

Antimicrobial Activity of Gallium Protoporphyrin IX against *Acinetobacter baumannii* Strains Displaying Different Antibiotic Resistance Phenotypes

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A paucity of effective, currently available antibiotics and a lull in antibiotic development pose significant challenges for treatment of patients with multidrug-resistant (MDR) *Acinetobacter baumannii* infections. Thus, novel therapeutic strategies must be evaluated to meet the demands of treatment of these often life-threatening infections. Accordingly, we examined the antibiotic activity of gallium protoporphyrin IX (Ga-PPIX) against a collection of *A. baumannii* strains, including nonmilitary and military strains and strains representing different clonal lineages and isolates classified as susceptible or MDR. Susceptibility testing demonstrated that Ga-PPIX inhibits the growth of all tested strains when cultured in cation-adjusted Mueller-Hinton broth, with a MIC of 20 $\mu\text{g/ml}$. This concentration significantly reduced bacterial viability, while 40 $\mu\text{g/ml}$ killed all cells of the *A. baumannii* ATCC 19606^T and ACICU MDR isolate after 24-h incubation. Recovery of ATCC 19606^T and ACICU strains from infected A549 human alveolar epithelial monolayers was also decreased when the medium was supplemented with Ga-PPIX, particularly at a 40- $\mu\text{g/ml}$ concentration. Similarly, the coinjection of bacteria with Ga-PPIX increased the survival of *Galleria mellonella* larvae infected with ATCC 19606^T or ACICU. Ga-PPIX was cytotoxic only when monolayers or larvae were exposed to concentrations 16-fold and 1,250-fold higher than those showing antibacterial activity, respectively. These results indicate that Ga-PPIX could be a viable therapeutic option for treatment of recalcitrant *A. baumannii* infections regardless of the resistance phenotype, clone lineage, time and site of isolation of strains causing these infections and their iron uptake phenotypes or the iron content of the media.

Immediately following the advent of antibiotics as therapeutic agents, resistance to these drugs emerged among pathogenic bacteria. Drug-resistant bacterial strains are selected for immediately after initiation of antibiotic regimens, and as a consequence of continued selective pressure, very few treatment options now exist for infections caused by resistant strains of some pathogens (1, 2). The continuing battle with resistance has led to the initial emergence of pathogens displaying multidrug-resistant (MDR) phenotypes, which has since been followed by the emergence of extremely drug-resistant (XDR) or totally drug-resistant (TDR) strains, an outcome that has recreated the preantibiotic era (3, 4). This crisis has involved major Gram-positive and Gram-negative pathogens, including *Enterococcus* spp., *Staphylococcus aureus*, members of the family *Enterobacteriaceae*, *Neisseria gonorrhoeae*, *Pseudomonas aeruginosa*, and *Acinetobacter* spp. (4). The emergence of MDR strains of each of these microorganisms has contributed to increased morbidity and mortality among patients, leading to extended lengths of hospital stay and exorbitant health care financial burdens that often go unremitted. Compounding this continuum of resistance evolution and treatment failures is the stagnation in the development of new antimicrobial agents used to treat infectious diseases caused by multidrug-resistant organisms (MDROs) (3, 4). All of the aforementioned factors have resulted in a call to action from the Infectious Diseases Society of America (5) and highlight the critical need for alternative therapeutic options that could be used alone or in combination with standard antimicrobials for the treatment of infections caused by MDROs.

Iron acquisition has been considered an alternate target of antimicrobial agents, largely because of the critical role that iron

plays in the physiology of bacteria, including pathogens that cause severe human infections (6, 7). Thus, siderophore-mediated iron acquisition processes have been targeted for the development of sideromycins, derivatives in which siderophores have been covalently linked to antibiotics, such as albomycin, salmycin, ferrymycins, the siderophore monosulfactam BAL30072, and a biscatecholate-mono-hydroxamate-carbacephalosporin conjugate (8–10). This “Trojan Horse” strategy has also included the use of gallium(III) as an anti-infective agent. Gallium binds to virtually all biological complexes that normally contain Fe(III), but it is not reduced to Ga(II) under physiological conditions and thus disrupts essential redox-driven biological processes (11). Accordingly, Ga has been used as simple inorganic and organic salts or complexed with organic compounds, including bacterial siderophores and porphyrins, such as protoporphyrin IX (12, 13). For example, gallium nitrate and gallium desfer-

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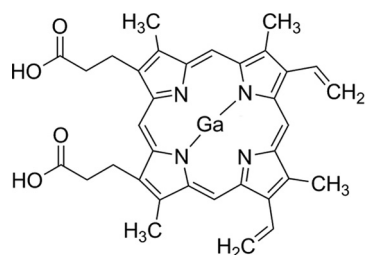


FIG 1 Molecular structure of gallium protoporphyrin IX (Ga-PPIX).

rioxamine showed antimicrobial activity against relevant human pathogens such as *P. aeruginosa* by killing free-living bacteria, blocking biofilm formation, and targeting iron metabolism (14, 15). Noniron metalloporphyrins have also been tested as antibacterials with Ga-protoporphyrin IX (Ga-PPIX) effecting the most potent action against Gram-positive and Gram-negative bacteria. The biological activity of this derivative (Fig. 1) seems to involve high-affinity uptake pathways, which depend on the expression of hemin/hemoglobin receptors, or low-affinity transport processes, whose mechanism of action are unknown (16). Furthermore, Ga-PPIX showed antimicrobial activity against *N. gonorrhoeae* when used topically in a murine vaginal infection model (17).

The Gram-negative obligate aerobic *Acinetobacter baumannii* is an opportunistic pathogen capable of causing a wide range of infections in humans, particularly in those with impaired immune systems (18, 19). The inherent and acquired resistance of clinical isolates of this pathogen complicates the severity of *A. baumannii* infections and their treatment because of the emergence of isolates that are resistant to all commercial antibiotics during the course of treatment (20, 21). This necessitates the investigation into novel chemotherapeutics needed for the treatment of infections, particularly those caused by MDR *A. baumannii* isolates obtained from patients with nosocomial infections and wounded patients. In spite of the increased occurrence of these infections, little is known about the pathobiology of this relevant pathogen. One of the best-understood interactions of this pathogen with the human host is its ability to acquire essential iron during the infection process. The results of genetic and functional analyses of the *A. baumannii* ATCC 19606 type strain showed that it acquires iron via the acinetobactin-mediated siderophore system, which proved to be critical for the infection of human alveolar epithelial cells, *Galleria mellonella* caterpillars, and mice (22). Furthermore, preliminary experimental data (23) and *in silico* genomic studies (24) indicate that *A. baumannii* expresses hemin acquisition systems that remain to be characterized experimentally. These observations suggest that Ga-containing derivatives could be effective microbicides against *A. baumannii* MDR strains. Accordingly, gallium maltolate and gallium nitrate showed antibacterial activity in mice and *G. mellonella* larvae experimentally infected with different *A. baumannii* clinical strains, including MDR isolates, although the *in vitro* activity of these derivatives depended on the free-iron content of the media (25–27). Furthermore, the analysis of *A. baumannii* LAC-4, which is considered a hypervirulent clinical isolate, showed that it is resistant to gallium nitrate except when it is in the presence of hemin, which it can use as an alternative iron source. Interestingly, the hemin utilization phenotype of the LAC-4 strain is abrogated when Ga-PPIX is added to medium

containing serum (28). In addition, the formation of Ga-siderophore complexes, such as those formed with staphyloferrin, could not result in effective antimicrobial activity (13). On the basis of these observations, we studied the effect of Ga-PPIX as an antimicrobial agent against a collection of *A. baumannii* clinical isolates that include strains obtained from military personnel with wound infections. Our results show that Ga-PPIX is an effective antimicrobial agent when tested under *in vitro*, *ex vivo*, and *in vivo* conditions independently of the antimicrobial resistance phenotype of the tested strains.

MATERIALS AND METHODS

Bacterial strains and growth conditions. All strains used in this work, which are listed in Table S1 in the supplemental material, were stored at -80°C as Luria-Bertani (LB) glycerol stocks. All strains were routinely cultured in LB agar or broth at 37°C . For antimicrobial susceptibility determinations, strains were passaged on 5% sheep blood agar prior to growth in cation-adjusted Mueller-Hinton broth (CAMHB; Oxoid, Basinstoke, Hampshire, England) as previously described (29). Iron-rich and iron-chelated conditions were achieved by adding $100\ \mu\text{M}$ FeCl_3 and the synthetic iron chelator 2,2'-dipyridyl (DIP), respectively, to the culture media. The identities of the *A. baumannii* isolates listed in Table 1 were confirmed by matrix-assisted laser desorption ionization–time of flight mass spectrometry (MALDI-TOF MS) using a Bruker microflex analyzer. The MALDI Biotyper v3.1 software package (Bruker Daltonics, Billerica, MA) was used to characterize MS fingerprints comprised of the means of 240 mass spectra profiles for each isolate analyzed. Score values greater than 2,000 were considered acceptable for species-level identification. All *A. baumannii* strains provided by D. Zurawski (Walter Reed Army Institute of Research [WRAIR]) were described previously (30). A 10-mg/ml stock solution of Ga-PPIX (Frontier Scientific, Logan, UT), for use whenever Ga-PPIX was required, was prepared using cell culture-grade dimethyl sulfoxide (DMSO) (Sigma-Aldrich, St. Louis, MO) and then sterilized using $0.2\text{-}\mu\text{m}$ nylon membrane filters (Pall Corp., Port Washington, NY).

AST. Strains were subjected to automated and manual antimicrobial susceptibility testing (AST) of standard antimicrobials and the experimental Ga-PPIX. MIC breakpoints and categorical interpretations for conventional antibiotics were mainly based on Clinical and Laboratory Standards Institute (CLSI) breakpoints (31). However, the European Committee on Antimicrobial Susceptibility Testing (EUCAST) breakpoints (http://www.eucast.org/fileadmin/src/media/PDFs/EUCAST_files/Breakpoint_tables/Breakpoint_table_v_3.1.pdf) were used to interpret the MIC of gentamicin for *A. baumannii* 19606^T. Currently, there are no breakpoints for tigecycline or Ga-PPIX.

Epsilometer (Etest)-based MIC tests (bioMérieux, Durham, NC) were performed and interpreted according to the manufacturer's instructions. Briefly, suspensions of the strains equivalent to a 0.5 McFarland standard were used to inoculate Mueller-Hinton agar plates (Remel, Lenexa, KS) prior to strip placement. The plates were incubated for 22 h in ambient air at 35°C prior to MIC interpretation. The MIC was interpreted by observing the concentration of the drug on the strip where the growth of the organism intersected the strip.

Isolate AST of Ga-PPIX and doxycycline was carried out by the Kirby-Bauer disk diffusion method according to standard procedures (32). Ga-PPIX stock solution was added to sterile 6-mm paper disks to deliver $50\ \mu\text{g}$ or $100\ \mu\text{g}$ of Ga-PPIX per disk and allowed to dry. Ga-PPIX paper disks and HardyDisk (Hardy Diagnostics, Franklin, OH) containing $30\ \mu\text{g}$ doxycycline were applied to the surfaces of Mueller-Hinton agar plates inoculated with bacteria. Zones of inhibition (ZOI) were measured after incubation in ambient atmosphere at 37°C for 24 h. ZOI of doxycycline were classified as resistant, intermediate, or susceptible based on CLSI standards.

For automated AST determinations, the Vitek 2 XL (bioMérieux) microbial identification system and the BD Phoenix (BD Diagnostic Sys-

tems, Sparks, MD) automated microbiology system were used according to the manufacturer's instructions. The bioMérieux susceptibility card AST-GN73 and Phoenix NMIC/ID-130 panels were used for each platform, respectively. Interpretive criteria were based on CLSI breakpoints (31).

Broth microdilution MICs. The MIC of Ga-PPIX was determined using the antimicrobial broth microdilution technique in the absence or presence of 10% heat-inactivated normal serum according to standard procedures (29). Briefly, broth microdilution plates were inoculated with 10⁵ CFU/ml of each test organism in CAMHB and incubated for 24 h at 37°C in ambient air. Twofold dilutions of Ga-PPIX were performed; the concentrations ranged from 5 µg/ml to 166 µg/ml. A sample with no added antibiotic was included as a growth control on each plate. A more defined Ga-PPIX MIC was determined with 1-µg/ml resolution covering a range of 10 µg/ml to 20 µg/ml for selected strains. The wells on the plates were read for turbidity at an optical density at 595 nm (OD₅₉₅) with a FilterMax F5 instrument using the multimode analysis software version 3.4.0.25 (Molecular Devices, Sunnyvale, CA). The MIC was determined as the microtiter plate's well with the lowest drug concentration at which there was no detectable growth. The MICs were determined at least three times in triplicate (n = 9) using fresh biological samples each time.

Ga-PPIX resistance frequency. CAMH broth samples were inoculated with a 1:100 dilution of 24-h cultures of ATCC 19606^T or ACICU strain and then incubated for 6 h at 37°C. Bacteria were plated onto CAMH agar plates containing 2× (40 µg/ml) and 4× (80 µg/ml) the MIC of Ga-PPIX. The population of bacteria treated with Ga-PPIX was quantified by plate count. Colonies were observed after 24 h and used to determine mutation frequency. The assay was performed twice in triplicate.

Time-kill assays. Time-kill kinetics assays were performed to determine the rapidity of bactericidal activity of Ga-PPIX. The time-kill kinetic assays were performed basically as described previously (33). Briefly, after routine growth, 10⁶ CFU/ml of *A. baumannii* ATCC 19606^T or ACICU were inoculated into CAMHB. Cultures were then treated with 0, 10, 20, or 40 µg/ml Ga-PPIX, concentrations that corresponded to 0, 0.5, 1, or 2 times the MIC, respectively. The numbers of CFU were determined at 0, 2, 4, 6, and 24 h postinoculation by plate count. Assays were performed twice in triplicate (n = 6) using fresh biological samples each time.

A549 ex vivo assays. To test the cytotoxicity of Ga-PPIX, an opaque white 96-well plate was seeded with 10⁴ A549 human pulmonary adenocarcinoma cells and incubated for 24 h in Dulbecco's modified Eagle's medium (DMEM) (Mediatech, Inc., Manassas, VA) supplemented with 10% heat-inactivated fetal bovine serum (HyClone, Logan, UT), penicillin, and streptomycin at 37°C in a 5% CO₂ atmosphere as previously described (22). Following incubation, the medium was exchanged for DMEM containing 10% heat-inactivated fetal bovine serum without antibiotics and supplemented with twofold dilutions ranging from 10 µg/ml to 640 µg/ml Ga-PPIX. The samples were incubated for 24 h, and cell viability was assayed using the CellTiter-Glo luminescent cell viability assay (Promega, Madison, WI). Luminescence was measured for 10 ms with a FilterMax F5 instrument. Experiments were performed twice in octuplet (n = 16). Groups were compared by analysis of variance (ANOVA) with Tukey's multiple-comparison test.

To test the antimicrobial effect of Ga-PPIX on the infection of submerged A549 cell monolayers, 24-well tissue culture plates were seeded with approximately 10⁴ cells per well and then incubated for 16 h as described before (22). Bacteria were cultured for 24 h in CAMHB at 37°C with shaking at 200 rpm, collected by centrifugation at 21,000 × g for 10 min, washed, resuspended, and diluted in DMEM with 10% heat-inactivated fetal bovine serum without antibiotics or supplemented with 20 or 40 µg/ml Ga-PPIX. The A549 cell monolayers were infected with 10⁴ CFU of the *A. baumannii* ATCC 19606^T or ACICU strain. Inocula were estimated spectrophotometrically at OD₆₀₀ and then confirmed by plate count. Infected-cell monolayers were incubated for 24 h at 37°C in 5% CO₂. The tissue culture supernatants were collected, the A549 monolayers were lysed with sterile distilled H₂O, and the lysates were added to the

TABLE 1 Antibiotic susceptibility profile of nonmilitary *A. baumannii* strains^a

| Strain | Penicillins | | | Cephalosporins | | | Carbapenems | | | Fluoroquinolones | | | Aminoglycosides | | | Tetracyclines | | | Miscellaneous | | |
|-------------------------|-------------|-----|-----|----------------|-----|-----|-------------|-----|-----|------------------|-----|-----|-----------------|-----|-----|---------------|-----|-----|---------------|---------|--|
| | AMP | SAM | TZP | CEZ | FEP | FOX | CAZ | CRO | IMP | MEM | CIP | LVX | AMK | GEN | TOB | TGC | DOX | CST | SXT | Ga-PPIX | |
| ATCC 19606 ^T | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| 17978 | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| LUH07672 | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| LUH08809 | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| LUH05875 | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| LUH13000 | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| RUIH0134 | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| RUIH0075 | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| AYE | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| SDF | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| A118 | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |
| ACICU | R | S | S | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | |

^a Antibiotic abbreviations: AMK, amikacin; AMP, ampicillin; CAZ, ceftazidime; CEZ, ceftazolin; CIP, ciprofloxacin; COX, ceftriaxone; CST, colistin; FEP, ceftepime; FOX, ceftoxitin; Ga-PPIX, gallium protoporphyrin IX; GEN, gentamicin; IMP, imipenem; LVX, levofloxacin; MEM, meropenem; SAM, ampicillin-sulbactam; SXT, trimethoprim-sulfamethoxazole; TGC, tigecycline; TOB, tobramycin; TZP, piperacillin-tazobactam.
^b The MIC values are shown in micrograms per milliliter.
^c The categorical interpretations (CI) are as follows: R, resistant; S, susceptible; I, intermediate; ND, not determined. Asterisks indicate the absence of interpretive standards.
^d MIC determined by Vitek-2 AST-GN73 card.
^e MIC determined by Elest.
^f MIC determined by BD Phoenix NMIC/ID-130 panel.
^g *, no interpretive standards available.
^h Susceptibility determined by Kirby-Bauer disk diffusion.
ⁱ MIC determined by microdilution.

cognate tissue culture supernatants. Combined lysate samples were serially diluted and then plated on nutrient agar. After overnight incubation at 37°C, the numbers of CFU were counted, and the viable-cell plate count for each sample was calculated and recorded. Experiments were performed twice in octuplet ($n = 16$) using fresh biological samples each time and compared by ANOVA with Tukey's multiple-comparison test.

***Galleria mellonella* in vivo assays.** Ga-PPIX's toxicity and efficacy to treat experimental infections were tested using final instar larvae of the greater wax moth, *G. mellonella* (Grubco Inc., Fairfield, OH) as described before (22). Briefly, larvae weighing between 0.25 g and 0.35 g were randomly assigned to groups of 10 for each trial and then injected into the last left proleg using a syringe pump (New Era Pump Systems Inc., Farmingdale, NY) and 26.5-gauge needles. Larvae were inspected for survival every 24 h for 5 days. Dead larvae were removed at the time of inspection. If more than two larvae died in any control group, the results of the cognate trial were excluded from analysis, and the trial was repeated. Trials were performed in triplicate ($n = 30$).

Ga-PPIX *in vivo* toxicity was tested by using mass-matched *G. mellonella* larvae (i.e., mean mass of each group was statistically indistinguishable by ANOVA from any other group). Larvae were injected with 5 μ l of twofold dilutions ranging from 0.2 mM to 25 mM (15.7 mg/ml to 126 μ g/ml) Ga-PPIX solubilized in 0.5 N NaOH and 1.5% DMSO. Control groups were either injected or not injected with carrier solution. Probit analysis was performed using R 3.0 (www.r-project.org). To test antimicrobial activity, larvae were injected with a 5- μ l bolus of 10^5 CFU of *A. baumannii* ATCC 19606^T or *A. baumannii* ACICU suspended in a phosphate-buffered saline (PBS) solution containing 2% DMSO (carrier) or carrier supplemented with either 20 μ g/ml or 40 μ g/ml Ga-PPIX. Uninjected larvae and larvae injected with the same volume of sterile carrier were used as controls. Kaplan-Meier survival analysis and odds ratios were determined using Prism 6.0 (GraphPad Software, Inc., La Jolla, CA).

RESULTS

Source and antibiotic susceptibility profiles of *A. baumannii* strains used in this study. Ga-PPIX proved to be an effective antimicrobial when tested against Gram-negative bacteria (*Yersinia enterocolitica*), Gram-positive bacteria (*Staphylococcus aureus*), and acid-fast bacilli (*Mycobacterium smegmatis*) (16). In spite of initial promising results, the application of Ga-PPIX to other pathogens has not been systematically tested. Thus, we used an approach similar to that described in the aforementioned publication (16) to test the Ga-PPIX susceptibility of an *A. baumannii* strain collection that included 30 *A. baumannii* strains isolated from U.S. wounded military personnel (strains 3132 to 3160 and 3284 in Table S1 in the supplemental material) and 12 nonmilitary isolates representing type strains, different international clonal lineages, strains whose genomes were fully sequenced and annotated, and strains that were classified as antimicrobial susceptible or MDR (strains 406 to 3022 and 3161 in Table S1). We initially tested the antibiotic susceptibility of all non-WRAIR strains, the identities of which were confirmed using a combination of standard bacteriological methods and modern technology, such as MALDI-TOF MS. This analysis showed that this set of strains includes isolates, such as SDF, that are susceptible to all tested antibiotics, which included more than six different classes of antimicrobials; isolates that display resistance to some antibiotics, such as ATCC 19606^T and ATCC 17978; MDR isolates, such as LUH07672, LUH08809, LUH05875, LUH13000, RUH00134, and RUH00875; and isolates that are resistant to practically all drugs, such as AYE and ACICU, with ACICU being the most resistant isolate used in this work (Table 1). The U.S. military strain collection of clinical isolates includes 30 strains, each of which are resistant to most of the 11 antibiotics tested (30).

Growth inhibitory effects of Ga-PPIX. All 42 tested strains displayed apparent growth when cultured in CAMHB at 37°C for 18 h using 96-well microtiter plates, although there was some growth variability among the strains (see Table S2 in the supplemental material). The addition of Ga-PPIX produced a dose-dependent growth response with several strains growing to higher optical densities in the presence of 5 μ g/ml to 20 μ g/ml of this nonferrous metalloporphyrin when assayed using an automated twofold microdilution method. However, bacterial growth was significantly reduced when the CAMHB contained more than 20 μ g/ml Ga-PPIX, making 40 μ g/ml the MIC for all 42 strains when tested using this automated approach, which resulted in inocula containing 10^6 bacteria per sample. A more detailed manual analysis of the nonmilitary isolates following the CLSI standards (using inocula of 10^5 bacteria and CAMHB) using a twofold Ga-PPIX serial dilution scheme resulted in a MIC value of 20 μ g/ml for all of these isolates. Furthermore, a 1- μ g/ml resolution scheme performed under the latter experimental conditions resulted in Ga-PPIX MIC values ranging between 11 μ g/ml and 20 μ g/ml (Table 1). The median and mean MIC of these nonmilitary isolates was 19 μ g/ml and 17.4 μ g/ml, respectively. The presence of 10% heat-inactivated normal human serum increased the MIC for the ATCC 19606^T and ACICU strains to 62.5 μ g/ml, a threefold increase compared with the values obtained in the absence of proteins, without causing the complete inactivation or sequestration of Ga-PPIX. The Ga-PPIX susceptibility of these isolates was also tested using disk diffusion assays. All strains tested produced detectable ZOI ranging between 13 mm and 23 mm in diameter when the seeded plates were exposed to sterile filter disks impregnated with 50 μ g or 100 μ g Ga-PPIX, respectively (Fig. 2), and there were no heteroresistant colonies noted in the ZOI, which is represented in the insets of Fig. 2A and B. This observation is in agreement with the failure to isolate Ga-PPIX-resistant colonies of ATCC 19606^T and ACICU when a population of greater than 9.8×10^9 bacteria were challenged to $2 \times$ (40 μ g/ml) or $4 \times$ (80 μ g/ml) the MIC of Ga-PPIX (data not shown).

Comparable results were obtained when the disk assays were conducted using LB agar plates that were supplemented with FeCl₃ or DIP to generate iron-rich or iron-chelated conditions, respectively (data not shown). Furthermore, comparable Ga-PPIX susceptibility responses were detected with the ATCC 19606^T strain and the isogenic derivatives ATCC 19606^T t6 (strain 1653 in Table S1 in the supplemental material) and ATCC 19606^T entA (strain 3069 in Table S1), which are affected in the uptake and biosynthesis of acinetobactin because of mutations in the genes coding for the BauA acinetobactin receptor protein and the EntA biosynthetic protein, respectively (34, 35). Additionally, the detailed analysis of *A. baumannii* AB5075 (strain 3156 in Table S1), a MDR isolate recently cultured from an infected wound that has been proposed as a model strain to study the pathobiology of this pathogen (30), showed a Ga-PPIX MIC of 20 μ g/ml when cultured in CAMHB under standard laboratory conditions following CLSI guidelines (Fig. S1A and S1B in the supplemental material).

These experimental data collected using two different methods show that the susceptibility of different *A. baumannii* clinical strains to Ga-PPIX, with a standard MIC of 20 μ g/ml, is independent of their time and site of isolation and their overall antimicrobial resistance phenotypes and clonal lineages. The data also indicate that the susceptibility to Ga-PPIX is independent of the

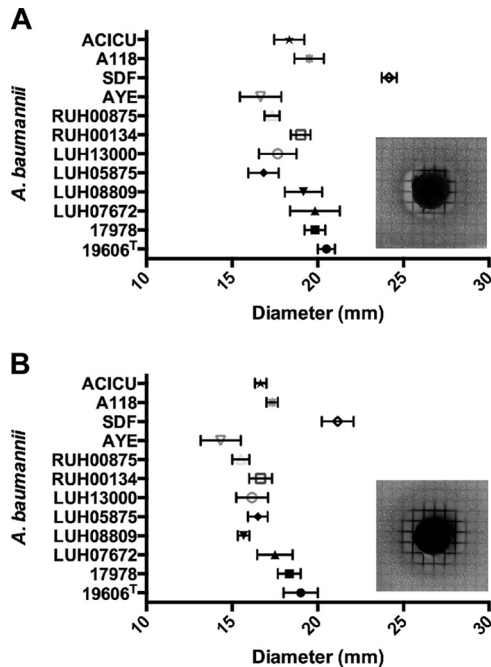


FIG 2 Ga-PPIX disk diffusion assays. *A. baumannii* bacteria were first seeded onto CAMH agar and then disks impregnated with 50 µg (A) or 100 µg (B) of Ga-PPIX were deposited on the surfaces of the plates. Growth inhibition halos were measured after incubation for 24 h at 37°C. Data are expressed as means \pm standard errors of the means (SEM) (error bars) from experiments done twice in triplicate. The insets show the growth inhibition halos observed when the CAMH agar plates were seeded with *A. baumannii* ATCC 19606^T and incubated for 24 h at 37°C.

free-iron content of the medium and the capacity of bacteria to produce or use acinetobactin, which seems to be the most common siderophore-mediated iron uptake system found in the genome of different *A. baumannii* clinical isolates (36).

Ga-PPIX time-kill kinetics. The antibacterial activity of Ga-PPIX was tested by determining its time-kill kinetics using the ATCC 19606^T and ACICU nonmilitary isolates, which are considered non-MDR and MDR strains, respectively, at a concentration of 0.5, 1, and 2 times the MIC as determined by the microdilution method. Both *A. baumannii* strains showed similar responses, with the addition of 10 µg/ml Ga-PPIX (0.5× MIC) causing only a small reduction in bacterial viability (Fig. 3). In contrast, the addition of 20 µg/ml (1× MIC) or 40 µg/ml (2× MIC) was effective in reducing the viable cells by >3 log units after 6 h of incubation. Furthermore, 1 × MIC did lead to a statistically significant reduction of 8×10^5 - and 2.6×10^3 -fold in viable cells after 24 h compared with the cognate untreated samples of ATCC 19606^T and ACICU, respectively. Interestingly, 40 µg/ml of Ga-PPIX caused complete bacterial cell death for both the ATCC 19606^T strain and ACICU MDR isolate. These results indicate that Ga-PPIX is both fast and effective in eliminating viable *A. baumannii* under *in vitro* conditions normally used to test the effectiveness of antimicrobial agents.

Ga-PPIX toxicity to eukaryotic cells and *G. mellonella* larvae. The toxicity of Ga-PPIX was tested using A549 alveolar epithelial tissue culture cells and *G. mellonella* caterpillars. Ga-PPIX exhibited no statistically significant toxicity for A549 cells at concentrations up to 160 µg/ml, 8 times the MIC, when tested using the

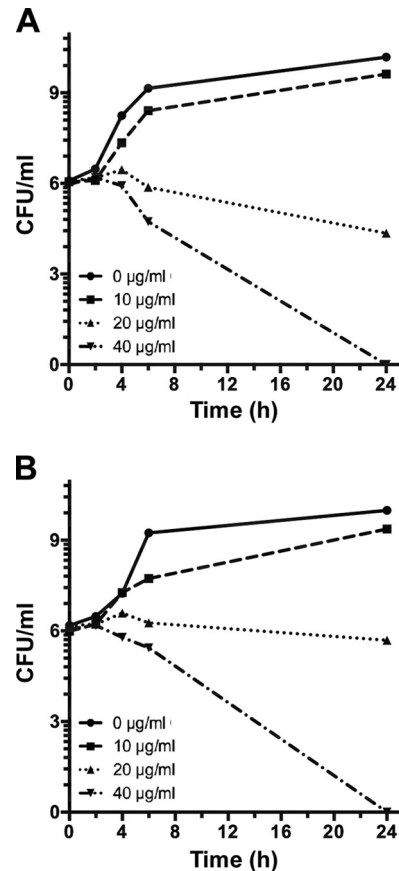
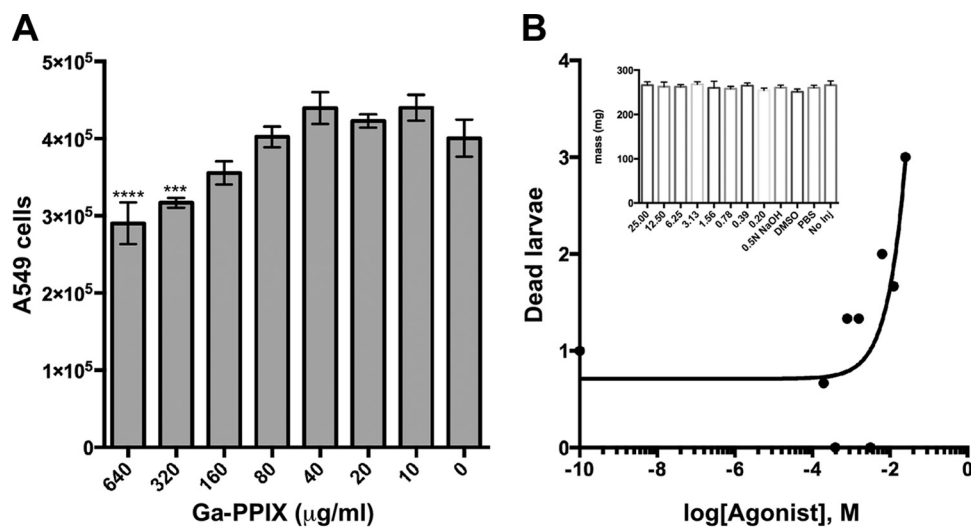


FIG 3 Ga-PPIX time-kill assays. Ga-PPIX time-kill kinetics for *A. baumannii* ATCC 19606^T (A) and ACICU (B) was determined after 0, 2, 4, 6, and 24 h of incubation by CFU count after exposure to 0 µg/ml (0× MIC), 10 µg/ml (0.5× MIC), 20 µg/ml (1× MIC), or 40 µg/ml (2× MIC) Ga-PPIX.

CellTiter-Glo luminescent cell viability assay. This assay also showed that there was a statistically significant reduction in A549 viability when the culture medium contained 640 µg/ml ($P \leq 0.0001$) or 320 µg/ml ($P = 0.004$) Ga-PPIX (Fig. 4A), concentrations that are far higher than the MIC values reported in the previous section. The *ex vivo* toxicity is corroborated by *in vivo* experiments using *G. mellonella* larvae (Fig. 4B) with comparable body mass (Fig. 4B, inset). After injection with Ga-PPIX over a 3-log-unit range (200 µM to 25 mM), the survival of larvae injected with 25 mM was significantly reduced than survival in the control group ($P = 0.0363$) (data not shown). The 50% lethal concentration (LC₅₀) for Ga-PPIX in *G. mellonella* was extrapolated by probit analysis from the log dose-response curve to be 157 mg/ml (Fig. 4B). It was also observed that the injection of high concentrations of Ga-PPIX resulted in immediate coloration of the larvae, and subsequently excreted feces were noticeably pink (data not shown).

These results demonstrate the ability of eukaryotic cells and organisms to tolerate Ga-PPIX at concentrations much higher than the MIC values determined using standard methods.

Ga-PPIX treatment of *ex vivo* and *in vivo* experimental infections. The antibacterial effectiveness of Ga-PPIX was determined by infecting A549 tissue cultures and *G. mellonella* larvae with the *A. baumannii* ATCC 19606^T strain or the ACICU MDR



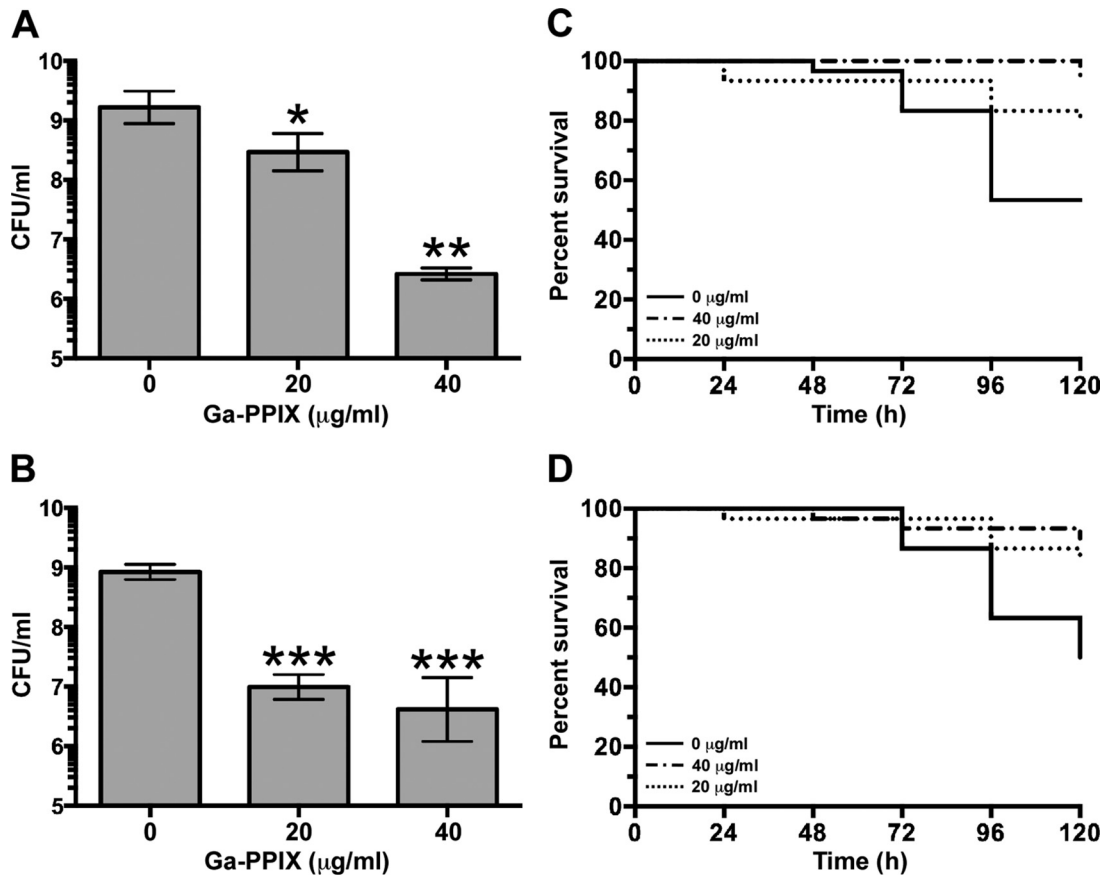


FIG 5 Antibacterial activity of Ga-PPIX against experimental infections. Monolayers of A549 human alveolar epithelial cells (A and B) and *G. mellonella* larvae (C and D) were infected with *A. baumannii* ATCC 19606^T strain (A and C) or ACICU strain (B and D) in the absence or the presence of either 20 µg/ml or 40 µg/ml of Ga-PPIX. Data shown in panels A and B represent the means (± SEM) of experiments performed twice in octuplet ($n = 16$) using fresh biological samples each time. Responses to the presence of Ga-PPIX were compared to those detected with bacteria cultured in the absence of this metalloporphyrin derivative. Values that are significantly different from the value for the control (0 µg/ml Ga-PPIX) are indicated by asterisks as follows: *, $P = 0.037$; **, $P = 0.014$; ***, $P = 0.0005$.

dent of the free-iron content of the media and the active expression of siderophore-mediated iron acquisition functions, while the presence of proteins in the *in vitro*, *ex vivo*, and *in vivo* susceptibility assays showed that they do not drastically affect the antibacterial activity of Ga-PPIX. Our data also showed that Ga-PPIX displays bactericidal activity relatively quickly, particularly when used at twice the MIC (40 µg/ml) under standard laboratory conditions, without causing any detectable cytotoxic effects on human cells and *G. mellonella* caterpillars. These findings are relevant since patients suffering from severe wound and soft tissue infections require immediate and effective antimicrobial chemotherapy. Equally concerning is the emergence of *A. baumannii* isolates with increased and/or altered virulence, such as the LAC-4 strain and isolates obtained from deadly cases of necrotizing fasciitis, respectively (37, 38). These infections require immediate and effective antimicrobial treatment not only because of the devastating nature of these infections but also because of the alarming MDR phenotype of the bacterial agents responsible for these diseases.

Our recent observation that the Ga-PPIX-susceptible strain AYE produces the hydroxamate siderophore baumannoferrin (39), but not acinetobactin or any catechol-derived siderophore due to a natural *entA* natural mutation (35), shows that the anti-

bacterial activity of this nonferroc metalloporphyrin extends beyond those strains that produce and use acinetobactin to acquire iron. More recently, it was reported that the *A. baumannii* ATCC 17978 strain, which could rely on the production and use of acinetobactin and fimsbactins to grow under free-iron limiting conditions (40), is also susceptible to Ga-PPIX. These observations further demonstrate that the antibacterial activity of Ga-PPIX is independent of the type or number of high-affinity iron chelators produced by a particular isolate. Whether these conditions also apply to the LAC-4 hypervirulent strain, which proved to be susceptible to Ga-PPIX (28), is currently unknown, since the siderophore-mediated iron acquisition system or systems expressed by this isolate is/are unknown.

The fact that Ga-PPIX displays bactericidal activity even in a medium such as LB agar, which has enough free iron to inhibit the expression of the acinetobactin system in the ATCC 19606^T strain (41), demonstrates the antibacterial effectiveness of this noniron metalloporphyrin derivative even under conditions that inhibit the expression of critical iron-regulated bacterial functions, including those required for iron acquisition via siderophore or hemin uptake processes. However, the mechanisms by which Ga-PPIX is transported into the bacterial cytoplasm and affects iron metabolism are not fully understood. Pioneering work done by

Stojiljkovic et al. showed that Gram-negative bacteria, such as *Yersinia enterocolitica*, are susceptible to Ga-PPIX when expressing hemin acquisition functions under low-iron conditions (16). Similarly, work done by Moriwaki et al. showed that the *Staphylococcus aureus* IsdH-NEAT3 hemin receptor interacts with Ga-PPIX in a fashion similar to that of the natural hemin ligand (42). However, these results contradict our observations showing that Ga-PPIX displays antibacterial activity against bacteria cultured in iron-rich media such as LB agar or broth, which should significantly inhibit the expression of hemin transport functions according to data collected with *Y. enterocolitica* and *Escherichia coli* (16). Taken together, these results argue in favor of the hypothesis that Ga-PPIX could reach the periplasmic and cytoplasmic spaces either by using unknown low-affinity or TonB-independent hemin receptors or passive diffusion through bacterial membranes (16). Those hypotheses are supported by our data showing that the *A. baumannii* strains AYE and ATCC 17978, which lack the eight-gene cluster coding for putative hemin oxygenase and hemin utilization proteins recently described by de Léséleuc et al. (28), and ATCC 19606^T strain, which does not have an identifiable *hemO* ortholog and therefore lacks this cluster, were as susceptible as the ACICU, SDF, and AB0057 strains, all of which harbor the aforementioned eight-gene cluster. The Ga-PPIX susceptibility of AYE, ATCC 17978, and ATCC 19606^T could be also explained by the presence of a 12-gene cluster (iron uptake gene cluster 2), which codes for predicted hemin uptake functions but lacks a canonical *hemO* ortholog, that is also present in the genomes of the ACICU, SDF, and AB0057 strains (24). Together, these observations indicate that the widespread susceptibility of *A. baumannii* to Ga-PPIX is mediated by more than one cellular process, a property that will most likely not favor the emergence of resistant derivatives at least in the short term as could be predicted from our work, which failed to detect a resistance phenotype using disk diffusion assays, broth MIC, or mutation rate studies.

In summary, our work shows that *A. baumannii* strains are susceptible to Ga-PPIX regardless of their clonal lineages, site and time of isolation, antimicrobial resistance phenotypes, expression of iron acquisition systems, iron content of the media, or presence of proteins in *in vitro* and *ex vivo* tissue culture susceptibility assays. All these properties make this nonferric metalloporphyrin derivative a potentially valuable and convenient antimicrobial agent that affects the viability of bacterial cells not only by interfering with critical physiological processes that depend on the redox properties of iron but also making bacterial cells more susceptible to oxidative stresses due to the presence of reactive oxygen species produced by bacteria growing under aerobic conditions as well as by the host in response to infection (16). However, the relatively low solubility of Ga-PPIX in aqueous solutions is one obstacle that must be overcome before this compound is introduced as a viable therapeutic option, either alone or in combination with current antibiotics, for the treatment of *A. baumannii* infections, particularly those caused by isolates that are highly resistant to current antimicrobial agents.

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