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The long-term survival of *Acinetobacter baumannii* ATCC 19606^T under nutrient-deprived conditions does not require the entry into the viable but non-culturable state

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Abstract Acinetobacter baumannii possesses a tremendous potential to thrive under hostile conditions. To learn more about its survival strategy and capacity to persist in the environment, we studied the effect of temperature, nutrient deprivation and dryness on the long-term survival of two A. baumannii strains (ATCC 19606^T and a clinical isolate). Our results revealed that both strains show a great persistence under stress that appears to involve a bust-andboom strategy. Bacterial survival was differentially affected by temperature and physical environment: Desiccation favored cell resistance to stress at 20 and 37 °C, while survival in aqueous environments was temperature dependent and led to changes in several cellular characteristics. In addition, we tested the ability of the A. baumannii ATCC 19606^T strain to form biofilms by monitoring the expression of adhesion-/biofilm-related genes (ompA, bfmR and csuAB). The observed downregulation of these genes suggests that the potential difficulties to adhere to solid surfaces and form biofilms likely limit the capacity of starved cells to spread and colonize abiotic surfaces.

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³ IKERBASQUE, Basque Foundation for Science, Bilbao, Spain **Keywords** Acinetobacter baumannii · Persistence · Survival strategy · Temperature · Dryness

Introduction

Microorganisms in the environment are often exposed to adverse physicochemical conditions. To increase their capacity of persistence and survival in hostile environments, bacteria can trigger a number of adaptation mechanisms. Rittershaus et al. (2013) highlighted three major strategies enabling to cope with the growth-limiting stress: bust-and-boom, cellular quiescence and true dormancy (sporulation). This last strategy can be neglected for nonspore-forming bacteria.

In the bust-and-boom model, most cells die upon starvation, and the few survivors subsist at the expense of the dead cells. These remaining cells have been defined as persisters (Zhang 2014). In the second model, cellular quiescence/dormancy or acquisition of viable but non-culturable (VBNC) state represents a common strategy enabling many non-differentiating bacteria to survive under various adverse conditions (Roszak and Colwell 1987; Barcina and Arana 2009; Oliver 2010). The entry into the VBNC state suggests that cells are unable to grow on the media that are normally used to culture them, although they retain metabolic activity and can possibly act as pathogens (Karunasagar and Karunasagar 2005), thus representing a hidden risk to public health. Moreover, since the VBNC response can occur simultaneously with the bust-and-boom strategy, Arana et al. (2007) suggested that VBNC cells could serve as a source of nutrients to sustain the survival of the persisting culturable cells until environmental conditions improve.

Over the last decade, Acinetobacter baumannii has emerged as an important nosocomial pathogen that has

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become endemic in some hospitals, causing serious opportunistic infections (Towner 2009; Lambiase et al. 2012; Roca et al. 2012). Hospital outbreaks of *A. baumannii* have been frequently reported worldwide from many medical, surgical and neonatal intensive care and burn units (Diancourt et al. 2010; Roca et al. 2012). Moreover, the emergence and rapid spread of multidrug-resistant isolates becomes a great concern because very few therapeutic options remain effective against them (Kempf et al. 2012; Eveillard et al. 2013).

Due to its clinical significance, during the last years several studies have contributed to understanding the pathogenesis of this organism. Although it is evident that this opportunistic pathogen has developed several mechanisms that control its persistence and spread, very little is known about the survival strategies that could explain the high persistence of this pathogen in adverse environments. Gayoso et al. (2014) have suggested that the bust-and-boom model could explain the recurrent outbreaks of *A. baumannii* found in intensive care units; however, the viability of nonculturable cells was not examined to assess the possible involvement of the VBNC state.

On the other hand, the environmental survival of A. baumannii has been suggested to involve its ability to colonize biotic and abiotic surfaces (Espinal et al. 2012). Among the biofilm-related virulence determinants, the pilus usherchaperone assembly system (encoded by the csuAB-A-B-C-D-E gene cluster) and the outer membrane protein OmpA appear to play the major role. The expression of the csuAB-A-B-C-D-E gene cluster is controlled by a two-component regulatory system comprising the sensor kinase BfmS and the response regulator BfmR encoded by the bfmRS operon (Tomaras et al. 2008; Gaddy and Actis 2009). Inactivation of *bfmR* results in a loss of expression of the *csuAB* operon and the subsequent inhibition of both pili production and biofilm formation on abiotic surfaces (Tomaras et al. 2008). OmpA is essential for attachment to biotic and abiotic surfaces and biofilm formation (Choi et al. 2008; Lee et al. 2008).

Study and characterization of the changes occurring during the long-term starvation process can contribute to a better understanding of *A. baumannii* survival in the environment and its persistence strategies, which could provide key information for its control. The aim of this study was to examine the effect of temperature and nutrient-deprived environments on *A. baumannii* ATCC 19606^T (model strain) and on a clinical isolate to assess how environmental conditions affect *A. baumannii* survival and persistence. Moreover, we also analyzed how these stressful conditions affect the expression of genes (*ompA*, *bfmR* and *csuAB*) implicated in adhesion and biofilm formation and known to play essential roles in colonization of abiotic environments.

Materials and methods

Acinetobacter baumannii strain and inocula preparation

Two Acinetobacter baumannii strains were used in this study: American Type Culture Collection (ATCC) strain 19606^T and a clinical isolate of *A. baumannii*, strain 06-2790, obtained from a skin ulcer of a patient at the Hospital Universitario Marqués de Valdecilla (Prof. J Ramos-Vivas, Santander, Spain). Both strains were separately grown overnight at 37 °C in Mueller–Hinton (MH) broth. Cells were transferred to fresh MH broth and grown at 37 °C with shaking (120 rpm) to stationary phase. Cells were collected by centrifugation (4500 g for 15 min) and washed three times with sterile saline solution (0.9 % w/v NaCl). Finally, the pellets were suspended in sterile saline solution.

Survival assays

Acinetobacter baumannii cells from stationary phase were incubated under nutrient deprivation at 20 or 37 °C in liquid or solid environments. Starvation was implemented by incubating cells in sterile saline solution or on cellulose acetate filters (Whatman, GE Healthcare Life Sciences) sterilized by a 20-min exposure to UV-C (approximately 253.7 nm; 70 mW/cm²).

For survival assays in the aqueous environment, experiments were carried out in Erlenmeyer flasks containing 300 ml of sterile saline solution inoculated to reach a density of 10^8 cells/ml. To avoid organic residues, the glass flasks were first cleaned with acid, rinsed with deionized water and kept at 250 °C for 24 h.

For survival assays on solid surfaces, cellulose acetate filters were inoculated with *A. baumannii* by filtering cellular suspensions (described above) to a density of approximately 10^8 cells/cm². The filters were incubated in sterile Petri dishes at 20 or 37 °C. Ambient humidity inside of Petri dishes was measured with a Fisher ScientificTM TraceableTM Digital Hygrometer/Thermometer (Fisher Scientific) and maintained at a relatively low level (21–27 %) at 20 and 37 °C. For each strain, the samples for subsequent analyses consisted of three randomly chosen filters that were individually placed in 10 ml of sterile saline solution and vigorously shaken for 2 min.

For the survival assays from both liquid and solid environments, subsamples were periodically collected to enumerate total, viable and culturable bacteria and analyzed directly their ability to adhere. Moreover, for *A. baumannii* ATCC 19606^T, the expression of *ompA*, *bfmR* and *csuAB* genes by reverse transcription quantitative PCR (RT-qPCR) was analyzed.

Bacterial counts

The total bacterial count (TBC) was determined by means of the standard acridine orange staining as described by (Hobbie et al. 1977). Viable bacteria were estimated as bacteria with intact cytoplasmic membranes (MEMB+). These MEMB+ bacteria were counted with the Live/Dead BacLightTM kit (Invitrogen Life Technologies) (Joux et al. 1997). Culturability, expressed as colony-forming units (CFUs), was determined by spreading aliquots of *A. baumannii* cells on Mueller–Hinton agar followed by incubation at 37 °C for 24 h.

Scanning electron microscopy (SEM)

A. baumannii ATCC 19606^T cells, periodically collected from survival assays in aqueous environments, were fixed with 2 % glutaraldehyde in 0.1 M sodium phosphate buffer (pH 7.4). The suspensions of the fixed cells were filtered through 0.22- μ m-pore-size membrane filters (GTTP02500 filters, Merck Millipore). The filters and the attached *A.* baumannii cells were further dehydrated by applying a series of increasing ethanol concentrations (30, 50, 70, 90 and 100 %). The filters were then overlaid with 1 ml of hexamethyldisilazane, incubated for 5 min and air-dried. Finally, the samples were coated with gold, and imaging was carried out by analyzing samples in a Hitachi S4800 scanning electron microscope.

Quantification of cell adherence to solid surfaces

To assess cell adherence, we employed the microtiter dish assay (O'Toole and Kolter 1998) with some modifications. Aliquots of cell suspensions (1 ml each), periodically collected from survival experiments, were added into the wells of sterile 24-well polystyrene plates (Corning Inc.) and incubated for 48 h at 37 °C without shaking. Sterile saline solution was used as a negative control. After incubation, plates were washed three times with sterile water and stained with 200 µl of 0.5 % (w/v) crystal violet solution. Then, plates were newly washed with sterile water (three times), and the dye bound to the attached cells was solubilized in 0.5 ml of 95 % ethanol. The optical density (OD) of each well was measured at 595 nm using an automated SynergyTM HT Multi-Detection Microplate Reader (BioTek Instruments, Inc., Vermont). During the survival experiments, cells were classified according to Stepanović et al. (2000). The cutoff OD (ODc) was defined as a value equal to three standard deviations above the mean OD of the negative control, and the following categories were established: $OD \leq OD_{C}$, non-adherent; $ODc < OD \leq (2 \times ODc)$, weakly adherent; $(2 \times ODc) < OD \leq (4 \times ODc)$,

moderately adherent; and $(4 \times ODc) < OD$, strongly adherent cells.

Along the survival experiments, the variation in the ability to adhere was expressed with an arbitrary value, OD decline values which were calculated by subtracting the respective initial OD from those obtained in each survival period.

Statistical data analysis

Statistical tests were carried out with the Stat View program (Abacus Concept, Inc.). All of the results presented are means of at least three experiments, and the coefficients of variation between replicate experiments were less than 12 %. The differences between the means were detected by a one-way analysis of variances. Probabilities less than or equal to 0.05 were considered significant. Logarithmic transformation for bacterial counts was used.

RNA preparation and reverse transcription quantitative PCR (RT-qPCR) analysis

Samples of cells collected from the survival experiments of A. baumannii ATCC 19606^T (aqueous and dry conditions) were mixed with stop solution (5 % phenol in ethanol) at a ratio of 8:1 and incubated on ice for 15-20 min, and the cells were collected by centrifugation (15 min, 4 °C, 4400 g). The pelleted cells were used to isolate total RNA using the Trizol® MaxTM Bacterial RNA Isolation Kit (Ambion). Each RNA sample was further purified using the PureLink® RNA Mini Kit (Ambion) and treated with DNase I (Invitrogen). RNA samples were subjected to RTqPCR at the General Genomic Service (SGIker) of the University of Basque Country (Spain). Namely, after verifying RNA quality and integrity by Lab-chip technology on an Agilent 2100 Bioanalyzer with Agilent RNA 6000 Nano Chips, RNA was used for cDNA synthesis using Affinity-Script Multiple Temperature cDNA Synthesis Kit (Agilent Technologies).

The expression of three different genes related to adhesion or biofilm formation (*ompA*, *bfmR* and *csuAB*) was analyzed by RT-qPCR. Specific primer pairs were designed using PrimerQuest tool available at IDT Integrated DNA Technologies, using universal primer design parameters for qPCR. Primer secondary structure analysis (hairpin and primer dimer formation analysis) was verified with Oligo-Analyzer tool, from IDT Integrated DNA Technologies, and primer specificity was checked using BLAST (NCBI). Specific primer pairs were designed for four candidate reference genes potentially suitable for normalization purposes: *gltA*, *gyrB*, *cpn60* and *gdhB* (Bartual et al. 2005; Hamouda et al. 2010). The target and candidate reference

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Gene Forward primer $(5'-3')$		Reverse primer $(5'-3')$	
ompA	GCTTCTCGTTTGTCTACTCAAGG	GCGAATACACGACGGTTCATAG	
bfmR	GTTCGCCCACACTATCATCAA	TAATCGTCTGCACCCATTTCC	
csuA/B	ACTGCTTCTGCTGATGTAGTTG	GTAGTGAACTGTTGTGGTTGGT	
gltA	GCGAGTTACCAACTGCTGAA	ACGACGGAAACCATTGAAGAA	
gyrB	GCCGTTTACGTGAGCTTTCT	GCCGCCTTCATAGTCATACAC	
cpn60	GGTAAAGAAGGCGTAATCACTGTA	GGTAAAGAAGGCGTAATCACTGTA	
gdhB	CCGACATATAGCACGACTTTGG	TACCTTCTGGACTAGCGATGAC	

gene specific primers used for RT-qPCR are listed in Table 1.

The RT-qPCR was carried out using Brilliant III Ultra-Fast SYBR Green QPCR Master Mix (Agilent Technologies), and 10 ng of cDNA and 500 nM of primers were added per reaction. RT-qPCR experiments were performed in 384-well plates in a 7900HT Fast Real-Time PCR System (Applied Biosystems) to determine the relative changes in the level of three transcripts. The PCR amplification conditions were 3 min at 95 °C followed by 40 cycles of 95 °C for 5 s and 60 °C for 20 s. The specificity of the primer pairs was verified by melting curve analysis following the last amplification cycle. No-template controls and minus reverse transcriptase controls (samples in which no reverse transcriptase was added) were also included. Having found no reference candidate genes with low variability, the amount of cDNA, quantified fluorometrically by Qubit 2.0 (Invitrogen), was used for data normalization. All reactions were carried out in triplicate for three biological replicates.

RT-qPCR data analysis

The analysis of RT-qPCR results was carried out using the SDS 2.4 software (Applied Biosystems). The qPCR efficiency calculation and correction, and data normalization with respect to the amount of cDNA were done using GenEx version 5.4 software (MultiD). Then, the relative changes in mRNA expression levels were determined with respect to the initial amounts of cDNA obtained by RT, and the significance of differences was assessed using Student's *t* test (p < 0.05).

Results

The effects of temperature (20 or 37 °C) and desiccation on culturability, viability and integrity of *A. baumannii* under nutrient deprivation are summarized in Fig. 1. For both strains, the total bacterial count (TBC) did not change during permanence in aqueous environments (Fig. 1a, b, e, f) or on solid surfaces (Fig. 1c, d, g, h), suggesting that *A. baumannii* cells preserved their integrity throughout the course of the experiment (at least 30 days). Moreover, culturability and viability also remained unchanged except for the populations incubated at 37 °C in saline solution (Fig. 1b, f). In this last case, the numbers of viable and culturable *A. baumannii* ATCC 19606^T cells decreased by 100-fold between 5 and 12 days of incubation (Fig. 1b) and *A. baumannii* clinical isolate 06-2790 by 15-fold during the same period (Fig. 1f). The concurrent loss of viability and culturability indicates that a fraction of population presented cellular membrane injured and the VBNC state was not induced under these conditions. For both strains, at the end of the experimental period, there were approximately 10^5 viable and culturable cells/ml.

Scanning electron microscopy (SEM) analysis of *A. baumannii* ATCC 19606^T populations during survival experiments (Fig. 2) revealed some cells with altered cell morphology (distorted cells). These cells were only observed in populations maintained in saline solution at 37 °C for ten or more days (Fig. 2c).

To learn more about the putative survival strategies employed by A. baumannii during its persistence in saline solutions, we also estimated the ability of populations maintained in aqueous and dry environments to adhere to solid surfaces (Fig. 3a). We found that the A. baumannii ATCC 19606^T strain behaves as a moderately adherent strain $(OD = 0.224 \pm 0.048)$ and its ability to adhere to solid surfaces progressively decreased along the survival period in both environments, being also temperature dependent (Fig. 3a). In the experiments in saline solution, this ability diminished quickly resulting in the complete loss of adhesiveness after 5 days (data not shown). Under dry conditions, the adhesiveness was also reduced but the populations retained a weak capacity to adhere at least for 15 days (data not shown). A. baumannii clinical isolate 06-2790, a strongly adherent strain $(OD = 0.529 \pm 0.065)$, showed a similar behavior (Fig. 3b).

During the survival experiments, concentration of total RNA per cell varied with conditions (Table 2). In general, despite slight fluctuations, the amount of total RNA recovered per cell in *A. baumannii* ATCC 19606^T populations maintained on the dry surfaces was constant over time. In contrast, variations in the amount of RNA were more evident in planktonic cells, especially at 37 °C, where the



Fig. 1 Evolution of *Acinetobacter baumannii* ATCC 19606^T (**a**–**d**) and clinical isolate 06-2790 (**e**–**h**) populations maintained in aqueous environment at 20 °C (**a**, **e**) and 37 °C (**b**, **f**) and on dry surfaces at

20 °C (**c**, **g**) and 37 °C (**d**, **h**). Total (*filled circle*), viable (*gray square*) and culturable (*inverted triangle*) cells

quantity of RNA recovered at the end of the experiment (after 8 days) was only 6 % when compared to the amount isolated from the cells present at the beginning of the experiments (Table 2).

The effects of temperature and desiccation under nutrient deprivation on the expression of the *ompA*, *bfmR* and *csuAB* genes are shown in Fig. 4. Regardless of the experimental conditions, for *A. baumannii* model strain, the expression of *ompA* and *bfmR* genes decreased progressively ($p \le 0.05$) from the beginning of the experiments. After 8 days of experimentation, lower expression was detected in cells exposed to dry environments at 37 °C. Both *ompA* and *bfmR* genes were downregulated by over threefold with respect to the initial values. *csuAB* gene showed greater variability in expression depending on environment. In contrast to slight variations detected for



Fig. 2 Scanning electron microscopy (SEM) analysis of the morphology and integrity of *Acinetobacter baumannii* ATCC 19606^{T} cells maintained in saline solution (aqueous environment). Panel **a** shows cells sampled at the beginning of the experiments and panels **b** and **c** after more than 10 days of permanence at 20 °C and 37 °C, respectively

populations maintained in saline solution at 20 and 37 °C, the *csuAB* gene was readily downregulated when the populations were maintained on dry surfaces.

Discussion

exposed to adverse conditions (nutrient deprivation, desiccation or non-optimal temperatures). To better characterize its ability to persist under stress (potentially facilitating infection outbreaks), we studied *A. baumannii* adaptation to starvation.

We found that its survival in liquid environments was clearly temperature dependent. Populations of A. baumannii ATCC 19606^T and clinical isolate 06-2790 both incubated at 37 °C displayed a loss of viability and culturability (Fig. 1b, f), resulting in a large fraction of non-culturable non-viable cells [i.e., cells showing altered cytoplasmic membrane permeability (Joux et al. 1997)] and the appearance of distorted cells (Fig. 2c). The loss of viability and physiological functions at 37 °C seems to correlate with a profound decrease in the amount of total RNA in the stressed cells (Table 2), thus indicating that this temperature can be detrimental under certain starvation conditions. As a result, only a small fraction (ca. 0.2 %) of the whole population remained culturable after 30 days. In contrast, no loss of culturability was observed when A. baumannii ATCC 19606^T was exposed to lower temperature (20 °C) at least for 30 days. The moderate decrease in the RNA content observed after 2 and 8 days could be associated with the overall decrease in the metabolism, thereby helping to save energy in favor of cell survival (Lever et al. 2015). Although there is no published work addressing the survival of A. baumannii in aqueous environments, several studies that were previously carried out on other mesophilic bacteria, including Escherichia coli and Pseudomonas fluorescens (Arana et al. 2010), support our observation that lower temperatures could be beneficial for prolong persistence of mesophilic bacteria under starvation.

Unlike survival in saline solutions, experiments with A. baumannii cells retained on dry surfaces revealed that the resistance to stress and survival under such conditions was even better. In other words, the viability and culturability of A. baumannii population did not experience significant changes for at least one month, and total RNA content remained constant for 8 days regardless of the temperature. Although several works (Wendt et al. 1997; Jawad et al. 1998; Espinal et al. 2012; Gayoso et al. 2014) have highlighted that A. baumannii survives long time on dry surfaces, these studies generally associate the resistance to desiccation to the longtime presence of only some culturable surviving bacteria. In this sense, our results differ from those obtained by Gayoso et al. (2014) and Wendt et al. (1997), who described the negative effect of dry conditions. With reference to these reports, it is even more remarkable that, in this study, the bacteria on dry surfaces at 37 °C could survive longer (and with higher percentage of survivors) than their planktonic counterparts. Webster et al. (2000) observed that a local strain of Acinetobacter species could persist on common clinical surfaces for relatively

Fig. 3 Time-dependent changes in the ability of *Acinetobacter baumannii* ATCC 19606^T (**a**) and clinical isolate 06-2790 (**b**) populations maintained in aqueous environment (*filled rectangle*) and under dry conditions (*open rectangle*) to adhere to solid surfaces at 20 °C and 37 °C. Relative values were calculated with respect to the initial time (0 days)



Table 2Total RNA (\pm SD)
content expressed as fg/cell and
determined during starvation
under aqueous and dry
conditions at 20 and 37 °C

Starvation conditions	Time (days)				
	0	2	4	8	
Aqueous conditions					
20 °C	42.01 ± 2.30	$74.56 \pm 6,.09$	23.93 ± 1.35	35.05 ± 5.18	
37 °C	42.01 ± 2.30	7.10 ± 0.58	6.20 ± 0.14	2.52 ± 0.56	
Dry conditions					
20 °C	50.12 ± 5.39	69.20 ± 3.75	32.66 ± 13.34	57.89 ± 4.37	
37 °C	50.12 ± 5.39	41.09 ± 14.60	54.14 ± 4.71	55.34 ± 11.68	

long periods, and this could partly explain that during the outbreaks the same *A baumannii* strain(s) continues to be recovered from both patients and environment. This result is consistent with our data and those obtained in other studies like the one conducted by Houang et al. (1998). The latter described the persistence (without loss of culturability) of a clinical isolate of *A. baumannii* maintained over 30 days at 22-24 °C under desiccation.

Different strategies have been envisaged to explain the mechanisms that allow bacteria to face stress conditions. The acquisition of the VBNC phenotype by Acinetobacter has been demonstrated only for A. calcoaceticus in aquatic systems (Lemke and Leff 2006). Gayoso et al. (2014) indicated that A. baumannii strain AbH12O-A2 maintained under prolonged desiccation conditions becomes dormant, a definition that corresponds to the VBNC state (Rittershaus et al. 2013), and that a few surviving cells embedded in biofilm followed a bust-and-boom strategy. In our work, populations of A. baumannii starved at 37 °C in a liquid environment (the only condition we found to decrease culturability) did not enter the VBNC state. This indicates that transition to the dormant state was not the major strategy for cells to maintain viability under these conditions. In contrast, the bust-and-boom strategy is compatible with our results: The persistence and subsequent colonization of other environments would be facilitated by mechanisms that allow the survival of the most persistent cells at the expense of dying cells.

Several authors have demonstrated that biofilm formation is closely linked to stress responses (Hall-Stoodley et al. 2004; Landini 2009) as it can increase the capacity to withstand environmental challenges, including desiccation (Costerton et al. 1999; Espinal et al. 2012; Marks et al. 2014). We observed that the survivability of A. baumannii strains used in this study was clearly dependent on the cell distribution in the environment (free cells in saline solution vs retained on filters). While cells on solid surfaces showed a high capacity for survival under all conditions, the response to aqueous environments was variable and dependent on temperature. We also found that planktonic bacteria under starvation possessed a lower adhesiveness in contrast to those retained on filters, thus indicating that non-planktonic bacteria could be more stable per se. Moreover, the expression of biofilm-related genes declined under all conditions, even though A. baumannii maintained viability for a long time. This suggests that starved populations could find difficulties to adhere to solid surfaces and develop biofilms and, therefore, were unable to spread and colonize abiotic surfaces. As suggested by Chang et al. (2014), the downregulation of genes playing less important roles in cell survival (e.g., ompA, bfmR and csuAB) should allow to save the energy required for the expression of those genes involved in adaption to environmental stress. Moreover, the low expression level of ompA, *bfmR* and *csuAB* does not seem to completely prevent cell adhesion and subsequent biofilm formation and therefore



Fig. 4 Time-dependent changes in the expression level of the *ompA*, *bfmR* and *csuAB* genes determined for *Acinetobacter baumannii* ATCC 19606^T populations maintained in aqueous environment (*filled rectangle*) and under dry conditions (*open rectangle*) at 20 and 37 °C. Relative values were calculated with respect to the initial time (0 days)

could explain why bacteria maintained under water-free conditions still preserved their adhesiveness, especially at 37 °C.

In summary, A. baumannii shows a great persistence under stress, although starvation and physical environment differentially affect its survival. The dissemination of this pathogen seems to be based on the ability of the bulk of the bacterial population to tolerate and overcome the negative effects of stress factors. We found that the adaptation process did not initiate the entry of A. baumannii cells into the VBNC state. Moreover, the populations maintained on dry surfaces were especially resistant to stress and were able to preserve not only cell culturability but also the cellular appearance and nearly unaltered RNA content. In contrast, the survival of bacteria in aqueous environments is temperature dependent and involves changes in viability, morphology and RNA content, as well as a loss of the ability to adhere to solid surfaces. Thus, while being a common stress factor, dryness seems to have a protective role for A. baumannii cells.

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