

Combatting Bacterial Infections by Killing Persister Cells with Mitomycin C

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ABSTRACT

Persister cells are a multi-drug tolerant subpopulation of bacteria that contribute to chronic and recalcitrant clinical infections such as cystic fibrosis and tuberculosis. Persisters are metabolically dormant, so they are highly tolerant to all traditional antibiotics which are mainly effective against 5 actively-growing cells. Here, we show that the FDA-approved anti-cancer drug mitomycin C (MMC) eradicates persister cells through a growth-independent mechanism. MMC is passively transported and bioreductively activated, leading to spontaneous crosslinking of DNA, which we verify in both active and dormant cells. We find MMC effectively eradicates cells grown in numerous different growth states (e.g., planktonic cultures and highly robust biofilm cultures) in both rich and minimal media. Additionally, ≤ 10 MMC is a potent bactericide for a broad range of bacterial persisters, including commensal *Escherichia* coli K-12 as well as pathogenic species of E. coli, Staphylococcus aureus, and Pseudomonas aeruginosa. We also demonstrate the efficacy of MMC in an animal model and a wound model, substantiating the clinical applicability of MMC against bacterial infections. Therefore, MMC is the first broad-spectrum compound capable of eliminating persister cells, meriting investigation as a new approach for the treatment recalcitrant infections. 15 of

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INTRODUCTION

There are 17 million new biofilm infections every year in the U.S.A., which lead to 550,000 fatalities (Wolcott and Dowd, 2011), and biofilms are difficult to treat due to the presence of persister cells (Lewis, 2008). Persisters arise due to metabolic inactivity (Lewis, 2007; Kwan *et al.*, 2013; Wood *et al.*, 2013) and are highly tolerant against all traditional antibiotic classes, which are primarily effective against actively growing cells. Bacterial persistence is a non-hereditary phenotype (Bigger, 1944) which occurs both stochastically (Balaban *et al.*, 2004) or through environmental influence (Dörr *et al.*, 2010; Möker *et al.*, 2010; Vega *et al.*, 2012; Kwan *et al.*, 2013; Hu *et al.*, 2014; Kwan *et al.*, 2014) in a small sub-population of all tested bacterial species (Lewis, 2008) (~1% during stationary phase and in biofilm cultures (Lewis, 2007, 2008)).

Few distinctly new antibiotics have been discovered recently (Mills and Dougherty, 2012), and current antibiotics are ineffective against persister cells. Thus, we searched for a compound which could eradicate persister cells. Mitomycin C (MMC) is used as an FDA-approved (Doll *et al.*, 1985) chemotherapeutic agent for a wide range of cancer treatments (e.g., bladder, gastric, and pancreatic) (Bradner, 2001). As an amphipathic molecule, MMC passively diffuses into cells (Byfield and Calabro-Jones, 1981). Bacterial cytoplasm is a reducing environment (Szybalski and Iyer, 1964), so after entering into cells, the quinone functional group of MMC is reduced spontaneously, initiating crosslinking of adjacent guanine residues in 5'-CG sequences to join two opposing strands of DNA (Tomasz, 1995). Because transport is passive and the reaction is spontaneous, we reasoned that MMC would be effective against cells in the persister state, a state of metabolic dormancy, since MMC activity would not require active metabolism.

We found that MMC is effective against persister cells in a broad range of bacteria including commensal *Escherichia coli* K-12 as well as pathogenic strains of *E. coli*, *Staphylococcus aureus* (frequently found in wounds), and *Pseudomonas aeruginosa*. We also demonstrated that MMC eradicates

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bacteria in biofilms, communities of notoriously difficult to treat cells present in a majority of infections. Furthermore, we verified that MMC kills persister cells by crosslinking DNA, and we demonstrated the efficacy of MMC in an animal model and in a wound model. Therefore, MMC has broad-spectrum activity against growing, non-growing, and persister cells, and should be used for the treatment of recalcitrant infections.

RESULTS AND DISCUSSION

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MMC kills active and persister cells in rich medium. MMC activity is decreased at high pH (Kennedy *et al.*, 1985); hence, we buffered the medium to avoid high pH fluctuations and to match the physiological resting pH of ~7.4 and exercising pH of ~7.1 and ~6.4 (Hermansen and Osnes, 1972). For this work, we compared MMC with ciprofloxacin, a fluoroquinolone that inhibits DNA replication and kills both growing and non-growing cells but not persister cells (Sanders, 1988) and which is commonly used in persister studies (Conlon *et al.*, 2013). Therefore, throughout this work, ciprofloxacin tolerance represents the baseline level of persistence. Additionally, antibiotic treatments were generally at least 5x the minimum inhibitory concentration (MIC) (Table 1) to ensure eradication of non-persisters and to minimize the survival of potential spontaneous resistant mutants.

As evidence that ciprofloxacin tolerance is due to persistence rather than spontaneous genetic resistance, we measured the tolerance of *E. coli* K-12 cultures after three rounds of ciprofloxacin treatment (5 µg/mL for 3 h) and subsequent regrowth of persisters in fresh media. There was no observable increase in ciprofloxacin survival after each round of regrowth (survival at 0.026 \pm 0.007%, 0.048 \pm 0.002%, and 0.044 \pm 0.001% for rounds 1, 2 and 3, respectively). In addition, no colonies were detectable with 5 µg/mL ciprofloxacin, showing absolutely that there were no resistant strains. These results confirm the reliability of using ciprofloxacin tolerance as an indicator of persistence.

We initially evaluated MMC with *E. coli* K-12 and found, compared to ciprofloxacin, that MMC was 2,300-fold more effective against exponentially-growing cells (**Fig. 1A**) and 150,000-fold more effective against mid-stationary-phase cells in buffered lysogeny broth (LB) medium (**Fig. 1B**). As evidence of the

ability of MMC to kill persister cells, we found that treatment of a late-stationary phase culture with MMC does not show the bi-phasic death curve that is characteristic of a persister population (**Fig. 1C**). We then utilized a rifampicin pretreatment which we previously demonstrated to induce high levels of persistence (~10-100%) (Kwan *et al.*, 2013) and found that MMC was highly effective against rifampicin-induced persister cells, in stark contrast to ciprofloxacin (**Fig. 1D**). Therefore, MMC kills non-persister cells and dormant persister cells.

MMC kills active and persister cells in minimal medium, in anaerobic cultures, and in biofilms. The previous assays were performed in planktonic cultures grown in rich medium; however, these growth conditions are a poor representation of ecological bacterial growth. Thus, we investigated MMC activity against cultures grown in minimal medium, in biofilms, and in anaerobic cultures. Exponential-phase cultures in M9-glucose were similarly susceptible to MMC and ciprofloxacin (**Fig. 1E**); however, during late-stationary phase in M9-glucose, we found that while the population was highly persistent against ciprofloxacin ($10 \pm 1 \%$) and the aminoglycoside gentamicin ($44 \pm 5 \%$), MMC eradicated cells (**Fig. 1F**). Biofilms more accurately model clinical bacteria growth with a high population of persisters (Lewis, 2008), and we found that MMC was effective against biofilms in M9-glucose, killing 100,000-fold more cells than ciprofloxacin and nearly eliminating cells after 24 h of treatment (**Fig. 1G**). Additionally, MMC did not cause biofilm dispersal, confirming that efficacy against biofilms was in fact due to eradication of cells (**Fig. S1**). Bacterial infections have a propensity to exist under anaerobic conditions, and we found that MMC eradicated anaerobic, late-stationary phase cells in rich medium beyond the limit of detection, in comparison to 0.44 \pm 0.08 % survival against ciprofloxacin treatment (**Fig. 1H**). Furthermore, anaerobic biofilm cultures in M9-glucose were 2,500-fold more susceptible to MMC than ciprofloxacin

(**Fig. 1I**).

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MMC kills viable but non-culturable cells. Numerous species of bacteria enter the viable but nonculturable (VBNC) state, another state of metabolic dormancy closely related to persistence (Ayrapetyan *et al.*, 2014), as a survival response to environmental stresses, and these cells do not resuscitate and become culturable unless exposed to suitable stimuli (Li *et al.*, 2014). VBNC cells exhibit high antibiotic

tolerance, similarly to persisters, and pose a risk to human health because they can avoid detection in goods, leading to infection (Li et al., 2014). Hence, we generated VBNC cultures by starving cells in saline solution for 36 days until there were \sim 1,000-fold more VBNC cells than culturable cells. Respiratory activity is a commonly used criterion for viability (Ramamurthy et al., 2014), so 95 RedoxSensor Green, a fluorescent dye for detection of actively respiring cells, was used to enumerate the VBNC population. Upon antibiotic treatment of these cultures, we found that MMC was 7-fold more effective at killing VBNC cells than ciprofloxacin (Fig. 1J), while also eradicating the culturable population (Fig. S2), unlike ciprofloxacin (0.40 ± 0.05 % survival). Therefore, we have demonstrated that 100 MMC is highly effective against metabolically dormant cells in both the persister and VBNC states.

MMC kills persister cells by crosslinking DNA. To verify that MMC kills bacteria via DNA crosslinks, we investigated MMC activity against single-gene deletion mutants lacking uvrA, uvrB, and uvrC. The UvrABC complex is part of the bacterial SOS response in E. coli (Michel, 2005) to repair DNA crosslinks (Weng et al., 2010), and contributes to MMC tolerance (Salem et al., 2009). We found that the 105 $\Delta uvrA$, $\Delta uvrB$, and $\Delta uvrC$ mutants were much more sensitive to MMC than the wild-type strain, and were rapidly eradicated (beyond the limit of detection) within less than 30 min of treatment (Fig. 2A). Additionally, we were able to complement the high sensitivity to MMC of a $\Delta uvrA$ mutant with production of UvrA via plasmid (Fig. 2B). These results confirm that DNA crosslinking is the basis for MMC bactericidal activity in actively-growing cells.

However, persister cells are dormant, thus having a different metabolic state than non-persisters. 110 Therefore, we sought to verify that MMC was in fact forming DNA crosslinks within persisters, rather than killing persisters through an unknown mechanism. Genomic DNA (gDNA) was isolated from both exponential phase cells (i.e., non-persisters) and rifampicin-induced persisters (Kwan et al., 2013) before and after MMC treatment. We hypothesized that crosslinks within DNA should inhibit amplification via 115 quantitative PCR (qPCR). qPCR was performed with primers designed to amplify a 234 nt region of rrsG and a 302 nt region of *murB*, containing 18 and 10 potential MMC crosslinking sites, respectively. As a positive control, qPCR was performed on gDNA crosslinked by MMC in vitro, verifying that DNA

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crosslinking inhibited amplification dramatically (*rrsG*: -1,456 \pm 1 fold and *murB*: -1,621 \pm 28 fold). Our *in vivo* results revealed the presence of gDNA crosslinks based on reduced quantities of PCR-amplifiable 120 DNA after MMC treatment for non-persisters (*rrsG*: -5.95 \pm 0.20 fold and *murB*: -5.13 \pm 0.14 fold) and persisters (*rrsG*: -5.01 \pm 0.51 fold and *murB*: -5.99 \pm 0.50 fold) (**Table S3**).

Crosslinked DNA runs differently from non-crosslinked DNA after denaturation because the crosslinks covalently bind the two strands, preventing separation (Matsumoto *et al.*, 1989). As confirmation of DNA crosslinking within persister cells found with qPCR, we isolated plasmid DNA (pDNA) from both exponential-phase cells (i.e., non-persisters) and rifampicin-induced persisters (Kwan *et al.*, 2013) before and after MMC treatment and performed agarose gel electrophoresis under denaturing conditions to allow uncrosslinked DNA to migrate as single strands. Under denaturing conditions, pDNA samples from cells without MMC treatment migrated as single-stranded DNA, while samples for both non-persisters and persisters treated with MMC showed a high percentage of crosslinking, based on migration as double-stranded DNA (**Fig. 2C**). DNA treated *in vitro* with MMC migrated in the expected manner as double-stranded DNA. Therefore, we have demonstrated by two independent means that MMC does in fact crosslink the DNA of persister cells.

MMC kills persister cells of pathogens. Clinical application of MMC is dependent on efficacy against pathogenic bacteria. Therefore, we tested the ability of MMC to kill *E. coli* O157:H7 (EHEC) a common pathogenic strain of Gram-negative *E. coli*. MMC was substantially more effective than ciprofloxacin against EHEC, eradicating both exponential (**Fig. 3A**) and mid-stationary phase cells (**Fig. 3B**) in rich medium beyond the limit of detection. Additionally, MMC eradicated late-stationary phase EHEC cells in M9-glucose beyond the limit of detection, while 1.2 ± 0.1 % of cells survived against ciprofloxacin (**Fig. 3C**). Biofilm cultures of EHEC were also eradicated (beyond the limit of detection) after 24 h of MMC treatment, in comparison to 3 ± 1 % of cells surviving ciprofloxacin (**Fig. 3D**).

MMC was also tested against Gram-positive *S. aureus* (methicillin sensitive) and Gram-negative *P. aeruginosa*, two other common species of human pathogens. Against planktonic cultures of *S. aureus* grown in rich medium, MMC was highly effective, eradicating both exponential (**Fig. 3E**) and mid-

stationary phase cultures (Fig. 3F), in comparison to ciprofloxacin (0.55 ± 0.04 % survival and 2.2 ± 0.1
% survival, respectively). MMC was also highly effective against biofilm cultures of *S. aureus* grown in minimal medium, eradicating cells beyond the limit of detection after 24 of treatment, compared with 18 ± 2 % survival against ciprofloxacin (Fig. 3G). MMC also killed cultures of *P. aeruginosa* PA14 grown planktonically in rich medium to exponential (Fig. 3H) and mid-stationary phase (0.0038 ± 0.0005 % survival) and in minimal medium to late-stationary phase (Fig. 3I), although the extent of killing activity
was similar to that of ciprofloxacin. Therefore, MMC is significantly more effective in eradicating EHEC and *S. aureus* and is similar to other potent antibiotics against *P. aeruginosa*, demonstrating the efficacy of MMC against several species of human pathogens.

MMC kills persister cells in a wound model. In clinical infections, bacteria are exposed to drastically different growth conditions from those generally used within laboratory cultures. The in vitro Lubbock 155 chronic wound pathogenic biofilm model was previously developed to closely represent growth conditions of polymicrobial infections (Sun et al., 2008). We used this in vitro wound model to test MMC activity against cultures of EHEC, S. aureus, and P. aeruginosa PAO1 as well as a co-culture of S. aureus and P. aeruginosa PAO1. Our strain of S. aureus is coagulase-positive (Hsueh et al., 1999), causing the medium to form a jelly-like mass consisting of insoluble fibrin (Zajdel et al., 1975). Cultures were grown statically so that coagulated plasma served as a scaffold for bacterial growth (DeLeon et al., 2014) in 160 cultures containing S. aureus, while biofilms formed at surface interfaces served as scaffolds for bacterial growth in cultures without S. aureus. We found that MMC was more effective than ciprofloxacin and ampicillin against all three species under wound-like conditions in mono- and co-cultures (Fig. 4A-D). These results show that MMC is a significantly more effective treatment than other antibiotics against pathogenic strains of several species (e.g., EHEC, S. aureus, and P. aeruginosa) grown using an in vitro 165 wound model. This substantiates the efficacy of MMC as a clinical treatment for clearing infections.

MMC is effective in an animal model. In order to test the efficacy of MMC treatment *in vivo*, we used an EHEC infection within the nematode *Caenorhabditis elegans*. *C. elegans* was fed on lawns of EHEC on nematode growth media (NGM) agar plates for 2 days in order to establish an infection. Nematodes were then exposed to MMC, ciprofloxacin, ampicillin, or no treatment, transferred to lawns of avirulent *E. coli* OP50, and monitored for viability. All three antibiotic treatments enhanced the survival of the EHEC-infected worms; however, survival with MMC was higher than with either ciprofloxacin or ampicillin based on four experimental replicates (10 worms per replicate) (Fig. 4E), likely because MMC eradicates persisters that can reestablish infection. Of note, we obtained similar results from four additional replicates performed with different antibiotic treatments and EHEC infection conditions. Therefore, MMC is consistently more effective than other antibiotics at clearing EHEC infection within an animal model.

CONCLUSION

Traditional antibiotics (e.g., fluoroquinolones, aminoglycosides, and β -lactams) are ineffective 180 against persister cells due to their mechanisms which rely on cellular activity. Here, we found that MMC is highly effective because of its unique mechanism of action, which is independent of the metabolic state, by demonstrating its activity against slow-growing, non-growing, and dormant (e.g., persister and VBNC) cells, as well as its activity on cells grown planktonically, in biofilms, in an in vitro wound model, and in an *in vivo* animal model. In comparison, several methods have been proposed for eradicating persister 185 cells, including increasing aminoglycoside uptake via glycolysis intermediates (Allison et al., 2011), altering membranes via Trp/Arg-containing antimicrobial peptides (Chen et al., 2011), activating ClpPmediated self-digestion via rifampicin and ADEP4 (Conlon et al., 2013), and converting persisters to nonpersisters via cis-2-decenoic acid (Marques et al., 2014). However, the potential application of these treatments against clinical infections is distant due to limited levels of *in vivo* testing. These treatments 190 are also likely limited to a small range of species that are susceptible to the compounds. In contrast, MMC has been an FDA-approved chemotherapeutic cancer drug for over forty years (Doll et al., 1985) with a well-characterized biochemical mechanism (Tomasz, 1995). Additionally, MMC passively diffuses into cells, and the DNA crosslinking activity of MMC is spontaneous, so MMC treatment should be effective against many bacterial species which cannot be fully cleared with traditional antibiotics such as 195 recalcitrant internal and external (wound) infections.

Throughout the majority of this study, MMC was effectively applied at a concentration of 10 μ g/mL, although the effective concentration should be much lower against strains with low MICs (e.g., EHEC and *S. aureus*; **Table 1**). For various cancer treatments, intravenously infused MMC dosages are often administered at concentrations between 0.5-2.0 μ g/mL (20-80 mg/m²) (Bradner, 2001) and topical dosages have been safely applied at concentrations up to 400 μ g/mL (Shields *et al.*, 2002). Therefore, the bactericidal concentrations of MMC are similar to the therapeutic concentrations that have been established for cancer treatments, which validates MMC as a readily applicable treatment for clinical infections.

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CONFLICT OF INTEREST STATEMENT

210 The authors declare no competing financial interests.

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Strain	ММС	Ciprofloxacin	Ampicillin	Gentamicin
<i>E. coli</i> K-12 BW25113	2 µg/mL	0.05 μg/mL	10 μg/mL	5 μg/mL
EHEC 86-24	1 μg/mL	0.05 µg/mL	$5 \ \mu g/mL$	
S. aureus ATCC29213	0.2 μg/mL	0.5 µg/mL	$2 \ \mu g/mL$	
P. aeruginosa PAO1	15 μg/mL	2 µg/mL	400 µg/mL	
P. aeruginosa PA14	2 µg/mL	0.1 µg/mL		

 Table 1.
 MICs for antibiotics used in this study.

FIGURE LEGENDS

Fig. 1. MMC eradicates metabolically dormant *E. coli* **K-12 cells in suspension and in biofilms.** Cell viability for exponential (turbidity of 2 at 600 nm) (**A**) and mid-stationary phase (turbidity of 4 at 600 nm) (**B**) cultures in buffered LB. (**C**) Time course of killing of late stationary-phase cells (16 h of growth) in buffered LB. (**D**) Cell viability of rifampicin-induced persister cultures in buffered LB (30 min pretreatment with 100 µg/mL rifampicin followed by resuspension in fresh media). Cell viability of exponential (turbidity of 0.4 at 600 nm) (**E**) and late-stationary phase (24 h of growth) (**F**) cultures in M9-glucose. Cell viability of biofilm cultures (24 h of growth) in M9-glucose (**G**), anaerobic late-stationary phase cultures (16 h of growth) in LB (**H**), anaerobic biofilm cultures (24 h of growth) in M9-glucose (**I**), and VBNC cultures (36 days of starvation in saline) (**J**). Cell viability is shown before (black) and after treatment (3 h for planktonic cultures, 24 h for biofilm cultures, and 16 h for VBNC cultures) with 5 µg/mL ciprofloxacin (green), 10 µg/mL gentamicin (purple), and 10 µg/mL MMC (red). * represents eradication beyond the limit of detection. Means ± s.d. are shown throughout ($n \ge 2$). MMC is mitomycin C and CIP is ciprofloxacin.

Fig. 2. MMC crosslinks DNA in *E. coli* K-12 persister cells. (A) Cell viability of *E. coli* K-12 wildtype, $\Delta uvrA$, $\Delta uvrB$, and $\Delta uvrC$ mid-stationary phase cultures (turbidity of 3 at 600 nm) in buffered LB treated for 30 min with MMC. (B) Cell viability of *E. coli* K-12/pCA24N, $\Delta uvrA/pCA24N$, and $\Delta uvrA/pCA24N$ -uvrA exponential phase cultures (turbidity of 2 at 600 nm) in buffered LB treated for 1 h with MMC. Cell viability is shown before (black) and after treatment with 10 µg/mL MMC (red). * represents eradication beyond the limit of detection. Means ± s.d. are shown for **A** and **B** ($n \ge 2$). (C) Denaturing gel electrophoresis for pDNA (4,518 nt) from *E. coli* K-12/pCA24N non-persisters (lanes 2 and 3) and rifampicin-induced persisters (lanes 4 and 5) before (lanes 2 and 4) and after (lanes 3 and 5) MMC treatment. Lane 1 is a positive control with *in vitro* crosslinked pDNA. "M" indicates the DNA ladder, the red arrow indicates migration as double-stranded DNA, and the blue arrow indicates migration as singlestranded DNA.

- **Fig. 3. MMC eradicates pathogens in suspension and in biofilms.** Cell viability of EHEC exponential (turbidity of 2 at 600 nm) (**A**) and mid-stationary phase (turbidity of 4 at 600 nm) (**B**) cultures in buffered LB, late-stationary phase cultures (24 h of growth) in M9-glucose (**C**), and biofilm cultures (24 h of growth) in M9-glucose (**D**). Cell viability of *S. aureus* exponential (turbidity of 0.8 at 600 nm) (**E**) and mid-stationary phase (turbidity of 3 at 600 nm) cultures in tryptic soy broth (TSB) (**F**), and biofilm cultures (24 h of growth) in modified M9-glucose (**G**). Cell viability of *P. aeruginosa* PA14 exponential phase cultures (turbidity of 2 at 600 nm) in buffered LB (**H**) and late-stationary phase cultures (24 h of growth) in M9-glucose (**I**). Cell viability is shown before (black) and after treatment (3 h for planktonic cultures and 24 h for biofilm cultures) with 5 µg/mL ciprofloxacin (green) and 10 µg/mL MMC (red). * represents eradication beyond the limit of detection. Means ± s.d. are shown throughout ($n \ge 2$).
- Fig. 4. MMC eradicates pathogens in clinically relevant wound and animal models. Cell viability of EHEC (A), *S. aureus* (B), and *P. aeruginosa* PAO1 (C) mono-cultures and *S. aureus* and *P. aeruginosa* PAO1 co-cultures (D) in an *in vitro* wound model (24 h of growth). Cell viability is shown before (black) and after 5 h treatment with ciprofloxacin (blue; 5 μ g/mL for EHEC and *S. aureus* or 10 μ g/mL for PAO1 mono- and co-cultures), ampicillin (green; 100 μ g/mL for EHEC and *S. aureus* or 2 mg/mL for PAO1 mono- and co-cultures), and MMC (red; 10 μ g/mL for EHEC and *S. aureus* or 15 μ g/mL for *P. aeruginosa* PAO1 mono- and co-cultures). * represents eradication beyond the limit of detection. (E) Survival of *C. elegans* after infection with EHEC (days -2 to 0) followed by 6 h exposure to 5 μ g/mL ciprofloxacin (green), 100 μ g/mL ampicillin (blue), 10 μ g/mL MMC (red), or no treatment (white). As a negative control, *C. elegans* was grown on OP50 without antibiotic treatment (black). Means ± s.d. are shown throughout ($n \ge 2$).



