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Revision for Antimicrobial Agents and Chemotherapy

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

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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 PABN 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. PABN 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 PABN 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.

1 INTRODUCTION

2

3 Pseudomonas aeruginosa is one of the so-called ESKAPE (Enterococcus faecium, Staphylococcus aureus, Klebsiella pneumoniae, Acinetobacter baumannii, Pseudomonas aeruginosa, and Enterobacter 4 species) pathogens (1) that harbor multiple mechanisms of drug resistance. It is considered by the 5 6 World Health Organization (WHO) as a critical priority species for the search of innovative therapies (2). 7 P. aeruginosa has acquired a series of resistance mechanisms to all classes of commonly used antibiotics (β -lactams, fluoroquinolones, aminoglycosides, and, more recently, polymyxins). Moreover, 8 9 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, 10 and MexXY-OprM, have been associated with increased MICs for a broad range of antimicrobial agents 11 12 (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 13 other Gram-negative bacteria, although they are capable of binding to their respective target in these 14 bacteria when tested in acellular systems (5, 6). 15

16

Another, often neglected, reason for antibiotic failure against P. aeruginosa is the capacity of 17 18 this bacterium to adopt specific lifestyles, such as intracellular survival. Although considered as an 19 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 20 21 macrophages in vivo (12, 13). In these intracellular niches, bacteria are protected from the host 22 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 23 dormant phenotypes that do not respond anymore to antibiotics, like persisters. These are 24 subpopulations of otherwise antibiotic-susceptible bacteria that switch to a transient non-dividing 25 phenotype surviving to high concentrations of bactericidal antibiotics (16). We have recently provided 26 evidence for another species, Staphylococcus aureus, that bacteria surviving to antibiotics 27 intracellularly are persisters, explaining why antibiotics fail to eradicate them (17). Preliminary data 28 29 suggest that the same may hold true for *P. aeruginosa* (18).

30

As P. aeruginosa resistance is worryingly increasing, few new antibiotics have been approved 31 over the last years, and are essentially new derivatives in existing classes with improved activity or 32 reduced susceptibility to resistance mechanisms (19). In this context, reviving old and disused 33 34 antibiotics may appear as an attractive strategy (20), which should be accompanied by an in-depth 35 reassessment of their efficacy according to the modern standards imposed by pharmacokinetic/pharmacodynamic (PK/PD) concepts, including in models of persistent infections like 36 intracellular survival. 37

One of the approaches proposed to increase the potency of drugs that are substrates for efflux 38 39 is to combine them with efflux pump inhibitors (EPIs)(19). PABN (MC 207,110; Phenylalanine-arginine β -naphthylamide (21)) and CCCP (Carbonyl cyanide m-chlorophenyl hydrazone (22)) have been widely 40 used in vitro to increase the accumulation of antibiotics within Gram-negative bacteria, but their 41 toxicity **R1.2**, and inadequate pharmacokinetic properties preclude clinical applications (23, 24). Efforts 42 have been made to synthesize new efflux pumps inhibitors with improved safety profile (25). Among 43 them, polyaminoisoprenyl compounds NV716 and NV731 (see chemical structures in Figure S1) 44 demonstrated their capacity to restore chloramphenicol activity against *Enterobacteria* spp (26) and 45 46 one of them (NV716), that of chloramphenicol and doxycycline against *Pseudomonas aeruginosa* (27), 47 or of florfenicol against Bordetella bronchiseptica (28).

48

The aim of the present study was to evaluate the effects of NV716 in comparison with that of 49 NV731 (less potent against *P. aeruginosa* (27)) and PAβN (reference EPI) in combination with old and 50 disused antibiotics against the intracellular forms of *P. aeruginosa*, using strains expressing or not 51 efflux pumps to infect THP-1 monocytes. We exploited a previously established in vitro 52 53 pharmacodynamic model (11) that allows comparing key pharmacological descriptors of antibiotic 54 intracellular activity, namely their relative potency and maximal efficacy (29). As antibiotics, we selected doxycycline, chloramphenicol (both substrates for efflux (see Table S1) and for which 55 preliminary data do exist regarding the beneficial effect of NV716 (27)), rifampin (not substrate for 56 efflux), and ciprofloxacin, selected as the most active drug against intracellular P. aeruginosa in this 57

- 58 pre-established *in vitro* model (11). The results were examined in the light of the effect of these EPIs on 59 antibiotic MIC and on the selection of persisters in broth culture.
- 60

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*.

67 **RESULTS**

68

69 Antimicrobial susceptibility

Table 1 shows the MICs of antibiotics against PAO1, PAO509 (deleted in the genes encoding 5 efflux 70 71 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 72 (10 µM) NV731 or NV716. These concentrations were selected because commonly used in previous 73 74 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 75 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 77 to chloramphenicol, while 205-2 was less susceptible to ciprofloxacin only. PABN and NV731 reduced 78 79 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) 80 important when testing strains that do not express efflux pumps or those that do, as well as when 81 using rifampin (not substrate for efflux) or other antibiotics (all substrates for efflux). 82

83

84 Cytotoxicity of inhibitors and antibiotics

In a first set of experiments, we evaluated the cytotoxicity of the EPIs and antibiotics in the conditions 85 of their use for intracellular infection by trypan blue exclusion test after 24 h incubation. Infection of 86 THP-1 cells by PAO1 caused 85% cell mortality in the absence of antibiotic. The inhibitors alone caused 87 concentration-dependent cell mortality, which was minimal at the concentration used for intracellular 88 infection studies (Figure 1, left panel). Doxycycline, chloramphenicol, and rifampin showed toxic effects 89 90 at high concentrations, with IC₅₀ (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 equivalent multiples of its MIC (which corresponds to much lower mass concentrations). IC₅₀ values for 92 toxicity of inhibitors, antibiotics or their combinations in non-infected and infected cells are detailed in 93 supplementary Figure S2 and Table S2 but do not differ from what is described in Figure 1. 94

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

Figure 2 shows the concentration-response curves against intracellular PAO1 in the absence of or in 98 the presence of NV716. The activity of antibiotics alone developed following a sigmoidal 99 concentration-response curve, as previously described (11). At low, subMIC concentrations, a ~3 log₁₀ 100 increase in CFUs was noticed over the 24 h incubation period. A static effect was observed at 101 extracellular concentrations close to the MIC of each drug. At the highest concentration tested, the 102 reduction in bacterial counts ranged from 2 log₁₀ CFU decrease for chloramphenicol, doxycycline, and 103 rifampin, to 2.6 log₁₀ CFU decrease for ciprofloxacin. In the presence of NV716 at 10 μ M, all curves 104 (except that of ciprofloxacin) were shifted to the left, meaning that the corresponding antibiotic was 105 more potent, a static effect being reached at lower extracellular concentrations. Furthermore, the 106 107 maximal reduction in intracellular counts was also increased, indicating that the antibiotics were more effective. 108

The same type of experiment was performed with the other EPIs and the other strains. Figure 3 109 110 illustrates the pharmacodynamic parameters calculated based on the Hill equation of these concentration-response curves (see methods for a definition of these parameters), namely the C_s (a 111 112 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 slightly higher than the MICs and systematically reduced of 1 doubling dilution or more by NV716, 114 except for ciprofloxacin against PAO1 and PA0509, which were already highly susceptible to this drug. 115 NV731 and PAβN did not show any significant effect on C_s values. Similarly, NV716 was the only EPI 116 capable of increasing the efficacy of the 4 drugs against all strains, with E_{max} values being at least 0.5 117 118 log₁₀ CFU more negative than those measured for the antibiotics alone.

119

Concentration-response curves of NV716 combined with fixed concentrations of antibiotics against 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

human serum (Table S1 for these values). The data are illustrated in Figure 4. Considering first the 125 126 activity of NV716 alone, a ~3 log₁₀ increase in CFU was observed after 24 h incubation with low NV716 127 concentrations (\leq 10 μ M) 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 ~1 log₁₀ CFU decrease for PAO1, 205-2 and BV-1, to ~2 log₁₀ CFU decrease for PAO509. These data are 130 coherent with those obtained in broth (Figure S3), showing that NV716 did not prevent PAO1 growth 131 132 for concentrations \leq 10 μ M but caused a rapid decrease in bacterial counts at 50 μ M. In the presence of antibiotics at their C_{min}, we noticed that the intracellular concentration-response curves were 133 globally parallel to those obtained with NV716 alone, reflecting thus essentially the activity of the EPI. 134 Three exceptions need however to be highlighted. First, the activity of chloramphenicol was increased 135 for NV716 concentrations > 3 μM against all strains. Second, an improvement in antibiotic activity was 136 137 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 µM), the reduction in CFU counts was 0.5-1.8 log₁₀ 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 0.5-2.2 log₁₀ at NV716 concentrations \leq 10 μ M to 1.4-5.6 log₁₀ at higher NV716 concentrations in 142 comparison with CFU counts measured for the antibiotic alone. 143

144

145 **Persister assay**

The fact that **NV716** increases the intracellular efficacy of antibiotics (improved killing capacity) 146 147 suggests it may decrease the proportion of antibiotic persisters in the population, recently identified as 148 a cause of failure to eradicate intracellular S. aureus with antibiotics (30). We therefore examined the 149 persister fraction surviving to ciprofloxacin (selected as highly bactericidal antibiotic) alone or 150 combined with **NV716** in stationary phases cultures. As shown in the left panel of Figure 5A, 151 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 152 inoculum. The same type of experiment was then performed with PAO1 and PAO509 exposed to 153

NV716 at increasing concentrations and used alone or combined with 50x MIC of ciprofloxacin. The 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 persisters for concentrations \leq 100 μ M but caused a marked reduction in bacterial counts at higher concentrations (3-5 log₁₀ CFU for concentrations > 300 μ M for PAO1 and > 100 μ M for PAO509). Strikingly, **NV716** at lower concentrations (10-100 μ M) was already capable to reduce up to 11- (PAO1) to 28- (PAO509) fold the percentage of persisters selected by ciprofloxacin.

R2.3 To investigate whether NV716 needs to be administered simultaneously with ciprofloxacin to 161 obtain a maximal activity, we repeated killing experiments with stationary phase cultures exposed to 162 50 x MIC of ciprofloxacin alone or combined with **NV716** added at different time points (Figure 5B) (31). 163 164 CFUs were significantly reduced after the addition of **NV716** at any time point, for concentrations \geq 10 165 µM and in both PAO1 and PAO509, indicating that **NV716** does not need to be added at the same time 166 as the drug to be able to increase its activity. In a next step, we aimed at examining whether NV716 action was related to a direct capacity to kill persisters or rather to resensitize them to the killing effect 167 of ciprofloxacin. To this end, we isolated the persisters surviving after 5 h of incubation with 168 169 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 170 171 persisters with ciprofloxacin alone caused only a marginal decrease in CFUs for both strains, confirming 172 the effective isolation of persister cells. When used alone, NV716 caused a significant decrease in CFUs only at the highest concentration tested (250 µM). In contrast, a significant reduction in CFUs was 173 174 already observed when it was combined at 10 μ M with ciprofloxacin for both strains.

176 **DISCUSSION**

177

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.

182

Considering first susceptibility data in broth, we observed a remarkable effect of NV716 on the 183 antibiotic intrinsic activity. While PABN and NV731 moderately reduced the MICs of antibiotics 184 185 (essentially those that are substrates for efflux), NV716 caused a marked decrease in the MICs of all drugs, comprising rifampin (not substrate for efflux; see Table S1) against all strains, including PAO509 186 that does not express efflux pumps or BV1 that expresses a non-functional MexAB-OprM efflux pump. 187 This suggests that its mode of action is not related to the inhibition of efflux only, but probably also 188 relies on its capacity to permeabilize the outer membrane of the *P. aeruginosa* (27). Of note, the 189 synergy with antibiotics is obtained at sub-inhibitory concentrations of NV716, R1.3. indicating that, if a 190 direct effect on bacteria takes place, it is not sufficient to cause bacterial death. PABN has also been 191 shown to permeabilize the outer membrane of Gram-negative bacteria by displacing Mg²⁺ and Ca²⁺ 192 (32), but at concentrations higher than the one we used here, established as adequate to inhibit efflux 193 in princeps studies with this molecule (21). The activity of NV716 and of the two other EPIs is observed 194 at concentrations that do not cause cytotoxicity for eukaryotic cells, suggesting no or minimal 195 196 interaction with eukaryotic membranes.

197

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.

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

219

220 Turning then our attention towards the intracellular maximal efficacy, we observed, for antibiotics used alone, reductions of 2-2.5 log₁₀ CFU with ciprofloxacin, in accordance with our previous 221 results (11), and of 1-2 log₁₀ CFU for the other antibiotics, the lowest effect being observed for 222 chloramphenicol and doxycycline against 205-2, indicating individual differences in responsiveness 223 among strains. In a broader context, the extent of intracellular killing may also differ depending on the 224 bacterial species. An intracellular E_{max} of -1.8 log₁₀ CFU was noticed for rifampin against a small colony 225 variant of S. aureus (36), but the activity of chloramphenicol and doxycycline has not been investigated 226 227 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* 229 tularensis after 24 h of incubation at a concentration close to its MIC (38). For fluoroquinolones, a 230 maximal reduction of 2 to 4 log₁₀ CFU was observed in concentration-response experiments similar to those performed here against S. aureus, Listeria monocytogenes, Legionella pneumophila, Burkholderia 231 232 thailandensis, Yersinia pseudotuberculosis, or Francisella philomiragia (34, 35, 39).

While PABN and NV731 did not affect the efficacy of any antibiotic against none of the strains, 233 234 **NV716** systematically improved it (more negative E_{max}), suggesting an effect on bacterial 235 responsiveness to antibiotics. We can most likely preclude interference related to the release of bacteria out of the cells before lysis or to the R1.6. remaining amount of antibiotics accumulated in cells 236 if considering that (a) NV716 did not increase the cytotoxicity of antibiotics, and (b) charcoal was 237 added to culture plates to adsorb residual antibiotic. Thus, the gain in intracellular efficacy observed in 238 the presence of NV716 is rather suggestive of improved bacterial responsiveness to the drugs. We 239 previously showed an increase in the intracellular efficacy of fluoroquinolones against an insertion 240 mutant in a putative deacetylase in PAO1, which showed a lower persister character in planktonic 241 cultures (18). Here, we found that NV716 was capable of decreasing the proportion of persisters 242 selected by ciprofloxacin in a stationary phase culture at concentrations for which it did not cause any 243 bacterial killing by itself R2.3 and also of reducing at subMIC concentrations the number of persisters 244 when combined with ciprofloxacin but not when used alone, suggesting a mode of action unrelated to 245 a direct bactericidal effect but rather to a resensitization of persisters to the antibiotic (31). Of interest, 246 other molecules capable of disrupting membrane integrity also reduced the residual fraction of 247 persisters upon exposure to antibiotics for different bacterial species (40-42). One of them (SPI009; 1-248 249 ((2,4-Dichlorophenethyl)Amino)-3-Phenoxypropan-2-ol)) was also shown to increase the efficacy of 250 ciprofloxacin against intracellular *P. aeruginosa* (43).

251

This work suffers from some limitations. Firstly, we used a single phagocytic cell line (THP-1 252 monocytes), which does not necessarily represent the repertoire of cells infected by P. aeruginosa in 253 254 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 255 256 accumulation of antibiotics and EPIs in our experimental conditions, but setting-up the necessary 257 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 258 259 pharmacodynamic evaluation of the possible interest of EPIs to restore the activity of old and disused 260 antibiotics against intracellular P. aeruginosa.

If we thus examine our data in a more clinically-oriented perspective, we see that **NV716** is capable of improving the activity of all antibiotics in the range of concentrations achieved in the serum of patients as soon as its extracellular concentration exceeds 3 μ M. Ten times higher concentrations allow to reach a 2-3 log₁₀ CFU decrease when combined with antibiotics at their human C_{max}, pleading for examining the possible interest of this compound *in vivo*.

268 MATERIALS AND METHODS

269

270 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 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.

Bacteria were **R2.m1** incubated overnight at 37°C in Tryptic Soy Agar (TSA; VWR; Radnor, PA). A single 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 (colonyforming unit). Cell culture media (RPMI-1640) and human and fetal bovine sera were from Gibco/ThermoFisher Scientific (Waltham, MA).

284

285 Antibiotics and Inhibitors

Chloramphenicol (potency, 98%), doxycycline (potency, 98%), and rifampin (potency, 98%) were 286 obtained as microbiological standards from Sigma-Aldrich (St Louis, MO), ciprofloxacin HCl (potency, 287 89%) from Bayer (Leverkusen, Germany) and gentamicin sulfate (potency, 60.7%), from PnReac 288 AppliChem (Darmstadt, Germany). Table S1 shows relevant properties of the antibiotics used in this 289 290 study. The reference efflux pump inhibitor Phe-Arg β -Naphthylamide (PA β N; potency, 98%) was 291 purchased from Sigma-Aldrich. The hydrosoluble hydrochloride polyamino-isoprenic salt derivatives 292 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 PABN or in 293 294 water at 10 mM [4 mg/L] for NV716 and NV731, respectively, and stored at -20°C until use.

296 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 $38 \,\mu$ M [20 mg/L](21) PA β N or of 10 μ M [4 mg/L] **NV731** or **NV716** (27).

300

301 Cytotoxicity assessment (Trypan blue exclusion assay)

THP-1 cells (7.5 x 10⁵ cells per mL) were incubated for 24 hours in 96-wells plates with the antibiotics 302 or inhibitors under study over a wide range of concentrations. At the end of the incubation, viability 303 was assessed using trypan blue exclusion test (vital colorant excluded from viable cells) using a 304 305 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 306 307 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 308 mm² for each single square). The percentage of cytotoxicity was evaluated based on the reduction in 309 310 the number of cells according following equation: living to the $\frac{Nb \text{ uncolored cells in controls}-Nb \text{ uncolored treated cells}}{x100}$. cytotoxicity (%) = 311 Nb uncolored cells in controls

312

313 Intracellular Infection

314 All experiments were performed using human THP-1 monocytes cultured in RPMI-1640 medium 315 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% 316 317 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 318 eliminated by incubation for 1 h with gentamicin at 50x MIC. After 3 washings with PBS, infected cells 319 320 were resuspended in the original volume of RPMI-1640 with 10% FBS. The post-phagocytosis inoculum was consistently 5 to 7 x 10^5 CFU/mg of cell protein (defined as time zero). Infected cells were then 321 322 incubated for 24 h with antibiotics over a broad range of concentrations (0.003-100x their MIC) in 12well plates. After 24 h of incubation, infected cells were pelleted by centrifugation, washed once with 323 324 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 325 326 supplemented with 2g/L charcoal (to avoid carry-over effect, especially in samples that had been incubated with high antibiotic concentrations [R1.5.no difference in colony counts was observed 327 between plates supplemented or not with charcoal for samples exposed to low antibiotic 328 concentrations, ruling out any interfering effect of charcoal in the assay]) and protein content using a 329 commercially available kit (Bio-Rad DC Protein Assay, Bio-Rad Laboratories, Hercules, CA). The activity 330 was expressed as the change in CFU (normalized by mg of cell protein) from the initial inoculum after 331 24 hours of incubation. The data were used to fit a sigmoidal function and calculate pharmacodynamic 332 parameters based on the corresponding Hill-Langmuir equation (apparent static concentrations $[C_s]$, i.e. 333 334 extracellular concentration resulting in no apparent intracellular growth, and maximal relative efficacy [E_{max}], i.e. maximal decrease in bacterial counts compared to the post-phagocytosis inoculum as 335 336 extrapolated for an infinitely large antibiotic concentration).

337

338 Persister assay and isolation

A single colony from overnight culture of PAO1 and PAO509 on TSA was R2.m1.added to in 200mL 339 MHB-CA in a 1 L flask and incubated for 24 h under agitation (130 rpm) at 37°C. 10 mL of the bacterial 340 341 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 342 on TSA supplemented with 2 g/L charcoal for CFU counting. The persister fraction was defined as the 343 number of CFU growing at the end of this experiment as compared to the initial inoculum (46). R2.3For 344 experiments using isolated persisters, stationary-phase cultures were incubated with ciprofloxacin (50x 345 MIC) during 5 hours, i.e. condition for which a plateau of killing was reached. The remaining bacteria 346 were washed twice with sterile Phosphate Buffered Saline (PBS), centrifuged (5,200 g, 15 min, 4°C) and 347 348 used for killing assays (31).

349

350 Curve fittings and statistical analyses.

351 Curve fittings were performed with GraphPad Prism (version 8.4.3) software for Windows (GraphPad 352 Prism Software, San Diego, CA). Pharmacodynamic parameters were calculated based on Hill equations

- 353 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.

356

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			MIC (mg/L) ^a and (fold-dilution decrease vs antibiotic alone)							ne)
Strains	Phenotype	Reference	Antibiotics ^b	AB	+ΡΑβΝ	ΡΑβΝ	+NV731	NV731	+NV716	<mark>R1.6</mark> NV716 alone (µM)
				alone	(38µM)	alone	(10µM)	alone	(10µM)	
				(mg/L)	[20mg/L]	(μM)	[4mg/L]	(μM)	[4mg/L]	
PAO1	Wild-type	(47)	CIP	0.25	0.06 (2)	>200	0.06 (2)	>250	0.03 (3)	50
			DOX	16	4 (2)		4 (2)		1 (4)	
			CHL	32	8 (2)		16 (1)		1 (5)	
			RIF	16	8 (1)		8 (1)		0.125 (7)	
PAO509	PAO1[Δ(MexAB-	(48)	CIP	0.016	0.008 (1)	>200	0.016 (0)	>250	0.008 (1)	25
	OprM), Δ(MexCD-		DOX	1	1 (0)		0.125 (3)		0.03 (4)	
	OprJ), Δ(MexJK),		CHL	2	2 (0)		1 (1)		0.125 (4)	
	Δ(MexXY), Δ(MexEF-		RIE	16	4 (2)		8 (1)		0.06(7)	
	OprN)]		NII .	10	4 (2)		0(1)		0.00(7)	
BV1	Clinical isolate	(44)	CIP	1	1 (0)	>200	0.5 (1)	>250	0.125 (3)	50
	(expressing		DOX	64	8 (3)		2 (5)		0.125 (9)	
	truncated, non-		CHL	8	4 (1)		0.5 (4)		0.06 (7)	
	functional MexAB-		RIF	32	16 (1)		4 (3)		0.008 (12)	
	OprM)		-	02	10 (1)		. (•)		(12)	
205-2	Clinical isolate (overexpressing functional MexAB)	(45)	CIP	4	2 (1)	>200	4 (0)	>250	1 (2)	50
			DOX	16	4 (2)		8 (1)		0.5 (5)	
			CHL	64	32 (1)		32 (1)		1 (6)	
			RIF	16	8 (1)		16 (0)		0.125 (16)	

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 PA β N 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 IC₅₀ 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 μ M for NV716 and 38 μ M for PA β N). All data are means ± 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 μ M; 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 μ M and PA β N, at a concentration of 38 μ M. The lower panel shows the E_{max} expressed in log₁₀ CFU reduction form the post-phagocytosis 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 C_{min} or C_{max} 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, post-phagocytosis 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 μ M). All data are means \pm 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 \pm 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).

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 \pm SEM (triplicates from 3 experiments). Statistical analysis: One way ANOVA with Dunnett's post hoc test comparing samples exposed to CIP alone: *, P \leq 0.05; ** p \leq 0.01; ****, P \leq 0.0001.

