

# Overlapping and Complementary Oxidative Stress Defense Mechanisms in Nontypeable *Haemophilus influenzae*

Alistair Harrison, Beth D. Baker, Robert S. Munson, Jr.

The Center for Microbial Pathogenesis at the Research Institute at Nationwide Children's Hospital and the Center for Microbial Interface Biology and Department of Pediatrics, The Ohio State University, Columbus, Ohio, USA

**The Gram-negative commensal bacterium nontypeable *Haemophilus influenzae* (NTHI) can cause respiratory tract diseases that include otitis media, sinusitis, exacerbations of chronic obstructive pulmonary disease, and bronchitis. During colonization and infection, NTHI withstands oxidative stress generated by reactive oxygen species produced endogenously, by the host, and by other copathogens and flora. These reactive oxygen species include superoxide, hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>), and hydroxyl radicals, whose killing is amplified by iron via the Fenton reaction. We previously identified genes that encode proteins with putative roles in protection of the NTHI isolate strain 86-028NP against oxidative stress. These include catalase (HktE), peroxiredoxin/glutaredoxin (PgdX), and a ferritin-like protein (Dps). Strains were generated with mutations in *hktE*, *pgdX*, and *dps*. The *hktE* mutant and a *pgdX hktE* double mutant were more sensitive than the parent to killing by H<sub>2</sub>O<sub>2</sub>. Conversely, the *pgdX* mutant was more resistant to H<sub>2</sub>O<sub>2</sub> due to increased catalase activity. Supporting the role of killing via the Fenton reaction, binding of iron by Dps significantly mitigated the effect of H<sub>2</sub>O<sub>2</sub>-mediated killing. NTHI thus utilizes several effectors to resist oxidative stress, and regulation of free iron is critical to this protection. These mechanisms will be important for successful colonization and infection by this opportunistic human pathogen.**

During aerobic respiration, the suboptimal metabolic reduction of molecular oxygen to water generates reactive oxygen species that are profoundly toxic to bacterial cells. Bacteria have therefore evolved multiple mechanisms to combat such lethal insults. These include enzymatic scavenging of oxidative stress-inducing agents by superoxide dismutases (SODs) that breakdown superoxide, as well as catalases, alkylhydroperoxides, periplasmic thiol peroxidases, and bacterioferritin-comigratory proteins that decompose peroxides. In addition to a bacterium's need to protect itself from endogenously generated reactive oxygen species, bacteria that reside within a host must also contend with the release of extracellular reactive oxygen species (ROS) by phagocytes, copathogens, and the host's flora. The mechanisms that abrogate the effects of externally derived oxidative stress within a host have been well described in the model organism *Escherichia coli* (1). These defenses overlap the defenses used to combat endogenously generated oxidative stress.

Bacteria can also use nonenzymatic mechanisms to protect against oxidative stress. Protection against the effects of free iron is especially critical. Hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) can react with intracellular free iron to generate toxic hydroxyl radicals via the Fenton reaction. The hydroxyl radicals generated can then damage DNA. For example, when an *E. coli* strain unable to scavenge H<sub>2</sub>O<sub>2</sub> was repeatedly passaged under aerobic conditions, DNA damage mediated by H<sub>2</sub>O<sub>2</sub> was readily observed (2, 3). As iron is critical for the generation of hydroxyl radicals via the Fenton reaction, bacteria protect themselves through the sequestration of iron. The role of ferritins and ferritin-like proteins, such as Dps, in this iron sequestration process is well understood (3–5). For example, Dps protects *E. coli* from the effects of H<sub>2</sub>O<sub>2</sub> in stationary phase (6). Moreover, there is strong conservation among Dps orthologs, with several core conserved amino acids forming an iron-binding core. Within this core, the bound ferrous iron is oxidized by H<sub>2</sub>O<sub>2</sub>, but critically, hydroxyl radicals are not produced (7). The expression of Dps is regulated by the anoxic redox control

regulator ArcA and protects anaerobically grown *Haemophilus influenzae* strain Rd from H<sub>2</sub>O<sub>2</sub> (8). Dps has also been shown to have a role in biofilm formation and *in vivo* survival of the nontypeable *H. influenzae* (NTHI) strain 86-028NP (9).

NTHI is a major cause of otitis media (OM), sinusitis, exacerbations of chronic obstructive pulmonary disease, and bronchitis, as well as other infections (10). Thus, illness due to NTHI is a major societal burden. NTHI resides as a commensal in the nasopharynx, along with host flora. In certain instances, for example following a viral infection, NTHI can then move to the middle ear, where it causes OM. Infection in the middle ear brings an increase in polymorphonuclear leukocytes (PMNs), with an increased possibility of insult by ROS. In addition, *Streptococcus pneumoniae*, a species that can cocolonize the nasopharynx, produces extracellular H<sub>2</sub>O<sub>2</sub> at concentrations that are bactericidal toward *H. influenzae in vitro*. *H. influenzae* therefore has a number of proteins that protect against oxidative stress. The genes that encode catalase and PgdX were previously identified in *H. influenzae* strain Rd. Subsequent analyses of the genome of the NTHI strain 86-028NP identified homologues of genes encoding catalase, PgdX, Dps, and the alkylhydroperoxidase TsaA (11, 12).

We hypothesized that expression of the numerous oxidative

Received 12 June 2014 Accepted 27 October 2014

Accepted manuscript posted online 3 November 2014

Citation Harrison A, Baker BD, Munson RS, Jr. 2015. Overlapping and complementary oxidative stress defense mechanisms in nontypeable *Haemophilus influenzae*. *J Bacteriol* 197:277–285. doi:10.1128/JB.01973-14.

Editor: V. J. DiRita

Address correspondence to Alistair Harrison, alistair.harrison@nationwidechildrens.org.

Copyright © 2015, American Society for Microbiology. All Rights Reserved. doi:10.1128/JB.01973-14

TABLE 1 Bacterial strains

Strain	Description	Reference or source
86-028NP	Nontypeable <i>H. influenzae</i> strain from a child with chronic otitis media	12
86-028NP <i>rpsL</i>	Streptomycin-resistant derivative of 86-028NP	18
86-028NP <i>rpsL</i> $\Delta$ <i>tsaA</i>	Derivative of 86-028NP <i>rpsL</i> with a deletion mutation of <i>tsaA</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i>	Derivative of 86-028NP <i>rpsL</i> with a deletion mutation of <i>hktE</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> ( <i>phktE</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> transformed with pSPEC1- <i>hktE</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>pgdX</i>	Derivative of 86-028NP <i>rpsL</i> with a deletion mutation of <i>pgdX</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>pgdX</i> ( <i>ppgdX</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>pgdX</i> transformed with pSPEC1- <i>pgdX</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>dps</i>	Derivative of 86-028NP <i>rpsL</i> with a deletion mutation of <i>dps</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> ( <i>pdps</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> transformed with pSPEC1- <i>dps</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> ( <i>phktE</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> transformed with pSPEC1- <i>hktE</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>fur</i>	Derivative of 86-028NP <i>rpsL</i> with a deletion mutation of <i>fur</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> $\Delta$ <i>pgdX</i>	Derivative of 86-028NP <i>rpsL</i> with deletion mutations of <i>hktE</i> and <i>pgdX</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> $\Delta$ <i>pgdX</i> ( <i>phktE</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> $\Delta$ <i>pgdX</i> transformed with pSPEC1- <i>hktE</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> $\Delta$ <i>pgdX</i> ( <i>ppgdX</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> $\Delta$ <i>pgdX</i> transformed with pSPEC1- <i>pgdX</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> $\Delta$ <i>hktE</i>	Derivative of 86-028NP <i>rpsL</i> with deletion mutations of <i>dps</i> and <i>hktE</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> $\Delta$ <i>hktE</i> ( <i>pdps</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> $\Delta$ <i>hktE</i> transformed with pSPEC1- <i>dps</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> $\Delta$ <i>hktE</i> ( <i>phktE</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> $\Delta$ <i>hktE</i> transformed with pSPEC1- <i>hktE</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> $\Delta$ <i>fur</i>	Derivative of 86-028NP <i>rpsL</i> with deletion mutations of <i>hktE</i> and <i>fur</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> $\Delta$ <i>fur</i> ( <i>phktE</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>hktE</i> $\Delta$ <i>fur</i> transformed with pSPEC1- <i>hktE</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>pgdX</i> $\Delta$ <i>fur</i>	Derivative of 86-028NP <i>rpsL</i> with deletion mutations of <i>pgdX</i> and <i>fur</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>pgdX</i> $\Delta$ <i>fur</i> ( <i>ppgdX</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>pgdX</i> $\Delta$ <i>fur</i> transformed with pSPEC1- <i>pgdX</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> $\Delta$ <i>fur</i>	Derivative of 86-028NP <i>rpsL</i> with deletion mutations of <i>dps</i> and <i>fur</i>	This study
86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> $\Delta$ <i>fur</i> ( <i>pdps</i> )	86-028NP <i>rpsL</i> $\Delta$ <i>dps</i> $\Delta$ <i>fur</i> transformed with pSPEC1- <i>dps</i>	This study

stress protection systems is essential for an NTHI cell, both as a commensal and during the course of infection (11). In the current studies, we generated mutant strains of 86-028NP that do not express catalase, TsaA, PgdX, or Dps. We also generated double mutants that do not express catalase and PgdX, catalase and Dps, or PgdX and Dps. Finally we generated mutations in catalase, PgdX, and Dps in a strain that lacked the ferric uptake regulator, Fur. These strains were tested for their ability to withstand the effects of oxidative stress induced by *in vitro* treatment with H<sub>2</sub>O<sub>2</sub>. Our data indicate that strain 86-028NP possesses multiple, overlapping oxidative stress defense mechanisms with critical roles in protection against iron-exacerbated oxidative stress. We hypothesize that regulation of these mechanisms is critical for NTHI survival *in vivo*.

## MATERIALS AND METHODS

**Bacterial strains and culture media used.** NTHI strain 86-028NP, recovered from the nasopharynx of a child with chronic OM, has been well characterized *in vitro* (13, 14) and in a chinchilla model of OM (15–17). The genome sequence has been published (12). A streptomycin-resistant derivative of strain 86-028NP with a mutation in *rpsL*, 86-028NP*rpsL*, was used as the parent strain in mutant construction, as described in Caruthers et al. (18).

For routine growth, NTHI strains were cultured on chocolate II agar plates (Fisher Scientific, Pittsburgh, PA). For counterselection, strain 86-028NP*rpsL* was cultured on chocolate agar plates containing 1,000  $\mu$ g streptomycin/ml. The complemented NTHI strains were cultured on chocolate agar plates containing 200  $\mu$ g spectinomycin/ml. For routine liquid culture, NTHI cells were grown in brain heart infusion broth supplemented with 2  $\mu$ g NAD/ml and 2  $\mu$ g heme/ml (sBHI). Unless stated otherwise, all growth was with shaking at 180 rpm at 37°C. Cell growth was tracked by measuring optical density at 600 nm.

**Construction of mutant strains.** All mutant strains were generated using a recombinering method developed for strain 86-028NP (18). Mutants were selected on medium containing 200  $\mu$ g spectinomycin/ml. The spectinomycin resistance antibiotic cassette was then removed by site-

specific recombination. After the removal of the spectinomycin resistance antibiotic cassette, additional mutations could be introduced using the same method. The identity of each mutant was confirmed by PCR and sequencing.

**Construction of complemented mutant strains.** Each complemented strain was generated as follows. The coding sequence, as well as the upstream sequence that contained a predicted OxyR-binding site, was amplified by PCR. The primer upstream from the gene contained a BamHI restriction site. The primer downstream from the gene contained an EcoRI restriction site. The PCR amplicon was cloned between the BamHI and EcoRI sites in pSPEC1, a derivative of the *Haemophilus-Actinobacillus pleuropneumoniae* shuttle vector pGZRS-39A, in which the kanamycin resistance gene was replaced by a spectinomycin resistance gene (16, 19). Each construct was transformed into its respective mutant by electroporation, and transformants selected on chocolate agar containing 200  $\mu$ g spectinomycin/ml. Strains are listed on Table 1.

**Quantitative RT-PCR.** Quantitative reverse transcription (RT)-PCR (qRT-PCR) was used to determine the expression of genes with predicted roles in combating oxidative stress. qRT-PCR was performed with a one-step QuantiTect SYBR green RT-PCR kit (Qiagen, Valencia, CA) as outlined by Harrison et al. (20). Three biological replicates and three technical replicates were performed for each gene analyzed. Fold changes were calculated. All threshold cycle ( $C_T$ ) values were normalized to the value for the endogenous control *gyrA*. Relative quantitation was calculated from the median  $C_T$  value using  $\Delta\Delta C_T$ , and statistical significance was determined using Student's two-tailed *t* test. A fold change in gene expression greater than 2-fold and with a *P* value of <0.05 was assessed as being significant.

***In vitro* sensitivity tests with hydrogen peroxide.** Unless stated otherwise, strain 86-026NP*rpsL* and each mutant to be tested were grown in sBHI with shaking at 180 rpm to mid-exponential phase. H<sub>2</sub>O<sub>2</sub> was then added to give a final concentration of 500  $\mu$ M. Growth was continued for 10 min with shaking at 180 rpm. Cells were then removed, serially diluted, and plated on chocolate agar to assess viability.

**Iron chelation tests with 2,2'-bipyridine.** To chelate iron from cultures, exponentially growing cells were treated with 500  $\mu$ M 2,2'-bipyridine, a treatment that we previously showed affected iron levels in sBHI

(20). Cells which did not receive 500  $\mu\text{M}$  2,2'-bipyridine had ethanol added to a final concentration of 0.2% to replicate the addition of ethanol used as the vehicle for 2,2'-bipyridine. Chelation was carried out concurrently with  $\text{H}_2\text{O}_2$  production, and cell viability was assessed on chocolate agar as described above.

**Quantitation of peroxidase activity.** Strain 86-026NP*rpsL* and each mutant to be tested were grown in sBHI with shaking at 180 rpm to mid-exponential phase. Cultures were then split into two 20-ml aliquots, and  $\text{H}_2\text{O}_2$  added to one aliquot of each strain to a final concentration of 500  $\mu\text{M}$ . After a further 30 min of growth, 5-ml amounts of cells, with or without  $\text{H}_2\text{O}_2$  treatment, were chilled on ice and then harvested by centrifugation at  $3,220 \times g$ , washed once in 5 ml cold 0.1 M potassium phosphate buffer, pH 6.8, resuspended in 1.7 ml cold 0.1 M potassium phosphate buffer, pH 6.8, and mechanically lysed using lysing matrix B (MP Biomedicals, Solon, OH). Lysates were cleared of lysing matrix by centrifugation at  $27 \times g$  for 5 min, and protein concentrations calculated using a bicinchoninic acid (BCA) assay (Thermo Scientific, Rockford, IL). Twenty-five-microliter amounts of the resulting lysates were then used in a catalase activity assay using an Amplex red catalase assay kit (Life Technologies, Grand Island, NY).

**Statistical analyses of  $\text{H}_2\text{O}_2$  sensitivity and peroxidase activity assays.** Data produced by the  $\text{H}_2\text{O}_2$  sensitivity and peroxidase activity assays were tested for significance using a multiple *t* test with Bonferroni correction. All analyses were carried out with Prism 6 (GraphPad Software, La Jolla, CA).

## RESULTS

**Roles of catalase, peroxiredoxin/glutaredoxin, and alkylhydroperoxidase in protection of NTHI against hydrogen peroxide-induced oxidative stress.** We had previously determined that the loss of OxyR, the master transcriptional regulator of genes with roles in protection against oxidative stress, had an effect on the viability of cells treated with both 250  $\mu\text{M}$  and 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$ . We thus hypothesized that one or more of these OxyR-regulated genes would be critical to the survival of strain 86-028NP when subjected to  $\text{H}_2\text{O}_2$ -induced oxidative stress. We have previously shown that the expression of *tsaA* was unresponsive to treatment with  $\text{H}_2\text{O}_2$ , and *tsaA* is not a member of the OxyR regulon in strain 86-028NP (21). In concert with these data, when the *tsaA* mutant and its parental strain were grown to late exponential phase and treated with 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$ , there was no significant difference in the rates of viability of the *tsaA* mutant and the parent (data not shown). Conversely, treatment with 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$  produced a slight but significant decrease in the viability of the *hktE* mutant compared to that of the parent strain (Fig. 1A). The catalase mutation was complemented with a plasmid-encoded parental copy of *hktE* downstream from its native 5' untranslated region (UTR), which contained a predicted OxyR-binding site. Restoration of OxyR-regulated *hktE* in the complemented *hktE* mutant abrogated the susceptibility of the *hktE* mutant to  $\text{H}_2\text{O}_2$  (Fig. 1A). This experiment was repeated with the *pgdX* mutant. The loss of PgdX resulted in increased viability compared to that of the parent strain after treatment with 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$ . This increased survival was reduced by complementation of the *pgdX* mutation (Fig. 1B). These data suggested the compensatory effect of an additional oxidative stress protection mechanism in the absence of PgdX. Possibly an increase in catalase was compensating for the loss of PgdX, as previously demonstrated in *H. influenzae* strain Rd (22). This hypothesis was supported by the observation that a strain that lacked both *hktE* and *pgdX* demonstrated reduced viability, albeit nonsignificant, compared to that of the parent after treatment with 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$  but that the loss of viability was abro-

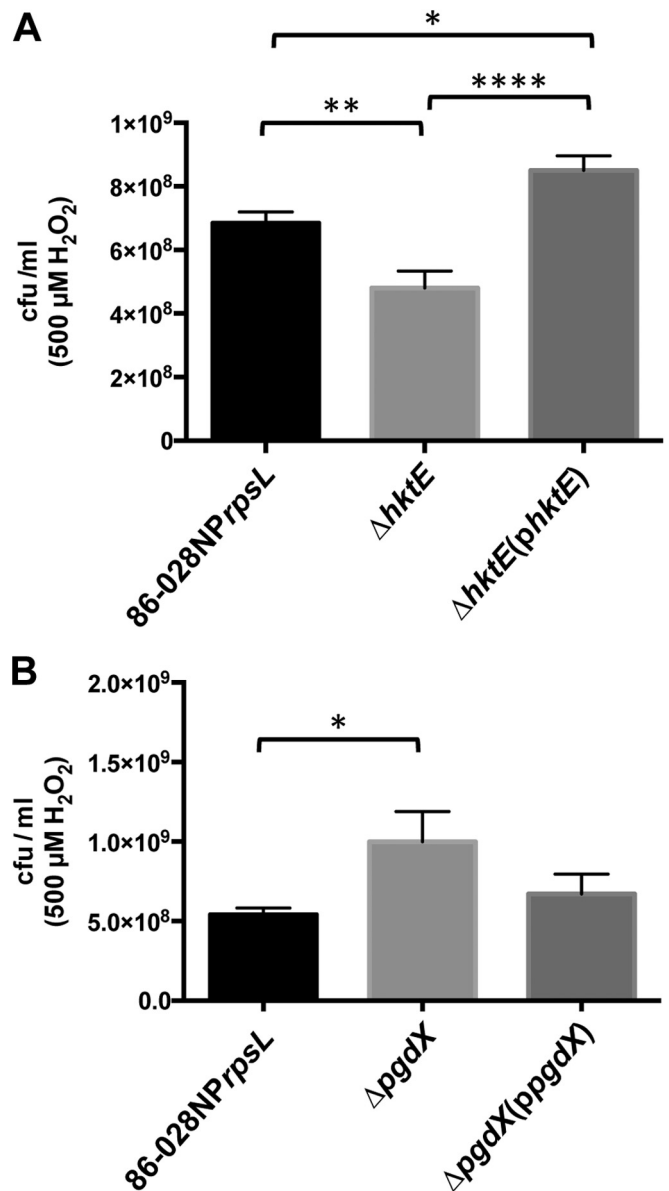
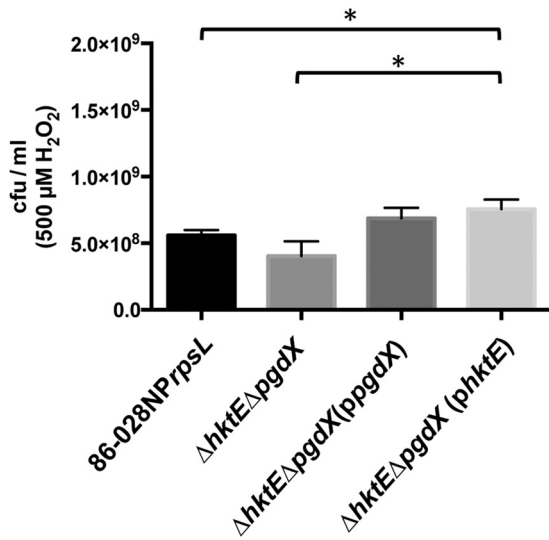


FIG 1 Both the catalase HktE and the peroxidase PgdX protect NTHI strain 86-028NP from  $\text{H}_2\text{O}_2$ -induced oxidative stress. Strains of 86-028NP that lacked either *hktE* (A) or *pgdX* (B) were grown exponentially and then treated with 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$  for 10 min. The cells were harvested and plated to assess viability relative to that of the parent strain. Loss of *hktE* produced a slight but significant loss in cell viability, while the strain that lacked *pgdX* exhibited increased viability. Complementation of the mutated gene restored the parental phenotype in both cases. Error bars show standard errors of the means. \*,  $P \leq 0.05$ ; \*\*,  $P \leq 0.01$ ; \*\*\*\*,  $P \leq 0.0001$  ( $n = 9$  [A];  $n = 6$  [B]).

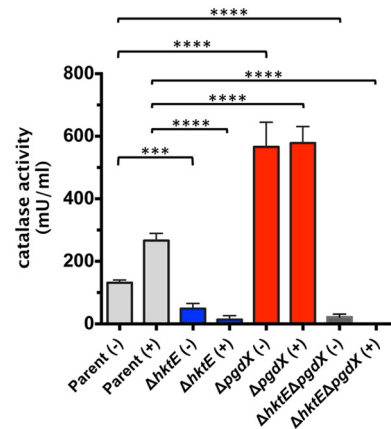
gated when the *hktE pgdX* double mutant strain was complemented with either *hktE* or *pgdX* (Fig. 2).

To determine whether there was increased *hktE* expression in the absence of PgdX, qRT-PCR analysis was used to quantify *hktE* expression in the *pgdX* mutant. Unexpectedly, when the mutant was treated with  $\text{H}_2\text{O}_2$ , the loss of PgdX led to a small, nonsignificant decrease in *hktE* expression that was reversed when the mutation was complemented. Little effect on *hktE* expression in the  $\Delta pgdX$  background was observed in the absence of peroxide-in-



**FIG 2** Catalase and the peroxidase PgdX together protect NTHI strain 86-028NP from  $H_2O_2$ -induced oxidative stress. A strain of 86-028NP that lacked both *hktE* and *pgdX* was grown exponentially and then treated with  $500 \mu M H_2O_2$  for 10 min. The cells were harvested and plated to assess viability. Loss of both *hktE* and *pgdX* produced a slight loss in cell viability. Complementation of the double mutant with either *hktE* or *pgdX* restored the parental phenotype. Error bars show standard errors of the means. \*,  $P \leq 0.05$  ( $n = 7$ ).

duced stress (data not shown). We further quantified the ability of the parent strain to decompose  $H_2O_2$ . The *hktE* mutant, the *pgdX* mutant, and the *hktE pgdX* double mutant strain were tested similarly. Each strain was grown to late exponential phase and then split into two aliquots. One aliquot was treated with  $500 \mu M H_2O_2$ , as in the sensitivity tests. The second aliquot was left untreated. Cell extracts were prepared and assayed using an Amplex red catalase activity kit from Life Technologies, which measures decomposition of  $H_2O_2$  as a surrogate for catalase activity. When extracts from cells untreated with  $H_2O_2$  were compared, the *hktE* mutant demonstrated an approximately 3-fold reduction in activity compared to the results for the parent strain. Strikingly, without  $H_2O_2$  pretreatment, the *pgdX* mutant decomposed  $H_2O_2$  with approximately 4-fold greater activity than the parent. However, this increase in enzymatic activity was lost in the strain that lacked both catalase and PgdX, suggesting that the increase in the ability to decompose  $H_2O_2$  in the *pgdX* mutant was due to increased catalase enzymatic activity (Fig. 3). This thesis was supported by data that compared the levels of  $H_2O_2$  decomposition activity in strains pretreated with  $H_2O_2$  with the levels in the strains that were left untreated. The parent exhibited an approximately 2-fold increase in activity when the cells were pretreated with  $H_2O_2$  compared to the activity in the cells that were untreated. When the *hktE* mutant was tested similarly, pretreatment with  $H_2O_2$  did not produce an increase in decomposition activity. Actually, pretreatment of the *hktE* mutant produced a slight but nonsignificant decrease in activity compared to that in the untreated cells (Fig. 3). However, when either the *pgdX* mutant or the  $\Delta hktE \Delta pgdX$  mutant was tested, pretreatment with  $H_2O_2$  had no significant effect on decomposition activity compared to the results for the untreated strains. These data are in agreement with data derived from *H. influenzae* strain Rd (22) and further suggest the presence of a third functional peroxidase in strain 86-028NP.



**FIG 3** A strain of 86-028NP that lacks the peroxidase PgdX has an increased ability to decompose  $H_2O_2$ . Strains of 86-028NP that lacked the genes encoding the catalase HktE, the peroxidase PgdX, or both HktE and PgdX were grown exponentially and then treated with (+)  $500 \mu M H_2O_2$  for 30 min. A control sample was left untreated (-). The cells were harvested and lysed, and their ability to decompose  $H_2O_2$  assessed using an Amplex red catalase assay kit. The addition of  $H_2O_2$  to the cells produced increased  $H_2O_2$  decomposition activity in the parent strain relative to the activity in the untreated cells. The *hktE* mutant showed a large reduction in  $H_2O_2$  decomposition activity. In contrast, the *pgdX* mutant demonstrated  $H_2O_2$  decomposition activity that was significantly greater than that of the parent. Slight activity remained in the strain that lacked both *hktE* and *pgdX*. Error bars show standard errors of the means. \*\*\*,  $P \leq 0.001$ ; \*\*\*\*,  $P \leq 0.0001$  ( $n = 8$ ).

**The loss of Dps increases NTHI's sensitivity to hydrogen peroxide-induced oxidative stress.** The toxic effects of  $H_2O_2$  are exacerbated by the presence of iron, due to the Fenton reaction. Ferric iron participates in the reductive decomposition of  $H_2O_2$ . This reaction leads to the production of hydroxyl radicals, which can profoundly damage DNA (23). Bacteria have therefore evolved mechanisms to abrogate the effects of the Fenton reaction. Aside from decomposition of  $H_2O_2$ , bacteria can sequester free iron. Notable among the proteins involved in iron sequestration is Dps, a ferritin-like protein whose function was first elucidated in *E. coli* (5). A *dps* mutant was generated in strain 86-028NP *rpsL*, and the importance of Dps in resistance to  $H_2O_2$ -induced oxidative stress was tested. Treatment of exponentially growing cells with  $500 \mu M H_2O_2$  produced an approximately 22-fold decrease in the viability of the *dps* mutant compared to that of the parent strain (Fig. 4). In contrast, a similarly treated catalase mutant exhibited a small decrease in viability compared to the viability of the parent strain (Fig. 1A). The *dps* mutation was complemented with a plasmid-encoded parental copy of *dps* downstream from its native 5' UTR, which contained a predicted OxyR-binding site. Restoration of OxyR-regulated *dps* in the complemented *dps* mutant abrogated the susceptibility of the *dps* mutant to  $H_2O_2$  (Fig. 4). We reasoned that the loss of Dps led to an increase in intracellular iron and, thus, in the presence of  $H_2O_2$ , to a decrease in cell viability due to the Fenton reaction. To determine the importance of  $H_2O_2$  in this process, the plasmid construct previously used to complement the catalase mutant (Fig. 1A) was introduced into the *dps* mutant. The sensitivity of this strain, 86-028NP *rpsL*  $\Delta dps$  (*phktE*), to  $500 \mu M H_2O_2$  was then assessed and found to be similar to that of the parental strain (Fig. 4). These data support the hypothesis that  $H_2O_2$ -mediated toxicity is due to the interplay between iron and  $H_2O_2$ , with the ability to chelate iron being critical to the

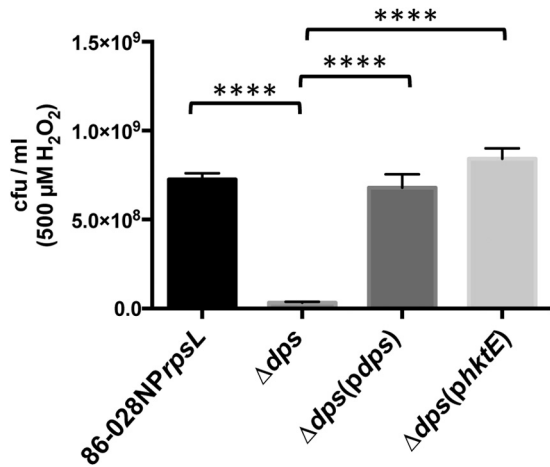


FIG 4 The ferritin-like protein Dps has a major role in the protection of NTHI strain 86-028NP from H<sub>2</sub>O<sub>2</sub>-induced oxidative stress. A strain of 86-028NP that lacked *dps* was grown exponentially and then treated with 500 μM H<sub>2</sub>O<sub>2</sub> for 10 min. The cells were harvested and plated to assess viability. Loss of *dps* produced a large and significant loss in cell viability relative to that of the parent strain. Complementation of the mutated gene with either *dps* or *hktE* restored the parental phenotype in both cases. Error bars show standard errors of the means. \*\*\*\*,  $P \leq 0.0001$  ( $n = 6$ ).

survival of the cell. We therefore generated an additional 86-028NP*rpsL* strain in which both *hktE* and *dps* were mutated. After treatment of exponentially growing cells with 500 μM H<sub>2</sub>O<sub>2</sub>, there was approximately a 110-fold decrease in the viability of cells due to the loss of both Dps and catalase (Fig. 5). When either Dps or catalase was introduced into the *hktE dps* double mutant strain via

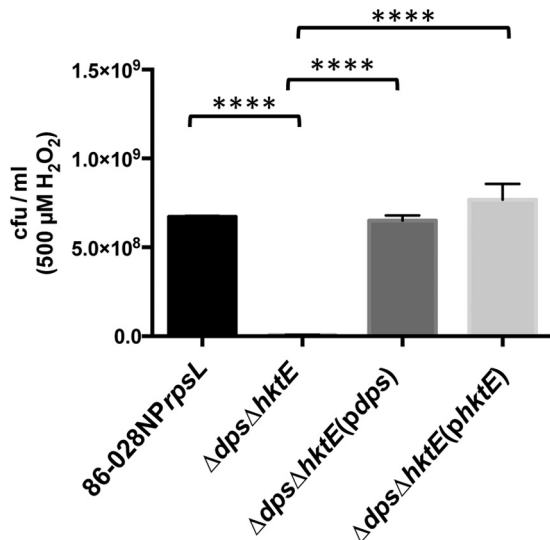


FIG 5 The activities of both the ferritin-like protein Dps and the catalase HktE are critical in the protection of NTHI strain 86-028NP from H<sub>2</sub>O<sub>2</sub>-induced oxidative stress. A strain of 86-028NP that lacked both *hktE* and *dps* was grown exponentially and then treated with 500 μM H<sub>2</sub>O<sub>2</sub> for 10 min. The cells were harvested and plated to assess viability. Loss of both *hktE* and *dps* produced a large and significant loss in cell viability relative to that of the parent strain. Complementation of the double mutant with either *hktE* or *dps* restored the parental phenotype. Error bars show standard errors of the means. \*\*\*\*,  $P \leq 0.0001$  ( $n = 5$ ).

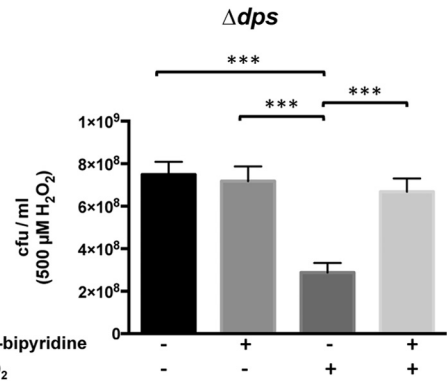


FIG 6 Minimizing the interplay between iron and H<sub>2</sub>O<sub>2</sub> is critical in the protection of NTHI strain 86-028NP from H<sub>2</sub>O<sub>2</sub>-induced oxidative stress. A strain of 86-028NP that lacked *dps* was grown exponentially and then treated with (+) or without (-) 500 μM H<sub>2</sub>O<sub>2</sub> for 10 min. The treatment with H<sub>2</sub>O<sub>2</sub> was carried out in the presence (+) or absence (-) of the iron chelator 2,2'-bipyridine. The cells were harvested and plated to assess viability. The *dps* mutant exhibited a loss of viability in the absence of iron chelation. In contrast, iron chelation protected the *dps* mutant from the effects of H<sub>2</sub>O<sub>2</sub>. Complementation of the *dps* mutation restored the parental phenotype under all conditions tested (data not shown). Error bars show standard errors of the means. \*\*\*,  $P \leq 0.001$  ( $n = 6$ ).

*pdps* or *phkTE*, the viability of the double mutant after H<sub>2</sub>O<sub>2</sub> treatment returned to parental levels (Fig. 5).

We further confirmed the importance of iron in H<sub>2</sub>O<sub>2</sub>-mediated killing of strain 86-028NP*rpsL* by comparing the toxic effects of H<sub>2</sub>O<sub>2</sub> on strain 86-028NP*rpsL* in iron-replete medium and iron-depleted medium. Briefly, exponentially grown cells were split into two aliquots, and one aliquot was treated with 500 μM 2,2'-bipyridine. Treatment with H<sub>2</sub>O<sub>2</sub> or with 2,2'-bipyridine had no significant effect on the viability of the parental strain (data not shown). Conversely, H<sub>2</sub>O<sub>2</sub> treatment significantly decreased the viability of the *dps* mutant compared to that of the *dps* mutant that was untreated (Fig. 6). The reduction in viability of the *dps* mutant due to H<sub>2</sub>O<sub>2</sub> treatment was abrogated when the iron in the medium was depleted with 2,2'-bipyridine (Fig. 6). Finally, when the experiment was repeated with the strain in which the *dps* mutation was complemented, there were no significant differences in the rates of viability of the strain under all treatments tested (data not shown).

The critical importance of the interplay between iron and H<sub>2</sub>O<sub>2</sub> in the induction of oxidative stress was further demonstrated using strains in which the mutations in catalase, *pgdX*, and *dps* were introduced into a strain with a deletion of the gene that encodes the ferric uptake regulator, Fur. We had previously generated a strain of 86-028NP carrying a marked deletion of *fur* (20). For this study, we generated a strain of 86-028NP*rpsL* carrying an unmarked deletion of *fur*. We could then easily introduce additional mutations into this strain. We previously showed that strain 86-028NP, which lacked Fur, had upregulated expression of many genes involved in iron import (20). Microarray analysis carried out as detailed in our previous studies (20) showed concordance between the Fur-regulated genes identified using the marked *fur* mutant and the unmarked *fur* mutant (data not shown). We therefore hypothesized that a *fur* mutant of strain 86-028NP*rpsL* would have elevated amounts of intracellular iron and so be more susceptible to the effects of H<sub>2</sub>O<sub>2</sub>. Previous killing assays were

carried out with exponentially growing cells that had been grown in sBHI with shaking at 180 rpm. However, the *pgdX fur* double mutant would not grow under these conditions (data not shown). We hypothesized that shaking at 180 rpm introduced oxygen into the medium at a level the mutants could not tolerate. This growth phenotype was partially overcome when shaking was reduced to 50 rpm. When shaken at 50 rpm, all strains except the *pgdX fur* double mutant had similar growth kinetics. The *pgdX fur* double mutant demonstrated a delayed entrance into exponential phase and had a lower rate of exponential growth than the other strains (data not shown). Despite the differences in growth kinetics, it was possible to test the viability of all strains after exposure to H<sub>2</sub>O<sub>2</sub>. The killing assays with 500 μM H<sub>2</sub>O<sub>2</sub> were thus repeated as detailed above, with the caveat that cells were grown to exponential phase while shaking at 50 rpm. Treatment with 500 μM H<sub>2</sub>O<sub>2</sub> produced a slight but nonsignificant increase in the survival of the *hktE* mutant (Fig. 7A). This stands in contrast with the results of the same experiment carried out with cells grown at 180 rpm, where there was a slight but significant decrease in the viability of the *hktE* mutant compared to that of the parent strain (Fig. 1A). When the *fur* mutant was treated similarly, there was a slight but nonsignificant reduction in the viability of the *fur* mutant compared to that of the parent (Fig. 7A). Conversely, when the *hktE fur* double mutant was treated with 500 μM H<sub>2</sub>O<sub>2</sub>, the viability of the mutant was reduced by approximately 26-fold compared to that of the parent (Fig. 7A). The sensitivity to H<sub>2</sub>O<sub>2</sub> of the *hktE fur* mutant was abrogated when *hktE* was introduced into the mutant via the complementing plasmid. This experiment was repeated with the *pgdX fur* double mutant. As shown by the results described above, the loss of PgdX resulted in increased viability after treatment with 500 μM H<sub>2</sub>O<sub>2</sub> compared to the viability of the parent strain (Fig. 7B). However, in contrast to the results for the *hktE fur* double mutant, which demonstrated a significant reduction in viability compared to that of the *fur* mutant, treatment with 500 μM H<sub>2</sub>O<sub>2</sub> produced a slight increase in the survival of the *pgdX fur* double mutant compared to the survival of the *fur* mutant (Fig. 7B). Loss of *fur* in the *pgdX fur* double mutant led to an approximately 5-fold decrease in survival compared to that of the *pgdX* mutant (Fig. 7B). Finally, we determined how the loss of both *fur* and *dps* affected cell viability. When the culture was growing exponentially, treatment of the *dps* mutant with 500 μM H<sub>2</sub>O<sub>2</sub> produced approximately a 5-log decrease in viability compared to that of the parent strain (Fig. 7C). This stands in contrast to the data derived from cells grown at 180 rpm, where loss of *dps* led to a decrease in viability of only 22-fold compared to that of the parent (Fig. 4). When the experiment was repeated with the *dps fur* double mutant, the number of viable cells recovered after treatment with 500 μM H<sub>2</sub>O<sub>2</sub> was below the limit of detection (Fig. 7C).

## DISCUSSION

*H. influenzae* can grow both aerobically and as a facultative anaerobe. During aerobic growth, the metabolism of *H. influenzae* will generate oxidative stress. *H. influenzae* will also endure oxidative stress derived from host defense cells (24, 25) and from copathogens (primarily *Streptococcus pneumoniae* [26]). In all niches, NTHI is also in an iron-restricted environment and so must have avid and well-regulated mechanisms to import iron. NTHI must therefore carefully balance the need to acquire scarce sources of iron while minimizing the generation of hydroxyl radicals via the

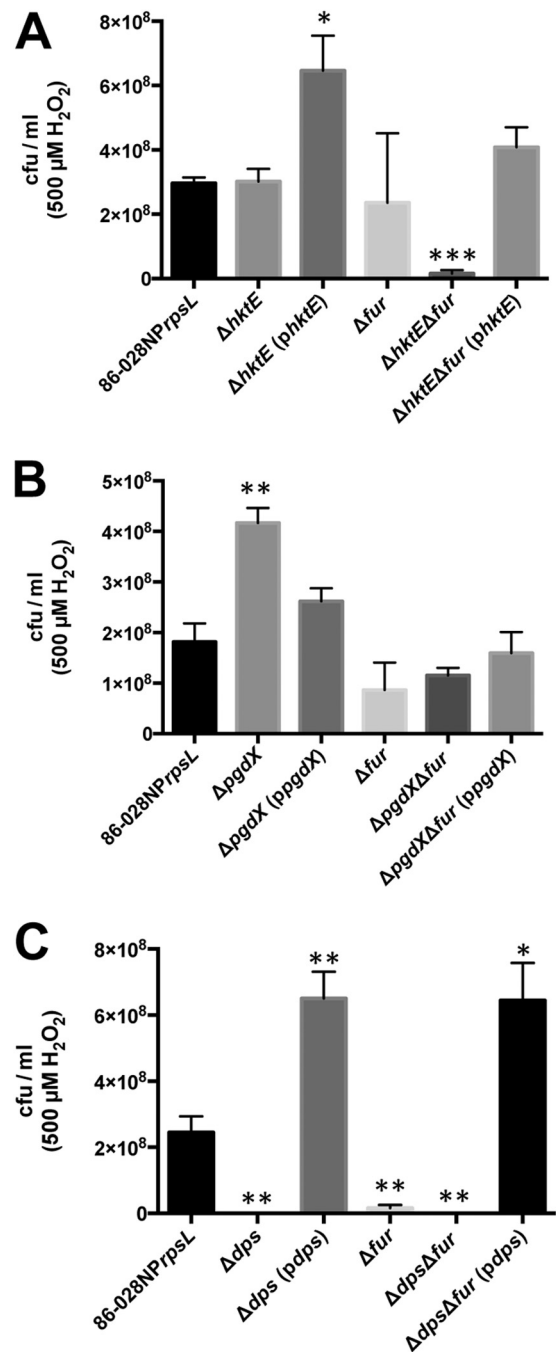


FIG 7 Unregulated import of iron into NTHI strain 86-028NP exacerbates the effects of H<sub>2</sub>O<sub>2</sub>-induced killing. The genes encoding catalase (*hktE*) (A), peroxiredoxin glutaredoxin (*pgdX*) (B), or the ferritin-like protein Dps (*dps*) (C) were deleted in a strain of 86-028NP that lacked the ferric uptake regulator, Fur. All strains were grown exponentially and then treated with 500 μM H<sub>2</sub>O<sub>2</sub> for 10 min. The cells were harvested and plated to assess viability. (A, B) Loss of the ability to decompose H<sub>2</sub>O<sub>2</sub> in the face of unrestricted iron import produced significant losses in cell viability relative to that of the parent strain or the respective single mutant strain. (C) Loss of the ability to bind iron in the *dps* mutant produced a nearly total loss of cell viability when iron import was unregulated and the cells were treated with H<sub>2</sub>O<sub>2</sub>. Complementation of the double mutants with *hktE*, *pgdX*, or *dps* restored cell viability. Significance was calculated relative to the results for the parent strain. Error bars show standard errors of the means. \*,  $P \leq 0.05$ ; \*\*,  $P \leq 0.01$ ; \*\*\*,  $P \leq 0.001$ ; \*\*\*\*,  $P \leq 0.0001$  ( $n = 3$  [A];  $n = 4$  [B, C]).

iron-mediated Fenton reaction. To mediate this balance, *H. influenzae* has evolved multiple overlapping defenses that allow the organism to survive in such a hostile environment.

**Enzymatic decomposition of H<sub>2</sub>O<sub>2</sub> is protective in NTHI.** To determine the respective roles of the products of genes presumed to protect against oxidative stress in NTHI strain 86-028NP, we generated strains that lacked the genes that encode catalase, PgdX, TsaA, and Dps. Using these strains, we demonstrated that TsaA had no role in protection against H<sub>2</sub>O<sub>2</sub>-induced oxidative stress. This is not unexpected, as we previously showed that the peroxide stress regulator OxyR does not regulate the expression of TsaA (21). The *E. coli* ortholog of TsaA, AhpC, has a primary role in the protection against endogenously generated oxidative stress. Concentrations of exogenously applied H<sub>2</sub>O<sub>2</sub> in excess of 20 μM overwhelm AhpC's ability to decompose the H<sub>2</sub>O<sub>2</sub>, so *E. coli* must rely on additional mechanisms of protection (27). NTHI strain 86-028NP lacks an ortholog of AhpC's cognate reductase AhpR. TsaA may well be reduced promiscuously by another reductase, but the lack of a cognate reductase and the insensitivity of *tsaA* expression to OxyR regulation suggest that TsaA's function in strain 86-028NP is different from that of AhpC in *E. coli*.

The results of hydrogen peroxide killing assays did, however, show a requirement for catalase, PgdX, and Dps. As expected, catalase protected against endogenously applied oxidative stress. However, the protection was not absolute. The catalase mutant only showed a slight reduction in viability compared to the viability of the parent. Other mechanisms generated an additional level of protection. This idea of an overlap in protective mechanisms against peroxide-induced stress was supported by data generated using the strain of 86-028NP that lacked *pgdX*. The *pgdX* mutant exhibited an increase in strain viability relative to that of the parent after H<sub>2</sub>O<sub>2</sub> treatment. Increased resistance to H<sub>2</sub>O<sub>2</sub> by a *pgdX* mutant was also observed by Pauwels et al. (22). In *H. influenzae* strain Rd, the absence of PgdX leads to suboptimal decomposition of H<sub>2</sub>O<sub>2</sub>, leading to an increased intracellular concentration of H<sub>2</sub>O<sub>2</sub>. This increase in H<sub>2</sub>O<sub>2</sub> produces a greater response in OxyR activity, with a concomitant upregulation of genes in the OxyR regulon (22). However, when the expression of *hktE* was assessed in the strain 86-028NP *pgdX* mutant, we observed no significant changes in *hktE* expression when oxidative stress was induced by H<sub>2</sub>O<sub>2</sub>. Similarly, there was no significant change in the expression of *dps* in the *pgdX* mutant (data not shown). *hktE* and *dps* are members of the OxyR regulon in strain 86-028NP (21). The lack of PgdX does not appear to lead to a change in gene expression of the members of the OxyR regulon in strain 86-028NP. The enhanced survival of the *pgdX* mutant undergoing oxidative stress must be due to additional regulatory mechanisms. The changes in peroxidase activity in the *pgdX* mutant, despite no increase in *hktE* transcripts, suggest that the activity of HktE may be either post-transcriptionally or posttranslationally regulated in strain 86-028NP. Alternatively, there may be a third peroxidase present in strain 86-028NP, as shown by the  $\Delta hktE \Delta pgdX$  mutant still showing resistance to H<sub>2</sub>O<sub>2</sub>, as well as the ability to decompose H<sub>2</sub>O<sub>2</sub>, albeit at a very low level. Finally, a disruption in the ability of strain 86-028NP to decompose H<sub>2</sub>O<sub>2</sub> may lead to a modulation of non-enzymatic protective mechanisms.

**Regulation of free iron is critical in the protection of NTHI from oxidative stress.** The exacerbation of H<sub>2</sub>O<sub>2</sub>-induced oxidative stress by the presence of free ferrous iron has been well described. The oxidation of iron from its ferrous to its ferric state

results in the generation of hydroxyl radicals that are injurious to DNA and many cellular proteins (3, 28). NTHI, like most bacteria, has an absolute requirement for extracellular sources of iron. NTHI has therefore evolved multiple mechanisms to take up both iron and iron-containing proteins, mechanisms tightly controlled by the ferric uptake regulator, Fur (20). NTHI specifically and bacteria in general must balance the need to take up iron with the requirement that iron is sequestered to prevent driving the Fenton reaction. The effects of a breakdown in this control can be readily observed through the use of strain 86-028NP mutants that lack both the ability to decompose H<sub>2</sub>O<sub>2</sub> and to sequester iron.

The deletion of *dps* from strain 86-028NP will produce a cell in which the ability to sequester free iron is substantially impaired. Introducing the *dps* mutation into strains in which either *hktE* or *pgdX* was mutated allowed us to determine the effect free iron had in the presence of externally applied H<sub>2</sub>O<sub>2</sub> that cannot be efficiently decomposed. Loss of *dps* alone did indeed increase the sensitivity of the cells to treatment with H<sub>2</sub>O<sub>2</sub>, but the loss of viability was greater when both *dps* and *hktE* were absent. The importance of the interplay between the ability to sequester free iron and the decomposition of H<sub>2</sub>O<sub>2</sub> was further demonstrated when a plasmid-encoded copy of *hktE* (*phktE*) was introduced into the *dps hktE* double mutant. When *phktE* was introduced into the *hktE* mutant, the complemented strain exhibited significantly increased resistance to H<sub>2</sub>O<sub>2</sub> compared to the resistance of the parent strain. The overexpression of *hktE*, by virtue of being on a multicopy plasmid, generated enhanced resistance to H<sub>2</sub>O<sub>2</sub>. The introduction of *phktE* into the *dps hktE* double mutant was similarly protective; the strain could decompose sufficient H<sub>2</sub>O<sub>2</sub> to restore the parental phenotype despite the absence of Dps' ability to chelate iron. The cells are therefore protected from H<sub>2</sub>O<sub>2</sub>-induced oxidative stress.

The importance of the ability to sequester iron as protection against H<sub>2</sub>O<sub>2</sub>-induced oxidative stress was further demonstrated when iron was chelated as the *dps* mutant was undergoing treatment with H<sub>2</sub>O<sub>2</sub>. We have previously demonstrated that treatment of strain 86-028NP growing exponentially in sBHI with 2,2'-bipyridine induces the expression of genes in the Fur regulon (20). Treatment with 2,2'-bipyridine chelates iron from sBHI, a treatment that induces the expression of genes whose expression is normally repressed by Fur. The loss of endogenous iron binding in the *dps* mutant is compensated by iron chelation, which minimizes the effect of the Fenton reaction and so protects the *dps* mutant from the effects of hydroxyl radical formation.

Finally, we showed that increased iron uptake in a mutant that lacks Fur also had a detrimental effect on resistance against H<sub>2</sub>O<sub>2</sub>-induced oxidative stress. An initial indication that increased uptake of iron through the loss of Fur was exacerbating sensitivity to H<sub>2</sub>O<sub>2</sub> was the observation that the *pgdX fur* double mutant would not grow when shaken at 180 rpm. Moreover, the *pgdX fur* double mutant still exhibited an impaired growth phenotype when shaken at 50 rpm. We suggest that the changes in the growth kinetics of the *pgdX fur* double mutant relate to the amounts of dissolved oxygen present in the culture medium. Using well-accepted conventions for the setup of shaken liquid cultures, we ensured that the medium never comprised more than 20% of the volume of the 250-ml flask. This allows adequate aeration of the medium with atmospheric gases. Flasks that are shaken at faster speeds have increased dissolved oxygen compared to the levels in those shaken at slower speeds (29, 30). Oxygen diffuses freely

across cell membranes. Thus, the oxygen concentration within bacteria is very similar to that within the growth medium (27). We would therefore presume that when the culture is grown at 180 rpm, the increased oxygen in the medium is such that the oxygen-dependent generation of reactive oxygen species within cells occurs at a rate that overwhelms the ability of the double mutants to combat the oxidative stress generated. Reducing the amount of dissolved oxygen in the medium through reduction of the speed the cultures are shaken at minimizes this effect. The exception was the *pgdX fur* double mutant, which still demonstrated an impaired growth phenotype when grown with shaking at 50 rpm. In *H. influenzae* strain Rd and *H. influenzae* type b, PgdX protects against the bacteriostatic effects of endogenously produced H<sub>2</sub>O<sub>2</sub> (31). When the culture was untreated, the loss of *pgdX* alone in strain 86-028NP had no effect on viability, while the loss of *fur* alone produced an approximately 2-fold reduction in viability compared to that of the parent. However, the untreated *pgdX fur* double mutant exhibited viability similar to that of the parent (data not shown). As with cells treated with H<sub>2</sub>O<sub>2</sub>, the loss of *pgdX* in the exponentially growing but untreated *pgdX fur* double mutant must lead to upregulation of the expression of an additional H<sub>2</sub>O<sub>2</sub>-protective mechanism(s). The loss of both *pgdX* and *fur* thus has a bacteriostatic effect on exponentially growing 86-028NP in the absence of treatment with exogenous H<sub>2</sub>O<sub>2</sub>, as the cells exhibited a much reduced exponential growth rate but had viability similar to that of the parent. Significant killing of the strain 86-028NP *pgdX fur* double mutant only occurred after treatment with exogenous H<sub>2</sub>O<sub>2</sub>. The upregulation of an additional mechanism(s) of H<sub>2</sub>O<sub>2</sub> decomposition in the absence of PgdX cannot compensate for the loss of PgdX in the face of unrestricted iron import. Furthermore, both the *hktE fur* and the *pgdX fur* double mutants were killed similarly and more readily than either the *hktE* or *fur* single mutants. These data again support our hypothesis that protection against oxidative stress in NTHI is reliant on both decomposition of H<sub>2</sub>O<sub>2</sub> and regulation of free iron concentration.

Finally, the most striking data were generated with the *dps fur* double mutant. When this strain underwent H<sub>2</sub>O<sub>2</sub> treatment, there was a nearly total loss of viability. As the expression levels of both *hktE* and *pgdX* were not significantly altered in the absence of Dps (data not shown), we would presume that the *dps fur* double mutant maintains its ability to efficiently decompose H<sub>2</sub>O<sub>2</sub>. It is apparent that the most critical mechanism in the protection of NTHI against oxidative stress is the regulation of free iron. Even in the face of an armament of mechanisms to decompose H<sub>2</sub>O<sub>2</sub>, any residual H<sub>2</sub>O<sub>2</sub> left intact can interact with free iron, generate hydroxyl radicals via the Fenton reaction, and kill the cell.

Elucidating the interplay between both enzymes, as well as any additional peroxidase(s), will be critical in understanding how NTHI persists so well within the host. It is clear that NTHI must be able to protect against oxidative stress mediated by peroxide and iron. NTHI must also be able to protect against changes in oxygen concentrations, fluxes that will also generate oxidative stress. NTHI must therefore employ its armament of proteins that can decompose H<sub>2</sub>O<sub>2</sub> and protect against the effects of iron-mediated damage as a critical part of the survival of this important human pathogen within its host.

## ACKNOWLEDGMENT

This work was funded by NIAID/NIH grant R01-AI077897 to R.S.M.

## REFERENCES

- Imlay JA. 2013. The molecular mechanisms and physiological consequences of oxidative stress: lessons from a model bacterium. *Nat Rev Microbiol* 11:443–454. <http://dx.doi.org/10.1038/nrmicro3032>.
- Imlay JA. 2003. Pathways of oxidative damage. *Annu Rev Microbiol* 57: 395–418. <http://dx.doi.org/10.1146/annurev.micro.57.030502.090938>.
- Park S, You X, Imlay JA. 2005. Substantial DNA damage from submicromolar intracellular hydrogen peroxide detected in Hpx- mutants of *Escherichia coli*. *Proc Natl Acad Sci U S A* 102:9317–9322. <http://dx.doi.org/10.1073/pnas.0502051102>.
- Le Brun NE, Crow A, Murphy ME, Mauk AG, Moore GR. 2010. Iron core mineralisation in prokaryotic ferritins. *Biochim Biophys Acta* 1800: 732–744. <http://dx.doi.org/10.1016/j.bbagen.2010.04.002>.
- Chiancone E, Ceci P. 2010. The multifaceted capacity of Dps proteins to combat bacterial stress conditions: detoxification of iron and hydrogen peroxide and DNA binding. *Biochim Biophys Acta* 1800:798–805. <http://dx.doi.org/10.1016/j.bbagen.2010.01.013>.
- Almiron M, Link AJ, Furlong D, Kolter R. 1992. A novel DNA-binding protein with regulatory and protective roles in starved *Escherichia coli*. *Genes Dev* 6:2646–2654. <http://dx.doi.org/10.1101/gad.6.12b.2646>.
- Calhoun LN, Kwon YM. 2011. Structure, function and regulation of the DNA-binding protein Dps and its role in acid and oxidative stress resistance in *Escherichia coli*: a review. *J Appl Microbiol* 110:375–386. <http://dx.doi.org/10.1111/j.1365-2672.2010.04890.x>.
- Wong SM, Alugupalli KR, Ram S, Akerley BJ. 2007. The ArcA regulon and oxidative stress resistance in *Haemophilus influenzae*. *Mol Microbiol* 64:1375–1390. <http://dx.doi.org/10.1111/j.1365-2958.2007.05747.x>.
- Pang B, Hong W, Kock ND, Swords WE. 2012. Dps promotes survival of nontypeable *Haemophilus influenzae* in biofilm communities *in vitro* and resistance to clearance *in vivo*. *Front Cell Infect Microbiol* 2:58. <http://dx.doi.org/10.3389/fcimb.2012.00058>.
- Murphy TF, Faden H, Bakaletz LO, Kyd JM, Forsgren A, Campos J, Virji M, Pelton SI. 2009. Nontypeable *Haemophilus influenzae* as a pathogen in children. *Pediatr Infect Dis J* 28:43–48. <http://dx.doi.org/10.1097/INF.0b013e318184dba2>.
- Harrison A, Bakaletz LO, Munson J, Robert S. 2012. *Haemophilus influenzae* and oxidative stress. *Front Cell Infect Microbiol* 2:40. <http://dx.doi.org/10.3389/fcimb.2012.00040>.
- Harrison A, Dyer DW, Gillaspay A, Ray WC, Mungur R, Carson MB, Zhong H, Gipson J, Gipson M, Johnson LS, Lewis L, Bakaletz LO, Munson RS, Jr. 2005. Genomic sequence of an otitis media isolate of nontypeable *Haemophilus influenzae*: comparative study with *H. influenzae* serotype d, strain KW20. *J Bacteriol* 187:4627–4636. <http://dx.doi.org/10.1128/JB.187.13.4627-4636.2005>.
- Bakaletz LO, Tallan BM, Hoepf T, DeMaria TF, Birck HG, Lim DJ. 1988. Frequency of fimbriation of nontypeable *Haemophilus influenzae* and its ability to adhere to chinchilla and human respiratory epithelium. *Infect Immun* 56:331–335.
- Holmes KA, Bakaletz LO. 1997. Adherence of non-typeable *Haemophilus influenzae* promotes reorganization of the actin cytoskeleton in human or chinchilla epithelial cells *in vitro*. *Microb Pathog* 23:157–166. <http://dx.doi.org/10.1006/mpat.1997.0145>.
- Bakaletz LO, Leake ER, Billy JM, Kaumaya PT. 1997. Relative immunogenicity and efficacy of two synthetic chimeric peptides of fimbriae as vaccines against nasopharyngeal colonization by nontypeable *Haemophilus influenzae* in the chinchilla. *Vaccine* 15:955–961. [http://dx.doi.org/10.1016/S0264-410X\(96\)00298-8](http://dx.doi.org/10.1016/S0264-410X(96)00298-8).
- Mason KM, Munson RS, Jr, Bakaletz LO. 2003. Nontypeable *Haemophilus influenzae* gene expression induced *in vivo* in a chinchilla model of otitis media. *Infect Immun* 71:3454–3462. <http://dx.doi.org/10.1128/IAI.71.6.3454-3462.2003>.
- Vogel AR, Szelestey BR, Raffel FK, Sharpe SW, Gearing RL, Justice SS, Mason KM. 2012. SapF-mediated heme iron utilization enhances persistence and coordinates biofilm architecture of *Haemophilus*. *Front Cell Infect Microbiol* 2:42. <http://dx.doi.org/10.3389/fcimb.2012.00042>.
- Carruthers MD, Tracy EN, Dickson AC, Ganser KB, Munson RS, Jr, Bakaletz LO. 2012. Biological roles of nontypeable *Haemophilus influenzae* type IV pilus proteins encoded by the *pil* and *com* operons. *J Bacteriol* 194:1927–1933. <http://dx.doi.org/10.1128/JB.06540-11>.
- Mason KM, Bruggeman ME, Munson RS, Bakaletz LO. 2006. The non-typeable *Haemophilus influenzae* Sap transporter provides a mechanism of antimicrobial peptide resistance and SapD-dependent potassium



- acquisition. *Mol Microbiol* 62:1357–1372. <http://dx.doi.org/10.1111/j.1365-2958.2006.05460.x>.
20. Harrison A, Santana EA, Szelestey BR, Newsom DE, White P, Mason KM. 2013. Ferric uptake regulator and its role in the pathogenesis of nontypeable *Haemophilus influenzae*. *Infect Immun* 81:1221–1233. <http://dx.doi.org/10.1128/IAI.01227-12>.
  21. Harrison A, Ray WC, Baker BD, Armbruster DW, Bakaletz LO, Munson RS, Jr. 2007. The OxyR regulon in nontypeable *Haemophilus influenzae*. *J Bacteriol* 189:1004–1012. <http://dx.doi.org/10.1128/JB.01040-06>.
  22. Pauwels F, Vergauwen B, Van Beeumen JJ. 2004. Physiological characterization of *Haemophilus influenzae* Rd deficient in its glutathione-dependent peroxidase PGdx. *J Biol Chem* 279:12163–12170. <http://dx.doi.org/10.1074/jbc.M312037200>.
  23. Imlay JA, Linn S. 1988. DNA damage and oxygen radical toxicity. *Science* 240:1302–1309. <http://dx.doi.org/10.1126/science.3287616>.
  24. Craig JE, Cliffe A, Garnett K, High NJ. 2001. Survival of nontypeable *Haemophilus influenzae* in macrophages. *FEMS Microbiol Lett* 203:55–61. <http://dx.doi.org/10.1111/j.1574-6968.2001.tb10820.x>.
  25. Naylor EJ, Bakstad D, Biffen M, Thong B, Calverley P, Scott S, Hart CA, Moots RJ, Edwards SW. 2007. *Haemophilus influenzae* induces neutrophil necrosis: a role in chronic obstructive pulmonary disease? *Am J Respir Cell Mol Biol* 37:135–143. <http://dx.doi.org/10.1165/rcmb.2006-0375OC>.
  26. Pericone CD, Overweg K, Hermans PW, Weiser JN. 2000. Inhibitory and bactericidal effects of hydrogen peroxide production by *Streptococcus pneumoniae* on other inhabitants of the upper respiratory tract. *Infect Immun* 68:3990–3997. <http://dx.doi.org/10.1128/IAI.68.7.3990-3997.2000>.
  27. Imlay JA. 2008. Cellular defenses against superoxide and hydrogen peroxide. *Annu Rev Biochem* 77:755–776. <http://dx.doi.org/10.1146/annurev.biochem.77.061606.161055>.
  28. Anjem A, Imlay JA. 2012. Mononuclear iron enzymes are primary targets of hydrogen peroxide stress. *J Biol Chem* 287:15544–15556. <http://dx.doi.org/10.1074/jbc.M111.330365>.
  29. Wittmann C, Kim HM, John G, Heinzle E. 2003. Characterization and application of an optical sensor for quantification of dissolved O<sub>2</sub> in shake-flasks. *Biotechnol Lett* 25:377–380. <http://dx.doi.org/10.1023/A:1022402212537>.
  30. Schiefelbein S, Frohlich A, John GT, Beutler F, Wittmann C, Becker J. 2013. Oxygen supply in disposable shake-flasks: prediction of oxygen transfer rate, oxygen saturation and maximum cell concentration during aerobic growth. *Biotechnol Lett* 35:1223–1230. <http://dx.doi.org/10.1007/s10529-013-1203-9>.
  31. Vergauwen B, Herbert M, Van Beeumen JJ. 2006. Hydrogen peroxide scavenging is not a virulence determinant in the pathogenesis of *Haemophilus influenzae* type b strain Eagan. *BMC Microbiol* 6:3. <http://dx.doi.org/10.1186/1471-2180-6-3>.