“On demand” redox buffering by H₂S contributes to antibiotic resistance revealed by a bacteria-specific H₂S donor†

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Understanding the mechanisms of antimicrobial resistance (AMR) will help launch a counter-offensive against human pathogens that threaten our ability to effectively treat common infections. Herein, we report bis(4-nitrobenzyl)sulfanes, which are activated by a bacterial enzyme to produce hydrogen sulfide (H₂S) gas. We found that H₂S helps maintain redox homeostasis and protects bacteria against antibiotic-triggered oxidative stress “on demand”, through activation of alternate respiratory oxidases and cellular antioxidants. We discovered, a hitherto unknown role for this gas, that chemical inhibition of H₂S biosynthesis reversed antibiotic resistance in multidrug-resistant (MDR) uropathogenic Escherichia coli strains of clinical origin, whereas exposure to the H₂S donor restored drug tolerance. Together, our study provides a greater insight into the dynamic defence mechanisms of this gas, modes of antibiotic action as well as resistance while progressing towards new pharmacological targets to address AMR.

Introduction

Maintenance of redox homeostasis is fundamental to cellular growth and survival. Induction of dysfunctional redox environment is a common mechanism used against pathogens by immune cells. In the past several decades, it has been well established that gases such as hydrogen sulfide (H₂S) and nitric oxide (NO) affect cellular redox balance. Bacteria-derived-H₂S through microbiota contribute significantly to repair mechanisms and are vital for the health of the gastrointestinal tract. Bacterial H₂S is also implicated as a cytoprotective agent against antibiotic-induced stress, thereby enhancing antibiotic tolerance. Oxidant remediation by bacterial H₂S is operational, but precise mechanisms of protection remain to be completely elucidated. Mapping out these cytoprotective mechanisms will help progress towards new strategies to combat the growing threat of antimicrobial resistance (AMR). Due to the dwindling arsenal of antibiotics, AMR is possibly the biggest problem that this current generation will face. In order to address this complex socioeconomic public health problem, multiple methodologies are necessary including a better understanding of the mechanisms of antibiotic action and factors contributing to antibiotic resistance. Herein, we systematically investigated the dynamic effects of H₂S in protecting bacteria from antibiotic-induced stress and the role of H₂S in modulating AMR.

Being a gaseous species, reliable detection as well as controlled and site-specific generation of H₂S within cells is fundamental to understanding its biology. Numerous donors of H₂S (ref. 3) are in development but none, to our knowledge, distinguish one type of cells over others. Enzymes, as metabolic triggers for activation of donors, offer distinct advantages as they facilitate localization of H₂S. A H₂S generating functional group is tethered to a substrate for an enzyme that is normally expressed in cells of interest (Fig. 1a). Upon entry into cells, metabolism by the target enzyme frees up the active H₂S generator inside cells thus achieving localized delivery. Recently, two esterase-activated H₂S donors were reported with wide potential applications in cellular studies (Fig. 1b). However, generating H₂S specifically in certain cells over others might be problematic when using esterase as a trigger. We chose E. coli nitroreductase (NTR), an oxygen-insensitive bacterial enzyme that is frequently expressed in bacteria but not in mammalian cells. Geminal dithiols are reported to undergo hydrolysis in buffer to produce...
H$_2$S; 4-nitroaryl groups are known substrates for NTR. We hence designed 1, an NTR-activated H$_2$S donor (Fig. 1b).

Results and discussion

Synthesis of 1 is achieved by the reaction of a variety of ketones (2) with 4-nitrobenzyl thiol (3) (Table 1). A H$_2$S-sensitive dye BODIPY-azide 4a was employed to detect H$_2$S. BODIPY-azide is known to be reduced by H$_2$S to produce a fluorescent amine 4b.

Compounds 1a–1g were independently exposed to NTR and all compounds were found to generate H$_2$S under these conditions (Fig. 2a). The cyclopentyl derivative 1c was found to be slightly better than the cohort of donors tested, and this compound was used for further studies.

A monobromobimane (mBBr, 6a) assay (with some modifications) was next used to confirm the production of H$_2$S. The electrophile mBBr reacts with sulfide anion to produce a thioether, which contains two bimane units (6b). When 6a was treated with Na$_2$S in a pH 7.4 buffer, as expected, 6b was formed (Fig. 2b). Under similar conditions, when 1c was co-incubated with 6a and NTR, we found evidence for the formation of 6b again supporting 1c as a source of H$_2$S when incubated with NTR (Fig. 2b). Next, NBD-fluorescein, a H$_2$S sensitive dye, was synthesized and incubated in the presence of 1c and NTR. Again, we found a distinct increase in fluorescence attributable to H$_2$S generation (Fig. S1, ESI†). Thus, the formation of H$_2$S was validated by several independent assays suggesting that this compound is a reliable donor of H$_2$S.

$$\text{H}_2\text{S}$$

The H$_2$S donor 1c was able to maintain elevated levels of H$_2$S over 45 min (Fig. 2c). The formation of a hydroxylamino- or amino-aryl derivative (Fig. 1c, Scheme S2, ESI†), which self-immolates to generate a geminal dithiol, was likely. This geminal dithiol should hydrolyze to produce H$_2$S.

Accordingly, when 1e was incubated in buffer in the presence of

Table 1

<table>
<thead>
<tr>
<th>Entry</th>
<th>R$^1$</th>
<th>R$^2$</th>
<th>Ketone</th>
<th>Prod</th>
<th>Yield %</th>
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<tbody>
<tr>
<td>1</td>
<td>CH$_3$</td>
<td>CH$_3$</td>
<td>2a</td>
<td>1a</td>
<td>80</td>
</tr>
<tr>
<td>2</td>
<td>Et</td>
<td>Et</td>
<td>2b</td>
<td>1b</td>
<td>78</td>
</tr>
<tr>
<td>3</td>
<td>Cyclopentyl</td>
<td></td>
<td>2c</td>
<td>1c</td>
<td>70</td>
</tr>
<tr>
<td>4</td>
<td>Cyclohexyl</td>
<td></td>
<td>2d</td>
<td>1d</td>
<td>71</td>
</tr>
<tr>
<td>5</td>
<td>Ph</td>
<td>CH$_3$</td>
<td>2e</td>
<td>1e</td>
<td>60</td>
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<tr>
<td>6</td>
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<td>CH$_3$</td>
<td>2f</td>
<td>1f</td>
<td>42</td>
</tr>
<tr>
<td>7</td>
<td>Thiophene</td>
<td>CH$_3$</td>
<td>2g</td>
<td>1g</td>
<td>38</td>
</tr>
</tbody>
</table>
Zn and ammonium formate, acetophenone was formed, supporting the proposed mechanism (Fig. S2, ESI†).

Having established that 1c generated H₂S in cell-free conditions in the presence of a bacterial enzyme, the ability of this compound to permeate cells to be metabolized by NTR to generate H₂S was evaluated. First, an HPLC-based method was used: E. coli cells were incubated with the H₂S-sensitive dye 4a and 1c. Cells were lysed and HPLC analysis of the lysate revealed the formation of 4b supporting H₂S generation (Fig. S3, ESI†); a similar result was recorded for a variety of bacteria supporting the broad utility of this donor. Next, flow cytometry analysis revealed the generation of H₂S inside intact bacterial cells when treated with 1c, supporting the ability of this donor to enhance H₂S levels in live bacterial cells (Fig. 2d). Next, 4-nitrobenzylbenzoate 5 (a likely competitive inhibitor) was synthesized using a previously reported method.† This compound was by itself incapable of generating H₂S in the presence of NTR and also inhibited H₂S generation from 1c (Fig. 2a). The negative control 5 did not generate H₂S within the bacteria, suggesting that the metabolism of the nitro group does not contribute to H₂S production (Fig. 2d). The H₂S donor 1c was ineffective in generating H₂S in E. coli strains lacking NTR (Fig. S4, ESI†), confirming NTR-specificity in vivo. As NTR is predominantly produced in bacteria but not in mammalian cells, the H₂S donor 1c must selectively enhance H₂S in bacteria. Human monocytic cells (THP-1) were treated with 1c, and H₂S levels were assessed by flow cytometry. Herein, we find that while Na₂S was capable of enhancing H₂S levels within THP-1 cells, 1c remained completely ineffective (Fig. 2e). Thus, 1c was selective in its ability to enhance H₂S in bacteria over mammalian cells (Fig. S5, ESI†). To our knowledge, this is the first example of a H₂S donor with species selectivity. Thus, this study lays the foundation for novel methodologies for site-specific enhancement of H₂S using this class of H₂S donors, for example, selectively enhancing H₂S in microbiota to study the effects of this gas on colorectal cancer and other similar pathophysiologicals is possible.\(^2\)

To begin understanding the mechanisms of H₂S-mediated oxidation remediation, we used a non-invasive redox biosensor (roGFP2) and assessed dynamic changes in the cytoplasmic redox potential of E. coli in response to oxidative stress.\(^18\) An increase in 405/488 nm excitation ratio of roGFP2 indicates oxidative stress, while the reverse suggests reductive changes.\(^19\) We first exposed E. coli expressing roGFP2 to 1c and measured the roGFP2 biosensor response, and no significant changes in 405/488 ratio were observed (Fig. 3a). Hence, H₂S alone did not affect ambient redox-potential of E. coli. In contrast, oxidative challenge with H₂O₂, a reactive oxygen species (ROS), rapidly increased the 405/488 ratio, and pre-treatment with 1c significantly reversed this response (Fig. 3a). However, pre-treatment with either Na₂S or 5 had no influence on H₂O₂-induced oxidative changes in the biosensor response (Fig. S6, ESI†).

Importantly, protective influence of 1c on intrabacterial redox potential translated into significantly higher resistance displayed by E. coli against bactericidal concentrations of H₂O₂ (Fig. 3b). Interestingly, H₂S itself did not have any significant effect on the growth of E. coli. Thus, intervention by H₂S occurs when other endogenous oxidant-remediation systems are overwhelmed. This property is consistent with the lower reduction potential of H₂S, when compared with major cellular thiols such as glutathione,\(^19\) and thus affords H₂S a unique role in cellular redox chemistry. Furthermore, in contrast with other routinely used antioxidants in redox biology, such as thiourea and bipyridyl, 1c does not significantly affect bacterial growth (Fig. 3b), suggesting that this tool would be appropriate for studying H₂S-mediated response to dynamic redox alterations during antibiotic-induced stress and lethality.

The emerging model for antibiotic lethality involves the induction of complex redox and metabolic alterations as a consequence of drugs’ interaction with their specific targets.\(^2,20\)

Fig. 3 (a) Reduction–oxidation sensitive GFP (roGFP2) was used to measure dynamic changes in cytoplasmic redox potential of E. coli upon exposure to: H₂O₂, 1 mM; 1c, 100 μM; (b) time-kill analysis of E. coli treated with hydrogen peroxide (1 mM) and 1c (100 μM) during 40 min. (c) Dynamic changes in cytoplasmic redox potential of E. coli upon exposure to: Amp, 5 μg mL⁻¹; 1c, 100 μM; (d) time-kill analysis of E. coli treated with Amp (5 μg mL⁻¹) and 1c (100 μM) during 120 min. (e) Dynamic changes in cytoplasmic redox potential of E. coli upon exposure to: Amik, 20 μg mL⁻¹; 1c, 100 μM; (f) time-kill analysis of E. coli treated with Amik (20 μg mL⁻¹) and 1c (100 μM) during 120 min.
Thus, it is important to understand the dynamic effects of H$_2$S in mitigating antibiotic-induced redox stress.\textsuperscript{24} To accomplish this, we exposed \textit{E. coli} to clinically relevant concentrations of the bactericidal antibiotic ampicillin (Amp; cell wall targeting), and an oxidative shift was recorded (Fig. 3c).\textsuperscript{21a,22} More importantly, pre-treatment with 1c reduced the degree of oxidative shift induced by Amp, resulting in significant tolerance to antibiotics (Fig. 3d). Amp-mediated increase in roGFP2 ratios emerged earlier than the time points at which significant killing was observed, indicating that oxidative stress precedes death, and 1c-derived H$_2$S protects bacteria by maintaining cytoplasmic redox potential.

Similar results were recorded for amikacin (Fig. 3e–f), a translation inhibitor, and ciprofloxacin, a replication inhibitor (Fig. S7, ESI†). Altogether, these results demonstrate that elevating endogenous H$_2$S levels can arrest antibiotics-triggered redox stress and killing.

Mechanisms of H$_2$S-mediated protection from antibiotic-induced lethality are poorly understood. It has been shown that H$_2$S elevates cellular antioxidant capacity and suppresses iron load in order to mitigate antibiotic-linked ROS production.\textsuperscript{23} Since sulfide is a potent ligand of copper and heme moieties, H$_2$S efficiently inhibits aerobic respiration by targeting copper-heme containing cytochrome bo oxidase (CyoA).\textsuperscript{24} Under these conditions, respiration becomes primarily dependent upon less energy-efficient cytochrome bd oxidase (CydB).\textsuperscript{24} Interestingly, modulation of cytochrome oxidases expression is known to influence antibiotic toxicity.\textsuperscript{21b} Therefore, we assessed whether terminal oxidases are important contributory factors in H$_2$S-mediated antibiotic tolerance. First, quantitative reverse transcription-PCR (qRT-PCR) analysis of \textit{E. coli} cells in the absence or presence of 1c was conducted (see ESI†). A significant down-regulation of the genes encoding CyoA was observed with 1c-treated bacteria (Fig. 4a). The expression of alternate oxidases was however either maintained (cytochrome bd oxidase I (cydB)) or highly induced (cytochrome bd oxidase II (appY)) by H$_2$S (Fig. 4a). During growth under low-O$_2$ tension, \textit{E. coli} down-regulated cyo operon and upregulated cyd and app operons, indicating that H$_2$S triggered genetic and physiological changes comparable to O$_2$-limitation.\textsuperscript{25} Amp treatment reversed the influence of H$_2$S on the expression of cytochrome oxidases as cyoA and cydB transcripts were significantly induced and repressed, respectively, compared to untreated cells (Fig. 4a). The \textit{appY} transcript remained down-regulated in response to Amp. Data suggest that Amp treatment promotes respiration via energetically efficient CyoA, which is consistent with a recent study demonstrating acceleration in aerobic respiration in response to bactericidal antibiotics.\textsuperscript{21b}

Having observed divergent effects of H$_2$S and Amp on cytochrome oxidases gene expression, we next examined the outcome of H$_2$S and Amp combination on transcription. qRT-PCR analysis of \textit{E. coli} pre-treated with 1c, followed by exposure to Amp showed severely down-regulated expression cyoA, whereas expression of cydB and \textit{appY} was robustly maintained (Fig. 4a) compared to that of Amp alone. Thus, maintenance of a respiratory flux through cytochrome bd oxidase I/II in response to H$_2$S treatment may be a key trait that permits adaptation upon subsequent exposure to antibiotics. To examine this possibility, we assessed cell-killing in respiratory mutants lacking either \textit{cyoA} or \textit{cydB}. While 1c-pretreatment resulted in a significant attenuation of Amp lethality in the case of CyoA mutant (like WT strain), it was completely ineffective in protecting \textit{cydB} mutant (Fig. 4b). In LB medium, WT, cyoA, and \textit{cydB} strains showed comparable growth profiles in the absence or presence of 1c, indicating that the differences in Amp susceptibility are not a consequence of reduced growth rates. Sustenance of \textit{cydB} expression in response to H$_2$S-Amp combination, coupled with maintenance of H$_2$S-mediated antibiotic tolerance in \textit{cyo} mutant (where aerobic respiration is mainly carried out by CydB) but not in \textit{cydB} mutant, suggests that the H$_2$S effect is likely to be dependent upon \textit{cydB}.\textsuperscript{24} Along with its role in respiration, CydB from \textit{E. coli} has been shown to reduce H$_2$O$_2$ by displaying catalase and quinol peroxidase activities.\textsuperscript{26} Therefore, maintenance of \textit{cydB} expression by H$_2$S can potentiate antibiotic tolerance by bolstering the bacterial antioxidant capacity. In concurrence with this result, displayed heightened sensitivity to H$_2$O$_2$ compared to \textit{cyoA} and WT strains (Fig. S8, ESI†).

In addition, consistent with previous results,\textsuperscript{4d} we observed that H$_2$S is ineffective in protecting cells that lack cytoplasmic catalase and peroxidase (KatA, KatE, ahpCF; HPX-) from Amp-induced lethality (Fig. S9, ESI†). Altogether, data implicate a central role for cytochrome bd oxidase and oxidant-
and co-workers developed a CBS inhibitor that helps to prevent the urgent need to develop new therapeutics. Recently, Berkowitz and co-workers developed a CBS inhibitor that helps to prevent the deleterious effects of enhanced H$_2$S such as neuronal cell death during episodes of stroke.\textsuperscript{29} The inhibitors that were developed in their study showed a marked diminution in neuronal cell death compared with an untreated control. It is likely that inhibitors of 3-MST may find similar application in sensitizing resistant pathogens.\textsuperscript{30} Our results provide a sound pharmacological basis for the design of inhibitors of biosynthesis of H$_2$S as a possible adjuvant.\textsuperscript{29–31} Furthermore, we identified critical aspects of bacterial physiology that could be exploited as part of new potentiation strategies. For examples, targeting antioxidant enzymes and alternate respiratory complexes (Cyd/App) is likely to enhance the killing potential of antibiotics. A combination of molecules/drugs targeting H$_2$S, antioxidants, and respiration could have a remarkable impact on drug-resistance and clinical outcomes.

\section*{Conclusions}

In summary, we report a new H$_2$S donor that reliably and selectively enhances H$_2$S within bacteria. An application of our new tool clearly revealed that H$_2$S is a key player in the maintenance of intracellular redox balance of bacteria to counteract a lethal degree of oxidative stress induced by antibiotics. The critical role that H$_2$S played in modulating drug resistance is also shown. Antibiotic resistance is emerging as possibly the biggest global health challenge for this generation. Therefore understanding pathogen defence mechanisms and their consequences in drug resistance is critical. From the evolutionary perspective, H$_2$S generating enzymes are prevalent in most sequenced bacterial genomes including environmental bacteria, indicating a naturally conserved role of H$_2$S in ensuring survival. It is likely that H$_2$S producing capability is under the selective pressure in diverse environmental bacteria that arises due to antimicrobials secreted by other bacteria and fungi inhabiting the same niche. Our study presents significant advances towards a complete understanding of antibiotic-induced stress and cytoprotective mechanisms of H$_2$S.

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\section*{Notes and references}
