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To cite this version:

Jean Michel Brunel, Gang Wang, Jean-Michel Brunel, Jean-Michel Bolla, Françoise van Bambeke. The Polyaminoisoprenyl Potentiator NV716 Revives Old Disused Antibiotics against Intracellular Forms of Infection by Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2021, 65 (3), 10.1128/AAC.02028-20 hal-03367286

HAL Id: hal-03367286 <https://hal.science/hal-03367286v1>

Submitted on 11 Oct 2021

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Revision for Antimicrobial Agents and Chemotherapy

The polyaminoisoprenyl R2.1potentiator NV716 revives old disused antibiotics against intracellular forms of infection by *Pseudomonas aeruginosa*

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Paper metrics: Abstract: 250 words (max 250) Main text length: 4556 words (no limit) References: 48 Figures/Tables: 5/1 Supplementary material: 2 Tables and 3 Figures * Corresponding author: Avenue Mounier 73, B1.73.05, 1200 Brussels, Belgium Email: francoise.vanbambeke@uclouvain.be Phone: +32-2-764.73.78

Abstract:

Active efflux confers intrinsic resistance to multiple antibiotics in *Pseudomonas aeruginosa*, including old disused molecules. Beside resistance, intracellular survival is another reason for failure to eradicate bacteria with antibiotics. We evaluated the capacity of polyaminoisoprenyl **R2.1**.potentiators (designed as efflux pump inhibitors [EPIs]) NV716 and NV731 compared to PAβN to restore the activity of disused antibiotics (doxycycline, chloramphenicol [substrates for efflux], rifampin [not substrate]) in comparison with ciprofloxacin against intracellular *P. aeruginosa* (strains with variable efflux levels) in THP-1 monocytes exposed during 24h to antibiotics alone (0.003-100x MIC) or combined with EPIs. Pharmacodynamic parameters (apparent static concentrations $[C_s]$; maximal relative efficacy $[E_{max}]$) were calculated using the Hill equation of concentration-response curves. PAβN and NV731 moderately reduced (0-4 doubling dilutions) antibiotic MICs but did not affect their intracellular activity. NV716 markedly reduced (1-16 doubling dilutions) the MIC of all antibiotics (substrates or not for efflux; strains expressing efflux or not); it improved their relative potency and maximal efficacy (lower C_s ; more negative E_{max}) intracellularly. In parallel, NV716 reduced the persister fraction in stationary cultures when combined with ciprofloxacin. In contrast to PAβN and NV731 that act as EPIs against extracellular bacteria only, NV716 can resensitize *P. aeruginosa* to antibiotics whether substrates or not for efflux, both extracellularly and intracellularly. This suggests a complex mode of action that goes beyond a simple inhibition of efflux and reduces bacterial persistence. NV716 may appear as a useful adjuvant, including to disused antibiotics with low antipseudomonal activity, to improve their activity, including against intracellular *P. aeruginosa.*

INTRODUCTION

 Pseudomonas aeruginosa is one of the so-called ESKAPE (*Enterococcus faecium*, *Staphylococcus aureus*, *Klebsiella pneumoniae*, *Acinetobacter baumannii*, *Pseudomonas aeruginosa*, and *Enterobacter species*) pathogens (1) that harbor multiple mechanisms of drug resistance. It is considered by the World Health Organization (WHO) as a critical priority species for the search of innovative therapies (2). *P. aeruginosa* has acquired a series of resistance mechanisms to all classes of commonly used 8 antibiotics (β -lactams, fluoroquinolones, aminoglycosides, and, more recently, polymyxins). Moreover, it is also innately resistant to many antibiotics, mainly due to poor outer membrane permeability and/or active efflux (3). Four main efflux systems, named MexAB-OprM, MexCD-OprJ, MexEF-OprN, and MexXY-OprM, have been associated with increased MICs for a broad range of antimicrobial agents (4), bringing the active concentrations above the levels that can be achieved in the serum of patients. This explains why drugs like tetracyclines or chloramphenicol cannot be used against *P. aeruginosa* or other Gram-negative bacteria, although they are capable of binding to their respective target in these bacteria when tested in acellular systems (5, 6).

 Another, often neglected, reason for antibiotic failure against *P. aeruginosa* is the capacity of this bacterium to adopt specific lifestyles, such as intracellular survival. Although considered as an extracellular pathogen, *P. aeruginosa* has been shown by many researchers as capable of invading and surviving inside epithelial (7, 8) and phagocytic cells (9-11) *in vitro*, as well as in lung epithelial cells and macrophages *in vivo* (12, 13). In these intracellular niches, bacteria are protected from the host humoral immune defenses but also from antibiotics, which need to have access, accumulate, and express their activity in the infected subcellular compartment (14, 15). Moreover, they can adopt dormant phenotypes that do not respond anymore to antibiotics, like persisters. These are subpopulations of otherwise antibiotic-susceptible bacteria that switch to a transient non-dividing phenotype surviving to high concentrations of bactericidal antibiotics (16). We have recently provided evidence for another species, *Staphylococcus aureus,* that bacteria surviving to antibiotics intracellularly are persisters, explaining why antibiotics fail to eradicate them (17). Preliminary data suggest that the same may hold true for *P. aeruginosa* (18).

 As *P. aeruginosa* resistance is worryingly increasing, few new antibiotics have been approved over the last years, and are essentially new derivatives in existing classes with improved activity or reduced susceptibility to resistance mechanisms (19). In this context, reviving old and disused antibiotics may appear as an attractive strategy (20), which should be accompanied by an in-depth reassessment of their efficacy according to the modern standards imposed by pharmacokinetic/pharmacodynamic (PK/PD) concepts, including in models of persistent infections like intracellular survival.

 One of the approaches proposed to increase the potency of drugs that are substrates for efflux is to combine them with efflux pump inhibitors (EPIs)(19). PAβN (MC 207,110; Phenylalanine-arginine β-naphthylamide (21)) and CCCP (Carbonyl cyanide m-chlorophenyl hydrazone (22)) have been widely used *in vitro* to increase the accumulation of antibiotics within Gram-negative bacteria, but their 42 toxicity R1.2.and inadequate pharmacokinetic properties preclude clinical applications (23, 24). Efforts have been made to synthesize new efflux pumps inhibitors with improved safety profile (25). Among them, polyaminoisoprenyl compounds **NV716** and **NV731** (see chemical structures in Figure S1) demonstrated their capacity to restore chloramphenicol activity against *Enterobacteria* spp (26) and one of them (**NV716**), that of chloramphenicol and doxycycline against *Pseudomonas aeruginosa* (27), or of florfenicol against *Bordetella bronchiseptica* (28).

 The aim of the present study was to evaluate the effects of **NV716** in comparison with that of **NV731** (less potent against *P. aeruginosa* (27)) and PAβN (reference EPI) in combination with old and disused antibiotics against the intracellular forms of *P. aeruginosa*, using strains expressing or not efflux pumps to infect THP-1 monocytes. We exploited a previously established *in vitro* pharmacodynamic model (11) that allows comparing key pharmacological descriptors of antibiotic intracellular activity, namely their relative potency and maximal efficacy (29). As antibiotics, we 55 selected doxycycline, chloramphenicol (both substrates for efflux (see Table S1) and for which preliminary data do exist regarding the beneficial effect of **NV716** (27)), rifampin (not substrate for efflux), and ciprofloxacin, selected as the most active drug against intracellular *P. aeruginosa* in this

- pre-established *in vitro* model (11). The results were examined in the light of the effect of these EPIs on antibiotic MIC and on the selection of persisters in broth culture.
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 We showed that **NV716**, contrarily to **NV731** and PAβN, was capable of increasing the relative potency and maximal efficacy of these antibiotics against intracellular *P. aeruginosa*, whether substrates of not for efflux and in strains expressing or not efflux pumps. These effects were related to drastic reductions in MIC and in persister fractions. Altogether, these data suggest that **NV716** does not simply act by inhibiting efflux but could be a useful adjuvant, including to disused antibiotics with low antipseudomonal activity, to improve their activity also against intracellular *P. aeruginosa.*

RESULTS

Antimicrobial susceptibility

 Table 1 shows the MICs of antibiotics against PAO1, PAO509 (deleted in the genes encoding 5 efflux pumps), 205-2 and BV1 (two clinical isolates expressing respectively a non-functional and a functional MexAB-OprM efflux pump), in the absence of or in the presence of 20 mg/L (38 μM) PAβN, or 4 mg/L (10 μM) **NV731** or **NV716**. These concentrations were selected because commonly used in previous works (21, 27). In control conditions (no inhibitor), ciprofloxacin, doxycycline, and chloramphenicol showed much lower MICs against PAO509 than against PAO1, in accordance with the fact they are 76 substrates for efflux (see Table S1). Conversely, the MIC of rifampin, which is not a substrate, remained unchanged. BV1 was less susceptible than PAO1 to ciprofloxacin and doxycycline and more susceptible to chloramphenicol, while 205-2 was less susceptible to ciprofloxacin only. PAβN and **NV731** reduced antibiotic MICs in few cases, while **NV716** systematically decreased them, except for ciprofloxacin in PAO509, the MIC of which was already very low. Of note, the effect of **NV716** was as (or even more) important when testing strains that do not express efflux pumps or those that do, as well as when using rifampin (not substrate for efflux) or other antibiotics (all substrates for efflux).

Cytotoxicity of inhibitors and antibiotics

85 In a first set of experiments, we evaluated the cytotoxicity of the EPIs and antibiotics in the conditions of their use for intracellular infection by trypan blue exclusion test after 24 h incubation. Infection of THP-1 cells by PAO1 caused 85% cell mortality in the absence of antibiotic. The inhibitors alone caused concentration-dependent cell mortality, which was minimal at the concentration used for intracellular 89 infection studies (Figure 1, left panel). Doxycycline, chloramphenicol, and rifampin showed toxic effects 90 at high concentrations, with IC_{50} (i.e. concentration causing 50% cell mortality) reaching approx. 30 91 times their respective MIC in infected cells (Figure 1, right panel). Ciprofloxacin was much less toxic at 92 equivalent multiples of its MIC (which corresponds to much lower mass concentrations). IC₅₀ values for toxicity of inhibitors, antibiotics or their combinations in non-infected and infected cells are detailed in 94 supplementary Figure S2 and Table S2 but do not differ from what is described in Figure 1.

96 **Activity of antibiotics alone or combined with EPIs against intracellular** *P. aeruginosa* **in an** *in vitro* 97 **pharmacodynamic model**

98 Figure 2 shows the concentration-response curves against intracellular PAO1 in the absence of or in the presence of **NV716**. The activity of antibiotics alone developed following a sigmoidal 100 concentration-response curve, as previously described (11). At low, subMIC concentrations, a \approx 3 log₁₀ increase in CFUs was noticed over the 24 h incubation period. A static effect was observed at extracellular concentrations close to the MIC of each drug. At the highest concentration tested, the 103 reduction in bacterial counts ranged from 2 log₁₀ CFU decrease for chloramphenicol, doxycycline, and 104 rifampin, to 2.6 log₁₀ CFU decrease for ciprofloxacin. In the presence of **NV716** at 10 μM, all curves (except that of ciprofloxacin) were shifted to the left, meaning that the corresponding antibiotic was more potent, a static effect being reached at lower extracellular concentrations. Furthermore, the maximal reduction in intracellular counts was also increased, indicating that the antibiotics were more effective.

109 The same type of experiment was performed with the other EPIs and the other strains. Figure 3 110 illustrates the pharmacodynamic parameters calculated based on the Hill equation of these 111 concentration-response curves (see methods for a definition of these parameters), namely the C_s (a 112 measure of the relative potency of the drug or the drug combination) and the E_{max} (a measure the 113 relative maximal efficacy of the drug alone or in combination). C_s of antibiotics alone were close or 114 slightly higher than the MICs and systematically reduced of 1 doubling dilution or more by **NV716**, 115 except for ciprofloxacin against PAO1 and PA0509, which were already highly susceptible to this drug. 116 **NV731** and PAβN did not show any significant effect on C_s values. Similarly, NV716 was the only EPI 117 capable of increasing the efficacy of the 4 drugs against all strains, with E_{max} values being at least 0.5 118 log_{10} CFU more negative than those measured for the antibiotics alone.

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120 **Concentration-response curves of NV716 combined with fixed concentrations of antibiotics against** 121 **intracellular** *P. aeruginosa*

122 **NV716** being the only adjuvant capable of improving both the relative potency and the relative 123 maximal efficacy of antibiotics, we examined the concentration-dependency of its effect when 124 combined with antibiotics at fixed concentrations corresponding to their respective C_{min} and C_{max} in

125 human serum (Table S1 for these values). The data are illustrated in Figure 4. Considering first the 126 activity of **NV716** alone, a ~3 log₁₀ increase in CFU was observed after 24 h incubation with low **NV716** 127 concentrations (≤ 10 µM) for all strains except PAO509, for which a slight reduction in CFU was already 128 observed at 10 µM. A static effect was obtained for **NV716** concentrations close to its MIC in RPMI-129 1640 (25 µM) for all strains. At 50 μM, **NV716** caused a reduction in bacterial counts that ranged from 130 \degree ~1 log₁₀ CFU decrease for PAO1, 205-2 and BV-1, to ~2 log₁₀ CFU decrease for PAO509. These data are 131 coherent with those obtained in broth (Figure S3), showing that **NV716** did not prevent PAO1 growth 132 for concentrations $\leq 10 \mu$ M but caused a rapid decrease in bacterial counts at 50 μ M. In the presence 133 of antibiotics at their C_{min} , we noticed that the intracellular concentration-response curves were 134 globally parallel to those obtained with **NV716** alone, reflecting thus essentially the activity of the EPI. 135 Three exceptions need however to be highlighted. First, the activity of chloramphenicol was increased 136 for **NV716** concentrations > 3 µM against all strains. Second, an improvement in antibiotic activity was 137 noticed against PAO509, as soon as **NV716** concentration was > 3 µM for all drugs that are substrates 138 for efflux (ciprofloxacin, chloramphenicol and doxycycline). Third, at the highest concentration of 139 **NV716** tested (50 μ M), the reduction in CFU counts was 0.5-1.8 log_{10} more important than that 140 observed with **NV716** alone for all drugs, denoting some sort of synergy. Combining **NV716** with 141 antibiotics at their C_{max} caused a drastic improvement of activity, with CFU counts being reduced from 142 0.5-2.2 log_{10} at **NV716** concentrations \leq 10 μ M to 1.4-5.6 log_{10} at higher **NV716** concentrations in 143 comparison with CFU counts measured for the antibiotic alone.

144

145 **Persister assay**

 The fact that **NV716** increases the intracellular efficacy of antibiotics (improved killing capacity) suggests it may decrease the proportion of antibiotic persisters in the population, recently identified as a cause of failure to eradicate intracellular *S. aureus* with antibiotics (30). We therefore examined the persister fraction surviving to ciprofloxacin (selected as highly bactericidal antibiotic) alone or combined with **NV716** in stationary phases cultures. As shown in the left panel of Figure 5A, ciprofloxacin at 50x MIC caused a rapid drop in CFU counts of PAO1 over the 2 first hours of incubation, after which no further change was observed, leaving a persisting fraction of 0.005% of the initial inoculum. The same type of experiment was then performed with PAO1 and PAO509 exposed to

 NV716 at increasing concentrations and used alone or combined with 50x MIC of ciprofloxacin. The 155 right panels of Figure 5A show the proportion of persisters after 5 h of incubation, expressed in percentage of the CFU counts measured in the absence of drugs. **NV716** alone had no effect on 157 persisters for concentrations ≤ 100 µM but caused a marked reduction in bacterial counts at higher 158 concentrations (3-5 log_{10} CFU for concentrations > 300 μ M for PAO1 and > 100 μ M for PAO509). Strikingly, **NV716** at lower concentrations (10-100 µM) was already capable to reduce up to 11- (PAO1) to 28- (PAO509) fold the percentage of persisters selected by ciprofloxacin.

 R2.3To investigate whether **NV716** needs to be administered simultaneously with ciprofloxacin to obtain a maximal activity, we repeated killing experiments with stationary phase cultures exposed to 50 x MIC of ciprofloxacin alone or combined with **NV716** added at different time points (Figure 5B) (31). CFUs were significantly reduced after the addition of **NV716** at any time point, for concentrations ≥ 10 µM and in both PAO1 and PAO509, indicating that **NV716** does not need to be added at the same time as the drug to be able to increase its activity. In a next step, we aimed at examining whether **NV716** 167 action was related to a direct capacity to kill persisters or rather to resensitize them to the killing effect of ciprofloxacin. To this end, we isolated the persisters surviving after 5 h of incubation with ciprofloxacin at 50 x MIC and reexposed them to ciprofloxacin at 50 x MIC, **NV716** at different concentrations**,** or a combination thereof (Figure 5C) (31). As expected, incubation of the isolated persisters with ciprofloxacin alone caused only a marginal decrease in CFUs for both strains, confirming the effective isolation of persister cells. When used alone, **NV716** caused a significant decrease in CFUs 173 only at the highest concentration tested (250 μ M). In contrast, a significant reduction in CFUs was 174 already observed when it was combined at 10 µM with ciprofloxacin for both strains.

DISCUSSION

 In this study we show that, in contrast with PAβN and **NV731** that act as efflux inhibitors against planktonic bacteria only, **NV716** is capable to re-sensitize *P. aeruginosa* to antibiotics whether substrates (doxycycline, chloramphenicol, ciprofloxacin) or not (rifampin) for efflux, not only in broth but also intracellularly.

 Considering first susceptibility data in broth, we observed a remarkable effect of **NV716** on the antibiotic intrinsic activity. While PAβN and **NV731** moderately reduced the MICs of antibiotics (essentially those that are substrates for efflux), **NV716** caused a marked decrease in the MICs of all 186 drugs, comprising rifampin (not substrate for efflux; see Table S1) against all strains, including PAO509 that does not express efflux pumps or BV1 that expresses a non-functional MexAB-OprM efflux pump. This suggests that its mode of action is not related to the inhibition of efflux only, but probably also relies on its capacity to permeabilize the outer membrane of the *P. aeruginosa* (27). Of note, the synergy with antibiotics is obtained at sub-inhibitory concentrations of **NV716**, R1.3.indicating that, if a direct effect on bacteria takes place, it is not sufficient to cause bacterial death. PAβN has also been 192 shown to permeabilize the outer membrane of Gram-negative bacteria by displacing Mg²⁺ and Ca²⁺ (32), but at concentrations higher than the one we used here, established as adequate to inhibit efflux in princeps studies with this molecule (21). The activity of **NV716** and of the two other EPIs is observed at concentrations that do not cause cytotoxicity for eukaryotic cells, suggesting no or minimal interaction with eukaryotic membranes.

 This contrasts with the cytotoxicity noticed for the three old and disused antibiotics, which caused significant loss of viability of THP-1 monocytes at the highest concentrations tested here, which are largely above those measured in the plasma of treated patients, owing to the high MIC of these drugs against *P. aeruginosa*. We had thus to limit the range of antibiotic concentrations tested in our concentration-response curves against intracellular bacteria in order to avoid a too massive loss of cells.

204 Considering then intracellular activity data, we noticed that all antibiotics displayed a 205 bacteriostatic activity at concentrations close or slightly higher than their MIC (C_s being considered as a 206 measure of the intracellular MIC), despite the fact they all accumulate to some extent in eukaryotic 207 cells (33) (Table S1). This is in line with our previous observations with other antibiotics in the same 208 model (11) or in similar models of intracellular infections by other bacterial species (34, 35), and was 209 interpreted as denoting a poor intracellular bioavailability. While PAβN and **NV731** did not increase this 210 relative potency (minimal modification in C_s), **NV716** caused a shift of the concentrations-response 211 curves to lower concentrations, with a commensurate reduction in the C_{s} , except for ciprofloxacin 212 against PAO1 and PAO509, the MIC of which was already rather low. Thus, **NV716** can decrease the 213 MIC of antibiotics not only extracellularly, but also intracellularly. We did not measure the cellular 214 concentration of **NV716** in the cells, but these activity data suggest that it has access to intracellular 215 bacteria in sufficient concentrations to exert its synergistic effect. The absence of gain in relative 216 potency observed with PAβN and **NV731** can be consecutive to their lower intrinsic effects on MICs (as 217 observed in broth) or possibly also to a lower penetration inside the cells, which has not been 218 investigated.

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 Turning then our attention towards the intracellular maximal efficacy, we observed, for 221 antibiotics used alone, reductions of 2-2.5 log_{10} CFU with ciprofloxacin, in accordance with our previous 222 results (11), and of 1-2 log_{10} CFU for the other antibiotics, the lowest effect being observed for chloramphenicol and doxycycline against 205-2, indicating individual differences in responsiveness among strains. In a broader context, the extent of intracellular killing may also differ depending on the 225 bacterial species. An intracellular E_{max} of -1.8 log_{10} CFU was noticed for rifampin against a small colony variant of *S. aureus* (36), but the activity of chloramphenicol and doxycycline has not been investigated in this model. Chloramphenicol demonstrated bacteriostatic activity against intracellular *Salmonella* 228 Typhimurium at 10x MIC (37) while doxycycline reduced of 1.5 log₁₀ the intracellular CFU of *Francisella tularensis* after 24 h of incubation at a concentration close to its MIC (38). For fluoroquinolones, a 230 maximal reduction of 2 to 4 log_{10} CFU was observed in concentration-response experiments similar to those performed here against *S. aureus, Listeria monocytogenes, Legionella pneumophila, Burkholderia thailandensis, Yersinia pseudotuberculosis,* or *Francisella philomiragia* (34, 35, 39).

 While PAβN and **NV731** did not affect the efficacy of any antibiotic against none of the strains, **NV716** systematically improved it (more negative E_{max}), suggesting an effect on bacterial responsiveness to antibiotics. We can most likely preclude interference related to the release of 236 bacteria out of the cells before lysis or to the **R1.6** remaining amount of antibiotics accumulated in cells if considering that (a) **NV716** did not increase the cytotoxicity of antibiotics, and (b) charcoal was added to culture plates to adsorb residual antibiotic. Thus, the gain in intracellular efficacy observed in the presence of **NV716** is rather suggestive of improved bacterial responsiveness to the drugs. We previously showed an increase in the intracellular efficacy of fluoroquinolones against an insertion mutant in a putative deacetylase in PAO1, which showed a lower persister character in planktonic cultures (18). Here, we found that **NV716** was capable of decreasing the proportion of persisters 243 selected by ciprofloxacin in a stationary phase culture at concentrations for which it did not cause any 244 bacterial killing by itself R2.3and also of reducing at subMIC concentrations the number of persisters 245 when combined with ciprofloxacin but not when used alone, suggesting a mode of action unrelated to 246 a direct bactericidal effect but rather to a resensitization of persisters to the antibiotic (31). Of interest, 247 other molecules capable of disrupting membrane integrity also reduced the residual fraction of persisters upon exposure to antibiotics for different bacterial species (40-42). One of them (SPI009; 1- ((2,4-Dichlorophenethyl)Amino)-3-Phenoxypropan-2-ol)) was also shown to increase the efficacy of ciprofloxacin against intracellular *P. aeruginosa* (43).

 This work suffers from some limitations. Firstly, we used a single phagocytic cell line (THP-1 monocytes), which does not necessarily represent the repertoire of cells infected by *P. aeruginosa in vivo*. The reason for selecting this permissive cell line was that it allows assessing the effect of antibiotics without interference of cell defense mechanisms. Secondly, we did not measure the cellular accumulation of antibiotics and EPIs in our experimental conditions, but setting-up the necessary methodologies would represent a work by itself. Thirdly, we did not study in details the occurrence of intracellular persisters, the aim of the study being rather oriented towards an in-depth pharmacodynamic evaluation of the possible interest of EPIs to restore the activity of old and disused antibiotics against intracellular *P. aeruginosa.*

262 If we thus examine our data in a more clinically-oriented perspective, we see that **NV716** is 263 capable of improving the activity of all antibiotics in the range of concentrations achieved in the serum 264 of patients as soon as its extracellular concentration exceeds 3 µM. Ten times higher concentrations 265 allow to reach a 2-3 log_{10} CFU decrease when combined with antibiotics at their human C_{max}, pleading 266 for examining the possible interest of this compound *in vivo*.

MATERIALS AND METHODS

Bacterial strains and culture media

 Table 1 shows the reference and clinical strains used in the study. R2.m2BV-1 has been isolated in 2006 272 at the Queen's University Belfast, Belfast, UK, from a patient with cystic fibrosis. It shows a truncation in MexB (672 amino acids instead of 1046) that drastically reduced the activity of efflux transporter and its impact of the MIC of the preferential MexAB-OprM antibiotic substrate temocillin (44). 205 has been isolated in 2012 at the University Hospital Münster, Münster, Germany, from a patient with cystic fibrosis (45), and shows a fully active MexAB-OprM efflux pump, based on the elevated MIC of temocillin.

278 Bacteria were R2.m1incubated overnight at 37°C in Tryptic Soy Agar (TSA; VWR; Radnor, PA). A single 279 colony was then **R2.m1**added in 10mL cation-adjusted Mueller-Hinton Broth (CA-MHB; BD Life Sciences, Franklin Lakes, NJ) and incubated at 37°C overnight under gentle agitation (130 rpm). TSA supplemented with 2g/L charcoal (Sigma-Aldrich, St Louis, MO) was used for CFU counting (colony- forming unit). Cell culture media (RPMI-1640) and human and fetal bovine sera were from Gibco/ThermoFisher Scientific (Waltham, MA).

Antibiotics and Inhibitors

 Chloramphenicol (potency, 98%), doxycycline (potency, 98%), and rifampin (potency, 98%) were obtained as microbiological standards from Sigma-Aldrich (St Louis, MO), ciprofloxacin HCl (potency, 89%) from Bayer (Leverkusen, Germany) and gentamicin sulfate (potency, 60.7%), from PnReac AppliChem (Darmstadt, Germany). **Table S1** shows relevant properties of the antibiotics used in this study*.* The reference efflux pump inhibitor Phe-Arg β-Naphthylamide (PAβN; potency, 98%) was purchased from Sigma-Aldrich. The hydrosoluble hydrochloride polyamino-isoprenic salt derivatives **NV716** and **NV731** were synthesized at Aix-Marseille University (26). Stock solutions of these inhibitors were prepared in dimethyl sulfoxide (DMSO) at concentrations of 96 mM [50 mg/mL] for PAβN or in water at 10 mM [4 mg/L] for **NV716** and **NV731**, respectively, and stored at -20°C until use.

Susceptibility Testing

 MICs of antibiotics against *P. aeruginosa* were measured by serial 2-fold microdilution in MHB-CA according to CLSI guidelines in control conditions or in the presence of 38 µM [20 mg/L](21) PAβN or of 10 µM [4 mg/L] **NV731** or **NV716** (27).

Cytotoxicity assessment (Trypan blue exclusion assay)

302 THP-1 cells (7.5 x 10⁵ cells per mL) were incubated for 24 hours in 96-wells plates with the antibiotics or inhibitors under study over a wide range of concentrations. At the end of the incubation, viability was assessed using trypan blue exclusion test (vital colorant excluded from viable cells) using a previously described method (29), with specific adaptations to the purpose of our study. Briefly, 50 μL of trypan blue reagent was added to 50 μL of cell suspension. After 10 min of incubation at 37°C, non colored (viable) cells were counted using a Fuchs-Rosenthal counting chamber (Tiefe 0.2mm). Cells were counted in the same surface for all conditions (1 big square made of 16 single squares (0.0625 mm² for each single square). The percentage of cytotoxicity was evaluated based on the reduction in the number of living cells according to the following equation: 311 cytotoxicity $(\%) = \frac{Nb \text{ uncolored cells in controls} - Nb \text{ uncolored treated cells}}{Nb \text{ uncolored cells in controls}} x100.$

Intracellular Infection

 All experiments were performed using human THP-1 monocytes cultured in RPMI-1640 medium supplemented with 10% fetal bovine serum (FBS) based on a previously established protocol (11). In brief, bacteria were opsonized by incubation during 1 h in RPMI-1640 medium supplemented with 10% human serum at 37°C with gentle agitation (130 rpm). Phagocytosis of opsonized bacteria was then allowed for 2 h using a bacterium:cell ratio of 10:1, after which non-phagocytosed bacteria were eliminated by incubation for 1 h with gentamicin at 50x MIC. After 3 washings with PBS, infected cells were resuspended in the original volume of RPMI-1640 with 10% FBS. The post-phagocytosis inoculum 321 was consistently 5 to 7 x 10^5 CFU/mg of cell protein (defined as time zero). Infected cells were then incubated for 24 h with antibiotics over a broad range of concentrations (0.003-100x their MIC) in 12- well plates. After 24 h of incubation, infected cells were pelleted by centrifugation, washed once with PBS at room temperature to eliminate extracellular bacteria and collected in 1 mL distilled water, to

 achieve complete cell lysis. Lysates were used for determining CFU counts by spreading on agar supplemented with 2g/L charcoal (to avoid carry-over effect, especially in samples that had been 327 incubated with high antibiotic concentrations $[R1.5]$ no difference in colony counts was observed between plates supplemented or not with charcoal for samples exposed to low antibiotic concentrations, ruling out any interfering effect of charcoal in the assay]) and protein content using a commercially available kit (Bio-Rad DC Protein Assay, Bio-Rad Laboratories, Hercules, CA). The activity was expressed as the change in CFU (normalized by mg of cell protein) from the initial inoculum after 24 hours of incubation. The data were used to fit a sigmoidal function and calculate pharmacodynamic 333 parameters based on the corresponding Hill-Langmuir equation (apparent static concentrations $[C_s]$, i.e. extracellular concentration resulting in no apparent intracellular growth, and maximal relative efficacy $[Em_{max}]$, i.e. maximal decrease in bacterial counts compared to the post-phagocytosis inoculum as extrapolated for an infinitely large antibiotic concentration).

Persister assay and isolation

339 A single colony from overnight culture of PAO1 and PAO509 on TSA was R2.m1.added to in 200mL MHB-CA in a 1 L flask and incubated for 24 h under agitation (130 rpm) at 37°C. 10 mL of the bacterial suspension were then incubated with 50x MIC ciprofloxacin for 5 hours at 37°C under agitation (130 rpm), in the presence of or in the absence of **NV716.** Aliquots were serially diluted in PBS and spread on TSA supplemented with 2 g/L charcoal for CFU counting. The persister fraction was defined as the 344 number of CFU growing at the end of this experiment as compared to the initial inoculum (46). R2.3For experiments using isolated persisters, stationary-phase cultures were incubated with ciprofloxacin (50x MIC) during 5 hours, i.e. condition for which a plateau of killing was reached. The remaining bacteria were washed twice with sterile Phosphate Buffered Saline (PBS), centrifuged (5,200 *g*, 15 min, 4°C) and used for killing assays (31).

Curve fittings and statistical analyses.

 Curve fittings were performed with GraphPad Prism (version 8.4.3) software for Windows (GraphPad Prism Software, San Diego, CA). Pharmacodynamic parameters were calculated based on Hill equations

- of concentration-response curves and statistical analyses were executed with GraphPad InStat 3
- version 3.10 (GraphPad Prism Software) or Graphpad Prism version 8.4.3.

Acknowledgements

We are thankful to V. Mohymont, K. Santos-Saial, and V. Yfantis for proficient technical assistance.

W.W.G. received a PhD grant from the China Scholarship Council (CSC). F.V.B. is Research Director from

the Belgian *Fonds de la Recherche Scientifique* (FRS-FNRS).

Funding

 This work was supported by the Belgian FRS-FNRS (grant T.0189.16). We thank Campus France and *Wallonie Bruxelles International* (WBI) for financing a PHC Tournesol program (France-Belgium 2018- 2019).

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Table 1. MICs of antibiotics and adjuvants (EPIs) alone or combined against reference strains and clinical isolates in broth.

^a Values in bold denote a decrease of at least 2 doubling dilutions vs. antibiotics alone

^b Abbreviations: AB: antibiotic; CIP, ciprofloxacin; DOX, doxycycline; CHL, chloramphenicol; RIF, rifampin. MICs were determined in at least two independent experiments.

Figure 1. Cytotoxicity of EPIs and antibiotics as determined by trypan blue exclusion test. Left: cytotoxicity of efflux inhibitors in non-infected THP-1 cells incubated with increasing concentrations of NV716, NV731, or PAβN during 24 h. Right: cytotoxicity of antibiotics in THP-1 cells infected by PAO1 and incubated with increasing concentrations of ciprofloxacin (CIP), doxycycline (DOX), rifampin (RIF), or chloramphenicol (CHL) during 24 h. R1.6. Antibiotic concentrations are expressed in multiples of their MIC against PAO1 (Log₁₀) scale)The tick horizontal dotted lines show the percentage of cell mortality measured in untreated infected cells. The thin dotted line shows a 50% cell mortality threshold, allowing to estimate IC_{50} values for each compound (see Figure S2 and Table S2 for these values). The vertical arrows in the left panel show the inhibitor concentrations used in intracellular experiments $R1.2.(10 \mu M)$ for NV716 and 38 μ M for PA β N). All data are means \pm SEM (triplicates from 3 independent experiments).

Figure 2. Concentration-response curves of antibiotics alone or combined with NV716 against intracellular PAO1 in a model of THP-1 monocytes. The graphs show the changes in CFU counts per mg of cell protein from the initial, post-phagocytosis inoculum after 24 h of incubation with increasing extracellular concentrations of antibiotics alone (ciprofloxacin [CIP], doxycycline [DOX], chloramphenicol [CHL] or rifampin [RIF]) or combined with a fixed concentration (10 µM; 4 mg/L) of NV716. The horizontal dotted line highlights a static effect and the vertical dotted line, the MIC of each antibiotic. All data are means ± SEM (triplicates from 3 experiments).

Figure 3. Intracellular pharmacodynamic parameters as calculated from the Hill equation of the concentration-response curves. The upper panel shows the C^s (in mg/L) for the 4 antibiotics (ciprofloxacin [CIP], doxycycline [DOX], chloramphenicol [CHL] or rifampin [RIF]) against the 4 strains, with horizontal dotted yellow-black lines corresponding to the MIC of the antibiotic in the absence of inhibitor. R.1.4.NV716 and NV731 were used at a concentration of 10 µM and PA β N, at a concentration of 38 µM. The lower panel shows the E_{max} expressed in log₁₀ CFU reduction form the postphagocytosis inoculum. All data are expressed as means ± SEM (triplicates from 3 experiments). Statistical analysis: One-way ANOVA with Dunnett post hoc test *vs* antibiotic alone (*: p < 0.05; **: p < 0.01; ***: p < 0.001 and ****: p < 0.0001).

Figure 4. Concentration-response curves of NV716 alone or combined with antibiotics at their human Cmin or Cmax against intracellular *Pseudomonas aeruginosa* **in a model of THP-1 monocytes.** The graphs show the changes in CFU counts per mg of cell protein from the initial, postphagocytosis inoculum after 24 h of incubation with increasing extracellular concentrations of NV716 alone or combined with ciprofloxacin [CIP], doxycycline [DOX], chloramphenicol [CHL] or rifampin [RIF] at concentrations corresponding to their human C_{min} or C_{max} (see values in Table S1). The combination with CIP at its C_{max} could not be studied, CFU counts being already close to the limit of detection with the antibiotic alone. The horizontal lines highlight respectively a static effect (black dotted line), the effect of the antibiotic alone at its C_{min} (red dotted line) or at its C_{max} (red plain line) and the vertical dotted line, the MIC of NV716 in RPMI-1640 (25 µM). All data are means ± SEM (triplicates from 3 experiments).

R2.3.Figure 5. Influence of NV716 on persisters selected by ciprofloxacin.

A. Persister fraction assay for ciprofloxacin (CIP) alone, NV716 alone or their combination against reference strains. Left: kill curve of a stationary phase culture of PAO1 exposed to ciprofloxacin at 50x MIC over time. The persister fraction is calculated as the difference in CFU counts between control and treated samples at 5 h, when a plateau value has been reached. Right: Percentage of persisters (as compared to the CFU counts of untreated samples) after 5 h of incubation with increasing concentrations of NV716 alone or combined with 50x MIC of ciprofloxacin. All data are expressed as means ± SEM (triplicates from 3 experiments). Statistical analysis: One-way ANOVA with Dunnett post hoc analysis comparing samples exposed to a given concentration of NV716 with the corresponding control (no drug or ciprofloxacin alone): $*$, $p < 0.05$; $**$, $p < 0.01$; $***$, $p < 0.001$; ****, $p < 0.0001$).

B. Kinetics of killing of stationary phase cultures of PAO1 (top) or PAO509 (bottom) by ciprofloxacin at 50x MIC alone or combined with NV716 at different concentrations. NV716 was added at different timings (0h [left], 5h [middle] or 24h [right]), highlighted by the vertical dotted line and the arrow. All data are expressed as means ± SEM (triplicates from 3 experiments).

C. Killing of persisters cells of PAO1 (left) or PAO509 (right) by ciprofloxacin, NV716 or their combinations. Persister cells were isolated after 5 hours of incubation with ciprofloxacin at 50x MIC (persisters), then incubated with either ciprofloxacin (CIP), NV716 at different concentrations, or their combination. All data are expressed as means ± SEM (triplicates from 3 experiments). Statistical analysis: One way ANOVA with Dunnett's post hoc test comparing samples exposed to CIP alone: *, $P \le 0.05$; ** $p \le 0.01$; ****, $P \le 0.0001$.

