QC results were within specified ranges as published in CLSI documents (M100-S23).

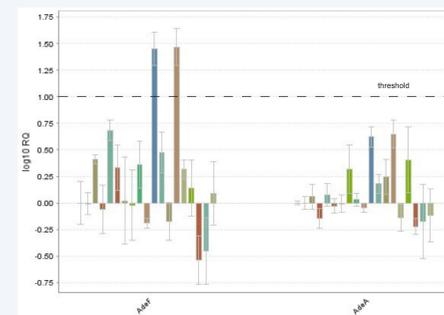
**Molecular typing.** *A. baumannii* isolates showing tigecycline MIC values at >2 mg/L, collected in 2011 (n=49) were epidemiologically typed by pulsed-field gel electrophoresis (PFGE) using previously described procedures. Genomic DNA was prepared in agarose blocks and digested with SmaI (New England, Beverly, Massachusetts, USA) and resolved in the CHEF-DR III (BioRad, Richmond, California, USA) using running conditions described elsewhere. Results were analyzed by GelCompar II software (Applied Math, Kortrijk, Belgium). Percent similarities were identified on a dendrogram derived from the unweighted pair group method using arithmetic averages and based on Dice coefficients. Band position tolerance and optimization were set at 1.2% and 0.5%, respectively.

**Expression of RND systems.** The expression of *adeA* and *adeF* was determined by quantitative real-time PCR (qRT-PCR) using DNA-free RNA preparations for 18 tigecycline-non-susceptible *A. baumannii* unique isolates from 2011. Total RNA was extracted from mid-log-phase bacterial cultures (cell density at OD<sub>600</sub> of 0.3-0.5) using RNA Protect Reagent and RNeasy Mini Kit (Qiagen, Hilden, Germany) in the Qiacube workstation (Qiagen) and residual DNA was eliminated with RNase-free DNase (Promega, Madison, Wisconsin, USA). Quantification of mRNA and sample quality was assessed using the RNA 6000 Pico kit on the Agilent 2100 Bioanalyzer (Agilent Technologies, Santa Clara, California, USA) according to manufacturer instructions. Only preparations with RNA integrity number (RIN) >6.5 that showed no visual degradation were used for experiments. Relative quantification of target genes was performed in triplicate by normalization to an endogenous reference gene (*rpoB*) on the StepOne Plus instrument (Life Technologies, Carlsbad, California, USA) using Power SYBR® Green RNA-to-CT™ (Life Technologies) and custom designed primers showing efficiency >98.0%. Transcription levels were considered significantly different if at least a 5-fold difference was noted compared with *A. baumannii* ATCC 19606.

## RESULTS

- A total of 244 (12.5%) of the *Acinetobacter* spp. isolates from Latin America hospitals displayed tigecycline MIC values of >1 mg/L. Tigecycline non-wildtype (WT) population MIC values (>1 mg/L) varied from 6.9 to 32.2% with an increasing trend in more recent years: 32.2% in 2011 and 14.6% in 2010 ( $p < 0.0001$ ; OR=0.16[0.09-0.28]); see EUCAST ECOFF tables <<http://mic.eucast.org/Eucast2/SearchController/search.jsp?action=performSearch&BeginIndex=0&Micdif=mic&NumberIndex=50&Antib=345&Specium=-1>>.

**Figure 1.** Relative quantification of transcriptional levels for *adeA* and *adeF* genes compared to those obtained from *A. baumannii* ATCC 19606.



**Table 1.** Susceptibility testing results, PFGE patterns and efflux pump expression results for 18 *A. baumannii* isolates displaying elevated tigecycline MIC values ( $\geq 2$  mg/L) selected for follow up studies.

Country	City	MIC in mg/L:							Relative Expression of efflux pumps <sup>a</sup>			
		Tigecycline	Doxycycline	Tetracycline	Minocycline	Ceftazidime	Imipenem	Levofloxacin	Colistin	PFGE	AdeABC	AdeFGH
Brazil	Florianopolis	8	>8	>8	2	>32	>8	>4	1	046B	0.600	0.290
		4	2	>8	2	>32	>8	>4	1	048A1	0.986	1.049
	Sao Paulo	8	4	>8	2	>32	>8	>4	1	048B	0.996	0.982
		8	4	>8	4	>32	>8	>4	>4	048B	1.084	2.293
		4	2	>8	2	32	>8	>4	1	048C	0.675	0.352
		4	4	>8	2	32	>8	>4	1	048D	0.935	2.136
		4	1	>8	1	>32	>8	>4	2	048E1	4.238	<b>28.237<sup>b</sup></b>
		4	1	>8	0.5	>32	>8	>4	0.5	048F	4.510	<b>29.464<sup>b</sup></b>
Chile	Santiago	4	1	>8	1	>32	>8	>4	1	048G	1.527	2.988
		8	2	>8	1	>32	>8	>4	2	048I	2.621	1.373
Mexico	Durango	4	4	>8	2	>32	>8	>4	>4	043A	1.760	0.673
		4	>8	>8	4	32	2	>4	0.5	126A1	0.770	1.228
	Guadalajara	4	>8	>8	>8	>32	0.5	>4	2	126A2	0.735	0.865
		4	4	>8	2	>32	>8	>4	1	115A1	1.233	4.819
Panama	Panama City	4	2	>8	2	>32	>8	>4	0.5	115B1	0.729	2.065
		4	>8	>8	>8	>32	>8	>4	0.5	115B1	0.940	0.647
Panama	Panama City	4	1	>8	1	>32	>8	>4	0.5	115C	1.148	2.576
		4	>8	>8	8	>32	>8	>4	2	346A	2.127	0.951

a. Relative expression level that corresponds to the critical threshold ( $C_t$ ) numbers determined by the detection system software as the amount of target was given as  $2^{-\Delta\Delta C_t}$ , where  $\Delta\Delta C_t$  is the difference between the target (*adeA* and *adeF*) and reference gene (*rpoB*)  $C_t$  values and was determined in comparison to the *A. baumannii* ATCC 19606 (relative expression = 1.000).

b. Values in bold were considered significant upregulation of the gene expression (>10-fold difference from the control ATCC strain).

## CONCLUSIONS

- Increased tigecycline resistance rates were secondary to the dissemination of clonal strains in two hospitals that participate in the surveillance program over all years sampled. Two isolates were recovered in hospitals only surveyed in 2011.
- We did not detect strains with elevated expression of AdeABC and only two sampled organisms had upregulation of AdeFGH. Additional studies are being carried out in an attempt to further explain the elevated tigecycline MIC values resulting from these Latin American isolates.
- More stringent infection control and antimicrobial stewardship practices appear needed to control increasing tigecycline resistance rates in these Latin American hospitals, where tigecycline and colistin might be prescribed for multidrug-resistant *Acinetobacter* infections.

## SELECTED REFERENCES

- Bratu S, Landman D, Martin DA, Georgescu C, Quale J (2008). Correlation of antimicrobial resistance with beta-lactamases, the OmpA-like porin, and efflux pumps in clinical isolates of *Acinetobacter baumannii* endemic to New York City. *Antimicrob Agents Chemother* 52: 2999-3005.
- Chu YW, Chau SL, Houang ET (2006). Presence of active efflux systems AdeABC, AdeDE and AdeXYZ in different *Acinetobacter baumannii* genomic DNA groups. *J Med Microbiol* 55: 477-478.
- Clinical and Laboratory Standards Institute (2012). M07-A9. *Methods for dilution antimicrobial susceptibility tests for bacteria that grow aerobically; approved standard: ninth edition*. Wayne, PA: CLSI.
- Clinical and Laboratory Standards Institute (2013). M100-S23. *Performance standards for antimicrobial susceptibility testing: 23rd informational supplement*. Wayne, PA: CLSI.
- Coyne S, Courvalin P, Perichon B (2011). Efflux-mediated antibiotic resistance in *Acinetobacter* spp. *Antimicrob Agents Chemother* 55: 947-953.
- EUCAST (2013). Breakpoint tables for interpretation of MICs and zone diameters. Version 3.0, January 2013. Available at: [http://www.eucast.org/clinical\\_breakpoints/](http://www.eucast.org/clinical_breakpoints/). Accessed January 2, 2013.
- Ruzin A, Immermann FW, Bradford PA (2010). RT-PCR and statistical analyses of *adeABC* expression in clinical isolates of *Acinetobacter calcoaceticus*-*Acinetobacter baumannii* complex. *Microb Drug Resist* 16: 87-89.