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1 Low concentrations of nitric oxide modulate *Streptococcus pneumoniae* biofilm

- 2 metabolism and antibiotic tolerance
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20 Running head: Nitric Oxide Treatment of *S. pneumoniae* Biofilms

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24 Abstract

25	Streptococcus pneumoniae is one of the key pathogens responsible for otitis media (OM),
26	the most common infection in children and the largest cause of childhood antibiotic
27	prescription. Novel therapeutic strategies that reduce the overall antibiotic consumption
28	due to OM are required because although widespread pneumococcal conjugate
29	immunization has controlled invasive pneumococcal disease, overall OM incidence has not
30	decreased. Biofilm formation represents an important phenotype contributing to the
31	antibiotic tolerance and persistence of <i>S. pneumoniae</i> in chronic or recurrent OM. We
32	investigated the treatment of pneumococcal biofilms with nitric oxide (NO), an endogenous
33	signaling molecule and therapeutic agent that has been demonstrated to trigger biofilm
34	dispersal in other bacterial species. We hypothesised that addition of low concentrations of
35	NO to pneumococcal biofilms would improve antibiotic efficacy and higher concentrations
36	exert direct antibacterial effects. Unlike in many other bacterial species, low
37	concentrations of NO, did not result in <i>S. pneumoniae</i> biofilm dispersal. Instead, treatment
38	of both in vitro biofilms and ex vivo adenoid tissue samples (a reservoir for S. pneumoniae
39	biofilms) with low concentrations of NO enhanced pneumococcal killing when combined
40	with amoxicillin-clavulanic acid, an antibiotic commonly used to treat chronic OM.
41	Quantitative proteomic analysis using iTRAQ (isobaric tag for relative and absolute
42	quantitation) identified 13 proteins that were differentially expressed following low-
43	concentration NO treatment, 85% of which function in metabolism or translation.
44	Treatment with low-concentration NO therefore appears to modulate pneumococcal
45	metabolism and may represent a novel therapeutic approach to reduce antibiotic tolerance
46	in pneumococcal biofilms. [245]

47 Introduction

48 *Streptococcus pneumoniae* is a Gram-positive bacterium that asymptomatically 49 colonizes the human nasopharynx. This opportunistic pathogen is responsible for invasive 50 diseases such as pneumonia, bacteremia and meningitis, and localized mucosal infections 51 such as otitis media and sinusitis. Globally, these infections represent a significant burden 52 of disease, particularly in the very young and the elderly. The World Health Organization 53 estimates that 1.6 million deaths occur annually due to pneumococcal infections, 54 accounting for around 11% of the mortality in children under 5 (1). The majority of deaths 55 occur in developing countries where invasive pneumococcal disease remains one of the 56 most common fatal childhood illnesses. 57 Pneumococcus is a leading pathogen in otitis media (OM), the most common 58 infection in young children and a principal reason for repeated physician visits. Upon 59 colonization with *S. pneumoniae* and the establishment of carriage in children, bacteria may 60 access the middle ear space by retrograde ascent from the nasopharynx due to the 61 presence of fluid and/or disruption of mucociliary clearance. Recurrent or chronic otitis 62 media causes much pain and morbidity at high economic cost to society (2). In spite of 63 concerns about the selection of antibiotic resistant bacteria, OM continues to be the primary reason for antibiotic prescription in children (3–5). In addition, although 64 65 pneumococcal conjugate vaccines (PCV) have reduced vaccine type invasive pneumococcal 66 disease, PCVs have not led to a decrease in the incidence of otitis media due to 67 pneumococci, most likely due to non-vaccine type replacement (1, 6, 7). Novel treatments 68 for pneumococcal infection are therefore needed to address the problem of recurrent 69 and/or chronic infections in children.

70	Infections occur following a breach of the mucosal epithelia subsequent to
71	colonization and despite being a prerequisite for infection, little is known about how
72	pneumococci colonize and persist in the nasopharynx. However, a growing body of
73	literature suggests that bacterial biofilm development plays a prominent role in
74	colonization and disease. In situ investigation of paediatric middle ear biopsies indicated
75	that pneumococcal biofilms were present on the middle ear mucosal epithelium in children
76	with chronic OM but not in children without chronic OM (8, 9). S. pneumoniae biofilms have
77	also been identified <i>in situ</i> on adenoid mucosal epithelia from children undergoing
78	adenoidectomy for the treatment of infective (chronic OM) or inflammatory (obstructive
79	sleep apnoea - OSA) otolaryngological disease consistent with the adenoid serving as a
80	reservoir of pathogens that may contribute to infection under circumstances that favor
81	middle ear infection (10, 11). More recently, pneumococcal biofilms have been investigated
82	with animal models and epithelial cell models (12–15).
83	Biofilms are highly adaptive surface-associated microbial aggregates that allow
84	bacteria to survive the diverse stressful conditions encountered in the host such as nutrient
85	limitation and host immune responses (16–19). The increased tolerance of biofilm bacteria
86	to conventional antibiotic therapeutic concentrations compared with their planktonic
87	counterparts also poses a significant problem in clinical settings, as does their propensity
88	to acquire further antimicrobial resistance via horizontal gene transfer, underscoring the
89	need for novel therapeutic strategies to limit the pneumococcal biofilm phenotype during
90	disease (20, 21).

91 Nitric oxide (NO) is an important signaling molecule that is ubiquitous in both
92 eukaryotes and prokaryotes, bridging the boundaries between host and pathogen. In the

94	inducible nitric oxide synthases (iNOSs) in epithelial and phagocytic cells with its
95	production resulting in damage to bacterial cell membranes and DNA (22). Bacteria have
96	also been shown to possess NOSs that oxidise L-arginine to produce the low concentrations
97	of NO observed in several species (23, 24). Multiple regulatory systems have been
98	identified that mediate the diverse responses of bacteria to NO, including conferring
99	protection from oxidative stress and playing a role in toxin biosynthesis (23, 25). In
100	biofilms, however, low concentrations of exogenous NO have been shown to trigger a
101	dispersal response in several bacterial species including Pseudomonas aeruginosa,
102	Escherichia coli and Staphylococcus epidermidis, a response associated with increased
103	antibacterial efficacy when used as an adjuvant in conjunction with antibiotics (26–28).
104	Furthermore, NO-releasing nanoparticles and gaseous NO have been shown to exert potent
105	antimicrobial effects against P. aeruginosa, Streptococcus pyogenes and Enterococcus
106	faecalis (29–31).
107	Previous investigations into the role of NO in pneumococcal infection have given
108	conflicting results. In a murine pneumonia model NO was found to be associated with
109	increased bacterial loads and reduced survival during bacteremia in wild-type mice but not
110	in NOS2-deficient mice. In contrast during pneumonia following intranasal infection NO
111	decreased pneumococcal viability in the lung in both mouse strains via a direct
112	antibacterial effect (32).
113	Because several studies using biofilm models have shown that low concentrations of
114	NO trigger release of bacteria from the biofilm in various bacterial species we hypothesized
115	that adjunctive NO treatment would also improve the efficacy of antibiotic killing of

human host, NO plays an important role in the innate immune response and is produced by

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116 pneumococci in biofilms. We tested this hypothesis using in vitro and ex vivo pneumococcal 117 biofilms. Biochemical studies and a high-throughput quantitative proteomic approach were 118 utilized to interrogate possible mechanisms of action.

119

120 Methods

121 Bacterial strains and growth conditions. Clinical isolates of Streptococcus pneumoniae 122 serotypes 14 (ST124), 19F and 23F (33), and the laboratory strain D39 (serotype 2) were 123 selected to evaluate the in vitro antibacterial efficacy of NO and antibiotic adjunctive 124 treatment. Strains were subcultured from frozen stocks onto Columbia blood agar (CBA) 125 plates (Oxoid, U.K.) as described (33). Briefly, cultures were incubated at $37^{\circ}C/5\%$ CO₂ and 126 colonies re-suspended in fresh Brain Heart Infusion (BHI) broth (Oxoid, U.K.) for

127 experiments.

128

129 Planktonic experiments. Flat-bottomed 96-well culture plates (Fisher Scientific, U.K.) 130 were inoculated with $\sim 1.0 \times 10^7$ bacteria per well (mid-exponential planktonic cultures) 131 grown in BHI. All treatments were prepared in BHI. Sodium nitroprusside dihydrate (SNP) 132 was added to wells for final concentrations ranging from 1 μ M to 10 mM; diethylamine 133 NONOate (DEA/NO; sodium salt), sodium nitrate (NaNO₃), sodium nitrite (NaNO₂) and 134 potassium cyanide (KCN; all Sigma-Aldrich, U.K.) were added at a final concentration of 1 135 mM, and carboxy-PTIO potassium salt and L-methionine (both Sigma-Aldrich, U.K.) were 136 added at final concentrations of 50 μ M and 1 mM respectively. BHI was added in place of 137 treatments for all untreated controls. BHI alone was used to measure background changes in absorbance. Cultures were incubated at 37°C/5% CO2 and absorbance (OD595) 138

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measured every 30 min over 2 hours using an EZ Read 400 spectrophotometer (Biochrom)(n=3).

141

142 In vitro biofilm experiments. For biofilm formation mid-exponential planktonic cultures 143 grown in BHI were used to inoculate individual wells of untreated polystyrene 6-well 144 plates (1 x 10⁸ cells per well) (Corning Incorporated, Costar, U.S.A.), and supplemented 145 with fresh BHI diluted 1:5 in distilled H₂O. Cultures were incubated at 37°C/5% CO₂ with 146 replacement of warm, fresh diluted 1:5 BHI daily for 2 to 7 days. 147 Prior to treatment medium was removed and biofilms washed twice using diluted 148 1:5 BHI. NO donor (SNP and DEA/NO) treatments were prepared fresh in diluted 1:5 BHI 149 and added to wells at final concentrations ranging from 100 nM to 1 mM. For adjuvant 150 experiments amoxicillin + clavulanic acid (AMC) was added at a final concentration of 151 300/60 µg/ml. Biofilms were incubated at 37°C/5% CO₂ for 2 hours after which the NO 152 donors or inhibitors were removed and the remaining biofilm rinsed twice with diluted 1:5 153 BHI. Biofilms were resuspended in Hank's balanced salt solution (HBSS) as described (20). 154 Briefly, biofilms were scraped and vortexed, both resuspended biofilms and removed supernatants were diluted in HBSS, spot plated onto CBA plates and incubated at 37°C/5% 155 156 CO_2 . To assess total biofilm biomass 100 µl of the resuspended biofilms were diluted 10-157 fold in 1:5 BHI and turbidity measured by absorbance (OD595) using a Jenway 6300 158 spectrophotometer. All assays were performed on 48 h biofilms using 2 technical replicates 159 of 2 biological replicates (n=4).

160

161 Confocal Laser Scanning Microscopy (CLSM). Mid-exponential planktonic cultures of 162 strain ST124 (n=3) were grown in BHI and used to inoculate 35 mm untreated glass 163 bottom CELLview cell culture dishes (Greiner Bio One, U.K.) and supplemented with fresh 164 1:5 BHI. Biofilms were grown under static conditions at 37°C/5% CO₂ for 48 h replacing 165 medium daily with fresh 1:5 BHI. Biofilms were treated with 1 mM SNP in 1:5 BHI or 1:5 166 BHI (untreated control) at $37 \circ C/5\%$ CO₂ for 2 hours. Treatments were removed and the 167 remaining biofilm rinsed twice with HBSS. Biofilms were stained with Live/Dead BacLight 168 Bacterial Viability Kit (Life Technologies, U.S.A.) according to manufacturer instructions. 169 Biofilms were examined immediately with an inverted Leica SP8 LSCM system using a 63x 170 oil immersion lens and sequential scanning of 1 µm sections (Leica Microsystems, Milton 171 Keynes, U.K.). To remove background eDNA staining, the Syto9 fluorescence intensity 172 threshold was set to that of planktonic pneumococci. Images were analysed using Leica LCS 173 Software.

174

175 Scanning Electron Microscopy (SEM). Serotype 14 (ST124) biofilms were grown for 48 h 176 in 6-well plates containing ethanol-sterilized 13 mm glass cover slips (V.W.R., U.K.). 177 Biofilms were treated with 1 mM SNP in 1:5 BHI or 1:5 BHI (untreated control) at 37°C/5% 178 CO_2 for 2 hours then processed for SEM as described (33). Biofilms were imaged using an 179 FEI Quanta 200 scanning electron microscope.

180

181 Protein extraction and iTRAQ (isobaric tag for relative and absolute quantitation)

182 labelling. Comparative analyses of protein expression between biofilms treated with 100

183 μ M SNP for 2 h at 37°C/5% CO₂ and untreated biofilms were performed on 3 technical

184 replicates of 3 biological replicates. Protein extraction and iTRAQ labelling were performed 185 as described (33).

186

187 Mass spectrometry, peak list generation, and database searching. Mass spectrometry, 188 peak list generation and database searching were performed as previously described (33). 189 Inclusion criteria for quantitative analysis were set at ≥ 3 peptide matches, ≥ 50 protein 190 score, $\geq 5\%$ sequence coverage (p < 0.05). Comparative protein data with >1.3 and <0.77 191 ratios were identified as having differential expression. For qualitative identification the 192 inclusion criteria were 2 peptide matches, \geq 50 protein score, and \geq 5% sequence coverage. 193

194 *Ex vivo* adenoid experiments. Adenoids were obtained from paediatric patients <12 years 195 of age undergoing adenoidectomy for the treatment of suspected inflammatory or infective 196 Ear, Nose and Throat (ENT) disease (n=11). Adenoids were collected on ice in sterile HBSS 197 + 5% fetal bovine serum (FBS; Sigma-Aldrich, U.K.), washed twice with HBSS to remove any 198 unattached bacteria, then dissected into four equal-sized sections with similar luminal 199 surface coverage. Tissue sections were normalized for weight and treated with $100 \,\mu M$ 200 SNP alone, 300/60 μg/ml AMC alone or 100 μM SNP + 300/60 μg/ml AMC in HBSS/10% 201 FBS for 2 h at 37°C/5% CO₂. Untreated control tissue was treated with HBSS/10% FBS 202 alone. Tissue sections were washed twice with 10 ml HBSS, macerated in 1 ml HBSS 203 through a 100 µm nylon cell strainer (Fisher Scientific, U.K.) and bacterial suspensions 204 serially diluted and spot plated onto CBA plates.

205

Statistical analyses. Statistical analysis of *in vitro* planktonic and biofilm data was
performed using one-way ANOVA and Tukey's multiple comparisons tests. Analysis of *ex vivo* adenoid data was performed using a Wilcoxon Signed Ranks test. Comparative data
reported as p<0.05 were considered statistically different.

210

211 Results

212 Treatment with the NO donor SNP decreased viability of planktonic cells and the cell 213 population remaining within in vitro biofilms. Since low-concentrations of NO have 214 been shown to result in the release or dispersal of other bacterial species from biofilms, we 215 first tested the hypothesis that low-concentrations of NO would have a similar effect on 216 pneumococcal biofilms. Established 48 hour biofilms were treated for 2 hours with a range 217 of NO concentrations generated from different concentrations of the NO-donor SNP (100 218 nM - 1 mM; Fig. 1). Measurement of biofilm biomass using turbidity, and viability using 219 colony forming unit (CFU) enumeration, respectively, indicated that treatment with low 220 concentrations of NO (100 nM to 100 µM SNP) did not have a similar effect on 48h 221 pneumococcal biofilms. Treatment with 1 mM SNP however, resulted in a significant 222 reduction in the biomass and a 3-log reduction in the number of viable cells remaining 223 within the biofilm ($p \le 0.001$). These results suggested that at 1mM SNP treatment may 224 either be triggering the release of *S. pneumoniae* from the biofilm or had a direct 225 bactericidal effect. To distinguish between these possibilities the number of viable cells 226 present in the biofilm supernatant was measured following treatment with 1 mM SNP. 227 Results indicated that there was a significant reduction in planktonic pneumococcal cells 228 suspended in the supernatant as well as in the biofilm following NO treatment (Fig. 2,

229	p≤0.05). Since NO treatment of biofilms formed by other bacterial species typically results
230	in increased numbers of bacterial CFUs in the supernatant (24) these data indicated that
231	treatment with higher concentrations of SNP had a direct antibacterial killing effect.
232	Furthermore, both SEM and CLSM imaging of biofilms treated with 1 mM SNP
233	demonstrated no significant change in biofilm ultrastructure following treatment
234	confirming the lack of dispersal (Fig. 3a-b, 3e-f). CLSM imaging did, however, demonstrate
235	a reduction in the number of viable cells remaining within the biofilm following treatment,
236	commensurate with the observed 3-log reduction in CFUs (Fig. 3c-d). To confirm that
237	higher NO concentrations were toxic for pneumococcus, mid-exponential planktonic
238	cultures were treated with the same range of concentrations of SNP. Cultures treated with
239	SNP concentrations between 1 μM to 10 mM for 2 hours showed a significant reduction in
240	growth with 500 μM to 5 mM SNP (p<0.05), and complete cessation of growth with
241	concentrations greater than 5 mM (Fig. 4).
242	
243	The response of S. pneumoniae to treatment with SNP was mediated by NO. Having

244 determined that both planktonic and biofilm pneumococci responded to SNP treatment, we 245 next wished to confirm that the response was indeed mediated by NO and not due to other 246 NO metabolites, intact SNP or SNP breakdown products other than NO. Treatment with 1 247 mM KCN, a control for the possible effect of cyanide anion liberation from the SNP 248 molecule, caused no reduction in pneumococcal growth confirming that the reduction in 249 viability compared with untreated bacteria was not due to cyanide toxicity (p=0.528; Fig. 250 5a). Moreover, treatment with 1 mM DEA/NO, an alternative NO-donor molecule that is 251 chemically and mechanistically distinct from SNP, resulted in a significant decrease in

253	with the NO-scavenger cPTIO abrogated the response to SNP (p=0.008, Fig. 5b). These
254	results indicated that the response to SNP treatment was NO-mediated.
255	In contrast, treatment with 1 mM nitrate (p=0.321) or nitrite (p=0.078) failed to
256	significantly reduce pneumococcal growth, indicating that the oxidative breakdown
257	products of NO were not responsible for the observed reduction in viability (Fig. 5a). Since
258	peroxynitrite (ONOO ⁻) is an extremely toxic molecule that can be produced by reaction of
259	NO with superoxide (O_2^{-}) to cause damage to DNA, proteins and lipids (24, 29), we further
260	tested whether ONOO ⁻ toxicity might be involved in reducing pneumococcal viability
261	during SNP treatment by using the ONOO ⁻ scavenger L-methionine. Indeed, the response to
262	SNP was also reduced by the presence of L-methionine ($p=0.005$) suggesting that
263	extracellular $ONOO^{-}$ formation secondary to reaction with O_{2}^{-} might be responsible for the
264	bactericidal effects of higher concentrations of NO on pneumococcal cells (Fig. 5c).
265	
266	NO adjunctive treatment of <i>in vitro</i> pneumococcal biofilms enhanced antibiotic
267	effectiveness. Others have shown that NO combined with antibiotic treatment resulted in
268	an additional reduction in the viability of biofilms in several types of bacteria (28). We
269	therefore tested whether NO treatment of pneumococcal biofilms could further reduce
270	bacterial viability when used as an adjunctive treatment in conjunction with a conventional
271	antibiotic used to treat otitis media. Serotype 2 strain D39 biofilms and biofilms from 3
272	different clinical isolates representative of serotypes 14 (ST124), 19F and 23F $$ and based
273	on their high isolation frequency in OM (20) and high rates of antibiotic recalcitrance were

growth, similar in extent to treatment with 1 mM SNP (p=0.013, Fig, 5a), whilst treatment

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274	used to assess NO adjunctive treatment (34, 35). Established biofilms were treated with 1
275	mM SNP and 300/60 $\mu g/ml$ AMC for 2 h and the viability of the remaining pneumococcal
276	biofilm bacteria assessed by CFU enumeration. Treatment of ST124, 19F and D39 biofilms
277	with the NO donor alone resulted in a 2-log reduction in viable bacteria, whereas treatment
278	of 23F biofilms resulted in a 3-log reduction (Fig. 6). AMC treatment alone resulted in a 3-
279	log reduction in ST124 and 23F biofilm viability, and a 2-log reduction in 19F and D39
280	biofilm viability (Fig. 6). Combined NO/AMC treatment, however, resulted in a 3-log
281	reduction in 19F and D39, a 5-log reduction in ST124, and complete killing of 23F when
282	compared with untreated biofilms (Fig. 6), and a significant reduction in viable
283	pneumococci in biofilms compared with antibiotic treatment alone ($p < 0.05$).
284	

285 Combined antibiotic and NO treatment enhanced ex vivo killing of S. pneumoniae on 286 adenoid tissue. Adenoids have been shown to act as a reservoir for S. pneumoniae and 287 biofilm bacteria, and may provide a source for infection in some cases of chronic otitis 288 media (10, 11, 36). Following research ethics committee approval and informed parental 289 consent (NHS REC 09/H0501/74) we subsequently examined adenoid tissue ex vivo that 290 was culture positive for *S. pneumoniae* from children <12 years of age undergoing 291 adenoidectomy for the treatment of suspected inflammatory or infective ENT disease to 292 evaluate the effect of NO on S. pneumoniae colonized mucosal epithelia (Fig. 7). Adenoid 293 sections (n=11) were treated with 100 μ M of SNP alone, 300/60 μ g/ml AMC alone, or 294 treated with both NO and AMC to determine if NO adjunctive therapy increased antibiotic 295 efficacy (Fig. 7). Treatment with 100 μ M of the NO-donor SNP alone did not significantly 296 reduce colonized pneumococci determined by CFUs (p=0.722) and AMC treatment alone

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Antimicrobial Agents and Chemotherapy resulted in a 2-log reduction compared with the untreated adenoid (p=0.005). However,
similar to results with *in vitro* biofilms, combined NO and AMC resulted in a significant
reduction of pneumococci CFUs on *ex vivo* adenoid tissue by nearly 3 logs (p=0.005)
compared with untreated adenoid tissue and further reduced the number of CFUs
compared with AMC alone (p=0.04).

302

303 Treatment of *in vitro S. pneumoniae* biofilms with NO induced a change in

304 translational and metabolic protein expression. Since low dose NO treatment of 48 h 305 pneumococcal biofilms did not appear to be cytotoxic or induce dispersal we used a high-306 throughput gel-free proteomic approach to investigate whether NO treatment induced 307 changes in protein expression to shed further light on the potential mechanisms involved. 308 Previous data from our lab demonstrated that a total of 112 proteins were differentially 309 expressed during biofilm development using iTRAQ (inclusion criteria \geq 3 peptide matches; 310 >5% sequence coverage and a 50+ protein score; p<0.05) (33). Of these, 13 proteins were 311 differentially expressed in established pneumococcal biofilms treated with NO for 2 h, 312 compared with biofilms treated with HBSS alone (Fig. 8). Eighty five percent (11/13) of 313 these proteins were involved in *S. pneumoniae* translation or metabolism (Figure 9a). Five 314 ribosomal proteins, all of which demonstrated significantly decreased expression in the 315 biofilm phenotype, exhibited increased expression following NO treatment, suggesting a 316 modulation of translational capacity similar to planktonic levels (33). Additionally, 6 317 metabolism-associated proteins were differentially expressed following NO treatment: 3-318 ketoacyl-(acyl-carrier-protein) reductase and PTS system fructose-specific II ABC

319 components were upregulated upon NO treatment, whereas arginine deiminase (ArcA), a

Antimicrobial Agents and Chemotherapy 320 PTS system mannose specific IID component, and 2 individual alcohol dehydrogenases 321 exhibited decreased expression following NO treatment. The significant reduction in ArcA 322 expression is of particular interest given its >4-fold increase in expression during biofilm 323 growth (33). Two other proteins, a SPFH domain-containing protein and a hypothetical 324 protein were also identified as having increased expression following NO treatment. These 325 data were further supported by the qualitative identification of 12 proteins with 326 differential expression following NO treatment including 7 associated with pneumococcal 327 metabolism (Figure 9b). 328 329 Discussion

330 Consistent with other studies, S. pneumoniae biofilms were more tolerant to 331 antibiotic treatment than planktonic pneumococci, including an antibiotic commonly used 332 to treat otitis media (13, 20, 37). However, pneumococcal biofilm antibiotic tolerance was 333 significantly diminished (by up to 2 logs) when accompanied by adjunctive treatment with 334 a low concentration of NO. Additionally, we demonstrated biochemically that reduced 335 viability of pneumococci in planktonic and biofilm growth conditions was mediated by NO, 336 and the oxidative breakdown products of NO, nitrite and nitrate, did not mimic this effect. 337 Although NO treatment reduced *in vitro* pneumococcal biofilm CFUs, our data suggest that 338 the anti-pneumococcal effect was not due to a dispersal of bacteria. Rather, higher 339 concentrations of NO demonstrated a direct antibacterial effect on pneumococcal growth. 340 Treatment with 1 mM SNP resulted in a decrease in biofilm viability by up to 3 logs, and in 341 the number of viable cells in the surrounding supernatant. These data were commensurate 342 with CLSM and SEM imaging which demonstrated no obvious changes in biofilm

343	ultrastructure, but did reveal a significant reduction in biofilm viability. A similar response
344	was observed using the structurally distinct NO donor, DEA/NO. The reduced antibacterial
345	response in the presence of the NO scavenger cPTIO, and the lack of response to nitrite and
346	nitrate, indicated that the antimicrobial effects were indeed NO-mediated, and not
347	associated with the formation of NO_3^- and NO_2^- which have also been shown to increase
348	antibiotic efficacy in <i>P. aeruginosa</i> biofilms (38). The specificity of the NO-mediated
349	response, along with the reduction in the planktonic growth rate observed with ${\geq}500~\mu M$ of
350	SNP, suggest a direct effect on growth and/or regulation of metabolism.
351	We also investigated the hypothesis that low concentrations of NO could enhance
352	antibiotic efficacy in the treatment of pneumococcal biofilms. In vitro S. pneumoniae
353	biofilms were more tolerant than planktonic pneumococci to AMC, an antibiotic commonly
354	used to treat pneumococcal infections, commensurate with other studies (20, 21). Results
355	indicated that the addition of 1 mM SNP significantly enhanced antibiotic efficacy by 1 to 2 $$
356	logs in each of four strains tested, three of which represent serotypes (14, 19F and 23F)
357	that are predominantly isolated from paediatric ENT patients and are associated with
358	developing antibacterial recalcitrance (34, 35). Thus, when combined with NO, antibiotic
359	tolerance within the biofilm was significantly diminished.
360	Pneumococcal interactions with epithelial cells have been shown to be important for
361	colonization and biofilm formation (12, 13), and pneumococci are commonly present on
362	adenoids from children with chronic OM or OSA (10, 11). We therefore used adenoid upper
363	respiratory mucosal epithelial tissue colonized with pneumococci to further determine if a
364	combination of NO and AMC might enhance pneumococcal killing using a lower
365	concentration than for <i>in vitro</i> biofilms. Treatment of <i>ex vivo</i> adenoid tissue culture

366	positive for pneumococcus and rinsed to remove unattached bacteria, resulted in a
367	significant reduction in pneumococcal CFUs on adenoid sections treated with antibiotic
368	alone, but not NO-donor alone on treated tissue from the same adenoid sample. When
369	combined however, NO/AMC treatment resulted in a significantly enhanced reduction in
370	CFUs. These results suggest that low-concentrations of NO rendered biofilm pneumococci
371	more susceptible to antibiotic killing. These results are also consistent with other data
372	showing that NO reduced <i>S. pneumoniae</i> viability <i>in vivo</i> using iNOS knock-out mice (32).
373	The difference in susceptibility to S. pneumoniae bacteremia and lung infection between
374	wild-type and iNOS ^{-/-} mice following intravenous infection versus intranasal infection may
375	be due to the differential effects of inducible and constitutive NO production by endothelial
376	and epithelial cells, suggesting that NO concentrations in the host are tissue dependent and
377	regulated locally, and that NO in different mucosal sites is important in anti-pneumococcal
378	host responses.

379 NO is constitutively synthesised in the respiratory epithelium and upregulated in 380 response to infection or inflammation (39). Since *S. pneumoniae* is highly adapted to the 381 upper airway, a compartment characterised by higher constitutive NO concentrations 382 compared with the lower airways as evidenced by exhaled breath analysis (40), it is likely 383 that this bacterium has the ability to respond to NO. Moreover, since epithelial cells also 384 produce iNOS, we speculate that exogenous NO may combine with endogenous NO levels to 385 achieve the higher concentrations of NO sufficient to produce an enhanced anti-386 pneumococcal response observed in vitro.

NO signaling has been shown to elicit different responses in bacterial biofilms,
however its role in mediating dispersal from a biofilm by the reversal of a genetically

389	determined program inducing biofilm development via cyclic di-GMP (c-di-GMP) has
390	garnered significant interest due to its potential as a treatment strategy for biofilm-
391	associated infections (24). In the model biofilm bacterium <i>P. aeruginosa</i> the NO dispersal
392	response results in increased motility and metabolic activity characteristic of the
393	planktonic (colonizing) phenotype allowing propagation to new sites within an
394	environmental niche (41). The single cell phenotype, as well as the increased metabolic
395	and replicative capacity associated with the dispersed planktonic bacteria, are
396	hypothesized to reduce biofilm antibiotic tolerance following NO treatment. Treatment of
397	pneumococcal biofilms with low concentrations of the NO donor, SNP (100 nM to 100 $\mu\text{M}),$
398	shown to disperse biofilms of other bacterial species resulted in no significant changes in
399	biomass or viability at these concentrations. This is unsurprising since S. pneumoniae is a
400	non-motile bacterium, which lacks proteins possessing the common EAL, GGDEF and HD-
401	GYP domains that are involved in the turnover of the secondary messenger c-di-GMP
402	known to mediate dispersal in other bacteria (27, 28).
403	Rather, proteomic analyses suggested that NO induced a shift to a planktonic-like
404	profile in a subset of proteins, notably those involved in metabolism and translation.
405	Proteomic analyses of S. pneumoniae remaining within biofilms following treatment with a
406	low concentration (100 μM) of NO indicated that 13 of 112 quantitatively identified
407	proteins were differentially expressed, indicating that NO was not directly cytotoxic at
408	concentrations of 100 μ M. The increased expression of five ribosomal proteins indicated
409	up-regulation of translational capacity, which was previously shown to be substantially
410	down-regulated in established pneumococcal biofilms (33). We previously hypothesized

411	that the decreased translation exhibited by the biofilm phenotype may contribute to
412	antibiotic tolerance in the biofilm mode of growth in <i>S. pneumoniae</i> (33, 41, 42).
413	Six additional proteins differentially expressed after NO treatment play a role in
414	pneumococcal metabolism. Arginine deiminase (ArcA) and two alcohol dehydrogenases
415	(Adh) were notably decreased. In our previous study, expression of these proteins was
416	markedly increased during biofilm formation, and may compensate for the dramatic
417	reduction in glycolytic activity observed in <i>S. pneumoniae</i> biofilms by up-regulating
418	arginine and pyruvate metabolism (33). Similar to the differential expression of
419	translational proteins, the increased expression of metabolic proteins following NO
420	treatment suggested that pneumococcus differentially modulates metabolism in planktonic
421	and biofilm modes of growth. Qualitatively, seven other metabolic proteins exhibited
422	differential expression following NO treatment, compared with untreated biofilms.
423	However these proteins were below the threshold of >3 peptides required for inclusion in
424	the quantitative iTRAQ dataset. Nonetheless, taken together these data suggest that while
425	high concentrations of NO elicit a direct antibacterial effect, low dose NO may be involved
426	in regulation of metabolism via a currently unknown signaling pathway.
427	The decreased expression of arginine deiminase following NO treatment is of
428	particular interest. Regulation of <i>S. pneumoniae</i> arginine metabolism appears to be distinct
429	from other bacteria and involves the regulators ArgR1, ArgR2 and AhrC (43, 44). ArgR1 is a
430	transcriptional regulator of the arginine deminase system (ADS) consisting of arginine
431	deiminase (<i>arcA</i>), ornithine carbamoyltransferase (<i>arcB</i>) and carbamate kinase (<i>arcC</i>) that
432	mediate arginine acquisition and virulence in pneumococcus. Abrupt changes in arginine
433	concentrations were recently shown to induce differential transcription of >450 genes in

434 *Streptococcus gordonii*, many of which were involved in adhesion and biofilm development
435 (45). Furthermore, the difference in the effect of NO on two of the strains of pneumococcus
436 used in our experiments is consistent with D39 and Serotype 14 having variable
437 disruptions in arginine regulators ArgR1 and ArgR2 (43, 44).

438 We speculate that NO may play a novel role in arginine metabolism and biofilm 439 development in *S. pneumoniae*, and in regulating growth in pneumococcus. Our results 440 suggest that low concentrations of NO modulate pneumococcal growth, possibly making 441 dormant bacteria within the biofilm metabolically active and more susceptible to antibiotic 442 killing. However, an alternative explanation is that the production of OONO⁻ may contribute 443 to the antibacterial effects of NO on pneumococcus since the antibacterial effect was 444 reduced in the presence of the peroxynitrite (OONO⁻) scavenger L-methionine. Elevated 445 OONO⁻ levels have also been found to be associated with the dispersal response and cell 446 death in *P. aeruginosa* (27). Peroxynitrite is a potent pro-oxidant and cytotoxic species 447 produced by the interaction of superoxide (O_2) and NO. Compromised pneumococcal 448 superoxide dismutase (SOD) activity may lead to enhanced formation of O₂- and 449 subsequent reaction with host NO leading to the production of OONO⁻, a reaction that 450 normally takes place in human macrophages (46). Peroxynitrite was a putative mediator of 451 NO induced cytotoxic damage in pneumococcal infected microglial cell cultures in vitro and 452 *in vivo* using pneumococcal mutants for pyruvate oxidase (*spxB*), and the arginine 453 metabolism mutant *carB* in mice (22). Intriguingly, these authors made the novel 454 observation that pneumococcus can release NO, suggesting that NO is an endogenous

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455	pneumococcal metabolite. Our previous proteomic analyses indicated pyruvate oxidase,
456	which produces H_2O_2 , was also markedly upregulated in pneumococcal biofilms (33).
457	A MerR-like transcriptional factor <i>NmlR_{sp}</i> required for NO defense was identified in
458	S. pnuemoniae D39 using the NO donor S-nitrosoglutathione (GSNO) (47). There was no
459	evidence of a MER-like transcriptional protein in our proteomic data, however the role of
460	$\mbox{Nm}\mbox{R}_{\mbox{sp}}$ was subsequently noted to have broader functional roles including a role in $\mbox{H}_2\mbox{O}_2$
461	production and in arginine biosynthesis (48).
462	The NO donor SNP has been widely utilized for a number of clinical applications,
463	primarily through its use as a vasodilator, however, prolonged treatment and/or high
464	doses have been suggested to pose a risk of cyanide-mediated cytotoxicity (49, 50). The
465	decomposition of SNP to cyanide has also been shown to be slow (<2.5% over 72 h) when
466	protected from direct exposure to high intensity/natural light (51). For the purpose of our
467	initial study SNP was used as a suitable NO-donor to explore the actions of NO on
468	pneumococcal biofilms since relatively low doses were applied for a short period (26, 27).
469	Furthermore, treatment with equimolar concentrations of KCN had no effect on
470	pneumococcal growth indicating that any observed responses to SNP were not the result of
471	cyanide toxicity (52). However, future studies investigating NO-mediated anti-
472	pneumococcal effects with alternative donors, such as Cephalosporin-3 $'$ -diazeniumdiolate
473	NO-Donor Prodrugs, which have been specifically designed to release NO at sites of
474	bacterial infection may offer better choices for clinical use (53).
475	The results of our study are consistent with other studies showing high
476	concentrations of NO were toxic to bacteria including pneumococcus (32). However, to our
477	knowledge our study is the first to show that: 1) planktonic and biofilm <i>S. pneumoniae</i>

478 responded differentially to low and high concentrations of NO; 2) the anti-pneumococcal 479 response was not induced by nitrite or nitrate, but was NO specific; 3) unlike other 480 bacteria, low concentrations of NO did not elicit a dispersal response by biofilm S. 481 pneumoniae; 4) a low concentration of NO altered the protein expression profile of biofilm 482 pneumococci; and 5) when accompanied by adjunctive treatment with NO, pneumococcal 483 sensitivity to antibiotic treatment was enhanced in vitro and ex vivo. These results suggest 484 that at lower concentrations, NO perturbs pneumococcal biofilm metabolism, but at higher 485 concentrations NO is toxic to S. pneumoniae. Targeted adjunctive NO treatment may be a 486 candidate novel therapy for reducing biofilm tolerance by pneumococcus. 487

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494

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662	Figure 1: SNP treatment of <i>in vitro</i> pneumococcal biofilms reduced biofilm viability
663	and biomass. 48h S. pneumoniae serotype 14 in vitro biofilms were treated with SNP for 2
664	hours, and the biomass assessed by absorbance (OD600) and viability measured by CFU/
665	cm ² . A significant reduction in total biomass and the number of viable cells remaining
666	within the biofilm was observed following 1 mM SNP treatment. *** $p \le 0.001$.
667	
668	Figure 2: SNP treatment of <i>in vitro S. pneumoniae</i> biofilms reduced the viable cell
669	population in the surrounding supernatant. The viability of 48h S. pneumoniae serotype
670	14 in vitro biofilm and supernatant populations was measured by CFU enumeration
671	following treatment with 1 mM SNP. SNP treatment significantly reduced both the biofilm
672	and supernatant populations. * $p \le 0.05$.
673	
674	Figure 3: In vitro S. pneumoniae biofilms treated with SNP demonstrated reduced
675	viability and no evidence of dispersal. 48h S. pneumoniae serotype 14 in vitro biofilms
676	were treated with 1 mM SNP for 2 hours then imaged using confocal microscopy and
677	Live/Dead staining. 1 mM SNP-treated biofilms (b) demonstrated no obvious change in
678	biomass when compared with untreated biofilms (a) , however, a reduction in the number
679	of Syto9-stained live bacteria in the 1 mM SNP-treated biofilms (d) was reduced in
680	comparison with untreated biofilms (c) , commensurate with CFU enumeration data.
681	Scanning electron microscopy with Alcian Blue staining further demonstrated no obvious
682	changes in biofilm ultrastructure between untreated (e) and 1 mM SNP-treated (f) biofilms
683	(4,000x magnification; scale bar: 10 μm).
684	

685 Figure 4: SNP treatment reduced the in vitro S. pneumoniae planktonic growth rate. S. 686 *pneumoniae* serotype 14 *in vitro* planktonic cultures were treated with SNP during 687 exponential growth phase, and the growth rate was measured by the change in absorbance 688 (OD595) over 2 hours, and compared with the untreated growth rate. A significant 689 reduction in growth rate was observed using 500 µM SNP, and complete cessation of 690 growth was observed with concentrations greater than 5 mM. $p \leq 0.05$; **** $p \leq 0.0001$. 691 692 Figure 5: The response of S. pneumoniae to treatment with SNP was NO-mediated. a) 693 S. pneumoniae serotype 14 exponential planktonic cultures were treated with the nitric 694 oxide (NO) donors SNP, DEA/NO, nitrate and nitrite, and the CN⁻ anion control potassium 695 cyanide (KCN) over 2 hours. Significant decreases in the growth rate were observed upon 696 treatment with two independent NO donors, SNP and DEA/NO, indicating that the response 697 was NO-mediated. KCN treatment had no effect on growth rate confirming the response to 698 SNP was not CN⁻ mediated (p=0.528). Sodium nitrate (p=0.321) and sodium nitrite 699 (p=0.078) treatments also had no effect on growth rate suggesting that nitrate and nitrite, 700 respectively, were not utilised as sources of NO. Finally, the addition of b) the NO-701 scavenger carboxy-PTIO, and **c)** the peroxynitrite scavenger L-methionine reduced the 702 response to SNP treatment suggesting the response may be mediated by either NO or 703 peroxynitrite. *≤0.05; **≤0.01; ***≤0.001. 704 705 Figure 6: Adjunctive treatment of S. pneumoniae in vitro biofilms with SNP enhanced

antibiotic efficacy. 48h *S. pneumoniae* serotype 14 (ST124), 19F, 23F and D39 *in vitro*

507 biofilms were treated for 2 hours and the remaining viable cells measured by CFU

709

710 significant reduction in viability. * $p \le 0.05$. 711 712 Figure 7: Adjunctive treatment of S. pneumoniae biofilms on ex vivo adenoid tissue 713 with SNP enhanced antibiotic efficacy. Adenoid tissue samples (n=11) were dissected 714 into four equal sections (each with a similar proportion of luminal surface) treated for 2 715 hours, and the viability of *S. pneumoniae* was measured by CFU enumeration. SNP 716 treatment alone had no significant effect on viable pneumococci (p=0.722), whereas AMC 717 treatment alone resulted in a significant reduction (p=0.005). Combined SNP and AMC 718 treatment however, resulted in enhanced antibiotic efficacy (p=0.041). * $p\leq0.05$ (Wilcoxon 719 Signed Ranks test). 720 721 Figure 8: Treatment of S. pneumoniae in vitro biofilms with SNP resulted in the 722 differential expression of a small subset of quantitatively identified proteins. 723 Comparative iTRAQ analyses of SNP treated (100 IM SNP/2 hours) and untreated S. 724 pneumoniae serotype 147-day old in vitro biofilms quantitatively identified 112 proteins of 725 which 13 were differentially expressed following treatment. 726 727 Figure 9: Treatment of S. pneumoniae in vitro biofilms with SNP resulted in a change 728 in metabolic and translation protein expression levels. Comparative iTRAQ analyses of 729 SNP treated (100 μ M SNP/2 hours) and untreated *S. pneumoniae* serotype 14 *in vitro* 730 biofilms a) quantitatively identified 13 differentially expressed proteins, and b)

enumeration. When used separately both SNP and AMC treatment reduced the viable

biofilm cell population, however, combined SNP and AMC treatment resulted in a further

- 731 qualitatively identified 12 differentially expressed proteins following treatment.
- 732 Quantitative inclusion criteria: ≥3 peptide matches, ≥50 protein score, ≥5% sequence
- 733 coverage (p<0.05). Qualitative inclusion criteria: 2 peptide matches, ≥50 protein score,
- 734 \geq 5% sequence coverage (p<0.05). Comparative protein data with >1.3 and <0.77 ratios
- 735 identified as having differential protein expression.



SNP Concentration

AAC





AAC



Sec. 16.6

a

b

20 z (µm)

d

730 20 z (µm) 10

180 200 220 240

200 220 240

180

160

140

120

x (µm)

100

120 x (µm) 140

Antimicrobial Agents and Chemotherapy

AAC



SNP Concentration





AAC







Antimicrobial Agents and Chemotherapy

AAC



Increased expressionDecreased expressionNo change

а								
					Expression Ratio			% Sequence
Function	Gene	Protein	Accession No.	(NO-treated/Untreated)			Matches	Coverage
	rpsB	30S ribosomal protein S2	YP_001836898	1	1.38		4	37.5
	rplQ	50S ribosomal protein L17	YP_001834961	1	1.46		5	33.6
Translation	rplM	50S ribosomal protein L13	YP_001835025	1	1.36		3	35.8
	rpIN	50S ribosomal protein L14	YP_001834945	1	1.87		4	50
	rpsO	30S ribosomal protein S15	YP_001836318	1	1.32		3	29.2
	arcA	arginine deiminase	YP_001836835		0.68	€	8	37.7
	fabG	3-ketoacyl-(acyl-carrier-protein) reductase	YP_001835136	1	1.49		4	28.4
Metabolism	adhE	alcohol dehydrogenase, iron-containing	YP_001836708	31	0.73	*	6	9.9
	adh	alcohol dehydrogenase, zinc-containing	YP_001836739		0.73	*	3	9.9
	manN	PTS system, mannose-specific IID component	YP_001835010		0.76	↓	6	23.1
	fruA	PTS system, fructose specific IIABC components	YP_001835543	1	1.32		4	13.8
Other	SPCG_2124	SPFH domain-containing protein	YP_001836841	1	1.80		3	18.1
	SPCG_1532	hypothetical protein SPCG_1532	YP_001836249	1	1.31		5	26.8

b

Antimicrobial Agents and Chemotherapy

AAC

				Expression Ratio			Peptide	% Sequence
Function	Gene	Protein	Accession No.	(NO-treated/Untreated)		Matches	Coverage	
Translation	rpsF	30S ribosomal protein S6	YP_001836244	1	1.32		2	26
	glmS	D-fructose-6-phosphate amidotransferase	YP_001834993		0.77	+	2	9.8
	tktA	transketolase	YP_001836712		0.75	*	2	6.5
	acoL	dihydrolipoyl deyhdrogenase	YP_001835853	1	1.37		2	14.3
Metabolism	dapH	2,3,4,5-tetrahydropyridine-2-carboxylate N-succinyltransferase	YP_001836779	1	1.46		2	13.8
	atpF	ATP synthase subunit B	YP_001836213		0.75	+	2	7.9
	accD	acetyl-CoA carboxylase beta subunit	YP_001835141		0.76	*	2	14.6
	metG	methioninetRNA ligase	YP_001835454	1	1.64		2	7.1
	SPCG_1897	hypothetical protein SPCG_1897	YP_001836614	1	1.34		2	11.8
Other	amiE	oligopeptide ABC transporter, ATP-binding protein	YP_001836579	1	1.64		2	10.1
	gidA	glucose-inhibited division protein A	YP_001834840		0.66	*	2	8
	SPCG_1659	Gfo/Idh/MocA family oxidoreductase	YP_001836376		0.67	*	2	7.6