# Alkyl Hydroperoxide Reductase Is the Primary Scavenger of Endogenous Hydrogen Peroxide in *Escherichia coli*

LAUREN COSTA SEAVER AND JAMES A. IMLAY\*

*Department of Microbiology, University of Illinois, Urbana, Illinois 61801*

Received 9 July 2001/Accepted 20 September 2001

**Hydrogen peroxide is generated during aerobic metabolism and is capable of damaging critical biomolecules. However, mutants of** *Escherichia coli* **that are devoid of catalase typically exhibit no adverse phenotypes during** growth in aerobic media. We discovered that catalase mutants retain the ability to rapidly scavenge  $H_2O_2$ **whether it is formed internally or provided exogenously. Analysis of candidate genes revealed that the residual activity is due to alkyl hydroperoxide reductase (Ahp). Mutants that lack both Ahp and catalase could not** scavenge  $H_2O_2$ . These mutants excreted substantial amounts of  $H_2O_2$ , and they grew poorly in air. Ahp is kinetically a more efficient scavenger of trace H<sub>2</sub>O<sub>2</sub> than is catalase and therefore is likely to be the primary scavenger of endogenous H<sub>2</sub>O<sub>2</sub>. Accordingly, mutants that lack Ahp accumulated sufficient hydrogen peroxide **to induce the OxyR regulon, whereas the OxyR regulon remained off in catalase mutants. Catalase still has an important role in wild-type cells, because the activity of Ahp is saturated at a low**  $(10^{-5} M)$  **concentration of**  $H_2O_2$ . In contrast, catalase has a high  $K_m$ , and it therefore becomes the predominant scavenger when  $H_2O_2$ **concentrations are high. This arrangement is reasonable because the cell cannot provide enough NADH for** Ahp to rapidly degrade large amounts of  $H_2O_2$ . In sum, *E. coli* does indeed generate substantial  $H_2O_2$ , but **damage is averted by the scavenging activity of Ahp.**

Virtually all aerobic organisms contain enzymes that convert superoxide and hydrogen peroxide to innocuous products. The ubiquity of these scavenging enzymes suggests that exposure to  $O_2$ <sup>-</sup> and  $H_2O_2$  is an inevitable part of the aerobic lifestyle and that these species can damage cells. It follows that organisms that cannot scavenge them will fare poorly in an aerobic habitat, and in 1971 McCord et al. proposed that some obligate anaerobes may be unable to grow in air at least in part because they lack sufficient levels of scavenging enzymes (17). This reasoning also suggested that aerobes would be much less oxygen tolerant if they lacked superoxide dismutase (SOD) and catalase. In 1985 this prediction was partly affirmed by studies of *sodA sodB* mutants of *Escherichia coli* (3). These strains suffered elevated rates of DNA damage, could not catabolize nonfermentable carbon sources, and did not grow at all without extensive amino acid supplements. However, contrary to expectation, mutants that lacked catalase grew as well as their wild-type parents and exhibited no increase in mutation rate (14, 29).

The fitness of the catalase mutants did not reflect invulnerability to  $H_2O_2$ . Low concentrations of exogenous  $H_2O_2$  (ca. 30)  $\mu$ M) are sufficient to inhibit the growth of *E. coli*. Although the growth-blocking cell lesions have not yet been identified,  $H_2O_2$ can oxidize thiols, which may inactivate enzymes that have active-site sulfhydryl residues. Methionine sulfoxide adducts (21) and protein carbonyls (5) may also be products of enzyme oxidation. In addition,  $H_2O_2$  can inactivate the exposed [4Fe-4S] clusters of aconitase B and fumarase B (S. M. Varghese, S. Korshunov, and J. A. Imlay, unpublished data). Finally, reactions between  $H_2O_2$  and intracellular iron generate hydroxyl radicals, which in turn attack DNA. Micromolar concentrations of  $H_2O_2$  are therefore mutagenic.

Thus, the robust performance of catalase mutants seemed to imply that these amounts of  $H_2O_2$  are not normally generated during aerobic growth. There are presently no data that firmly establish the amount of  $H_2O_2$  that is formed as a by-product of metabolism. No enzymes in *E. coli*, other than SOD, generate  $H<sub>2</sub>O<sub>2</sub>$  as a deliberate, stoichiometric product. However, in vitro studies have shown that  $H_2O_2$  can be formed by the adventitious oxidation of redox enzymes by molecular oxygen.

The main sources of  $H_2O_2$  inside the cell are probably flavoenzymes, both because they are abundant and because flavins are amenable to the univalent electron transfer reaction that initiates the production of superoxide and  $H_2O_2$  (6, 18). The rates at which some flavoenzymes generate these species have been measured in vitro, and extrapolations suggest that they may form about 10  $\mu$ M H<sub>2</sub>O<sub>2</sub> s<sup>-1</sup> in vivo (18). Gonzalez-Flecha and Demple reported evidence of endogenous  $H_2O_2$ production by *E. coli* (7). If these rates are correct, then the lack of phenotype of catalase mutants seems difficult to explain.

In this study, we discovered that mutants which are devoid of catalase still scavenge physiological concentrations of  $H_2O_2$  as rapidly as do wild-type cells. By identifying and eliminating the remaining scavenging enzyme, alkyl hydroperoxide reductase, we were able to directly quantify endogenous H<sub>2</sub>O<sub>2</sub> production and demonstrate that it is enough to poison scavengerless cells.

### **MATERIALS AND METHODS**

**Chemicals, enzymes, and media.** Cumene hydroperoxide, 5,5-dithio-bis(2 nitrobenzoic acid) (DTNB), horseradish peroxidase (type II), hydrogen peroxide, *o*-dianisidine, *o*-nitrophenyl-ß-D-galactopyranoside, plumbagin, and potassium cyanide were purchased from Sigma. Coomassie protein reagent was obtained from Pierce. Bovine liver catalase (20 mg/ml) was from Boehringer Mannheim, and Amplex red (AR) was from Molecular Probes. Dimethyl sul-

<sup>\*</sup> Corresponding author. Mailing address: Department of Microbiology, University of Illinois, Urbana, IL 61801. Phone: (217) 333-5812. Fax: (217) 244-6697. E-mail: jimlay@uiuc.edu.

TABLE 1. *E. coli* strains

Strain	Genotype	Source or reference
UM <sub>1</sub>	katE katG14 lacY rspL thi-1	14
<b>UM120</b>	katE12::Tn10 hfrH thi-1	Peter Loewen
<b>UM202</b>	$katGI7::Tn10 hfrH$ thi-1	Peter Loewen
<b>GK100</b>	$\Delta$ cydAB::cam $\Delta$ (cyoABCDE)456::kan	12
<b>KM38</b>	As UM1 plus $\Delta cydAB::cam$	$P1(GK100) \times UM1$
<b>KM39</b>	As UMI plus $\Delta$ (cyoABCDE)456:: kan	$P1(GK100) \times UM1$
SK2255	zbe-279::Tn10 thyA6 rps120 $decC1$	E. coli Genetic Stock Center
N9716	As GC4468 plus $\Delta$ oxyR::spec	Gisela Storz
AS430	$\Delta$ oxyR::spec $\Delta$ lacU169 rpsL	$P1(N9716) \times GC4468$
AB1157	$F^-$ thr-1 leuB6 proA2 his-4 thi-1 argE2 lacY1 galK2 rspL supE44 ara-14 xyl-15 mtl-1 tsx-33	10
MG1655	$F^-$ wild-type	E. coli Genetic Stock Center
JI360	As MG1655 plus katE12::Tn10	$P1(UM120) \times MG1655$
JI361	As MG1655 plus katG17::Tn10	$P1(UM202) \times MG1655$
JI362	As JI360 plus $\Delta(katE12::Tn10)1$ (Tet <sup>s</sup> )	Tet <sup>s</sup> derivative of JI360
JI364	As JI361 plus $\Delta(katGI7::Tn10)1$ (Tet <sup>s</sup> )	Tet <sup>s</sup> derivative of JI361
JI367	As JI364 plus katE12::Tn10	$P1$ (UM120) $\times$ JI364
JI370	As MG1655 plus $\triangle$ ahpCF' kan::'ahpF	$P1(MC4100\Delta ahpCF) \times MG1655$
JI372	As JI362 plus $\triangle$ ahpCF' kan::'ahpF	$P1(MC4100\Delta ahpCF) \times J1362$
JI374	As J1364 plus $\triangle$ ahpCF' kan::'ahpF	P1(MC4100 $\Delta$ ahpCF) $\times$ JI364
JI377	As J1367 plus $\triangle$ ahpCF' kan::'ahpF	P1(MC4100 $\Delta$ ahpCF) $\times$ JI367
MC4100	araD139 $\Delta(\text{arg}F-\text{lac})169$ $\lambda^-$ flhD5301 fruA25 relA1 rpsL150 $rbsR22$ deoC1	E. coli Genetic Stock Center
MC4100 $\Delta$ ahpCF	As MC4100 plus $\triangle$ ahpCF' kan::'ahpF	Gisela Storz
GS022	As MC4100 plus $\lambda$ RS45 $\Phi(katG::lacZ)$	Gisela Storz
LC70	As GS022 plus $\triangle$ ahpCF' kan::'ahpF	$P1(MC4100\Delta ahpCF) \times G5022$
LC74	As LC70 plus $\Delta$ oxyR::spec	$P1(AS430) \times LC70$
LC80	As GS022 plus katG17::Tn10	$P1(UM202) \times GSO22$
HDO3	As GS022 plus katE12::Tn10	$P1$ (UM120) $\times$ GS022
LC82	As LC80 plus $\Delta(katGI7::Tn10)2$ (Tet <sup>s</sup> )	Tet <sup>s</sup> derivative of LC80
LC84	As LC82 plus katE12::Tn10	$P1$ (UM120) $\times$ LC82
LC87	As MC4100 $\Delta$ ahpCF plus zbe-279::Tn10	$P1(SK2255) \times MC4100\DeltaahpCF$
LC89	As GS022 plus $\triangle$ ahpCF' kan::'ahpF zbe-279::Tn10	$P1(LC87) \times G5022$

foxide (DMSO) was purchased from Fisher. Water for buffers was purified with a Labconco Water Pro PS system using deionized water as the feedstock.

Luria broth (LB) contained (per liter) 10 g of Bactotryptone (Difco), 5 g of yeast extract (Difco), and 10 g of sodium chloride. To prevent the photochemical formation of hydrogen peroxide, LB medium was shielded from light and used within 24 h of its preparation. Glucose minimal medium consisted of minimal A salts (19) with 1 mM  $MgSO_4 \cdot 7H_2O$ , 5 mg of thiamine, and 2 g of glucose per liter. However, to minimize the chemical production of hydrogen peroxide, the glucose medium used in some experiments (as noted) was prepared immediately before use and contained only 0.5 g of glucose per liter. Tetracycline, kanamycin, spectinomycin, and chloramphenicol were used at 12, 40, 120, and 20  $\mu$ g/ml, respectively.

**Growth conditions and strains.** Cultures were routinely grown at 37°C. Aerobic cultures were grown in shaking flasks; anaerobic cultures were grown in a Coy chamber (Coy Laboratory Products, Inc.) under  $85\%$  N<sub>2</sub>–10% H<sub>2</sub>–5% CO<sub>2</sub>. The optical densities (OD) of all cultures were measured at 600 nm.

The strains used in this study were derived from *E. coli* K-12 and are listed in Table 1; isogenic strains were used in all experiments. Mutations were introduced into strains via P1 transduction (19). To avoid the outgrowth of suppressed strains, the *katG, katE*, *oxyR*, and Tn*10*-linked *ahp* mutations were transduced and selected under anaerobic conditions. The presence of *katG* null mutations was confirmed by an *o*-dianisidine assay for hydroperoxidase I (HPI) activity (below). *katE* mutants failed to form bubbles on plates when colonies were overlaid with a drop of 30%  $H_2O_2$ . Mutants lacking  $\alpha$ yR were identified by their inability to induce HPI when anaerobic, exponentially growing cultures at 0.1 OD in LB medium were exposed to 60  $\mu$ M H<sub>2</sub>O<sub>2</sub> for 45 min. Mutations in *gshA* were confirmed by measurements of intracellular thiol levels using DTNB (13).

Alkyl hydroperoxide reductase cannot easily be assayed in extracts, because its subunits dissociate (11, 26). However, *ahpCF* mutants formed a large zone of growth inhibition when 10  $\mu$ l of 3% cumene hydroperoxide was spotted onto a filter disk and laid onto a mutant-seeded plate (32). The excision of Tn*10* elements containing a tetracycline resistance marker was achieved at 42°C by standard methods (16).

 $H_2O_2$  **detection.** In the presence of  $H_2O_2$ , horseradish peroxidase (HRP) oxidizes AR to the fluorescent product resorufin. One milligram of AR was dissolved in 0.78 ml of DMSO, and 0.75 ml of this solution was then diluted into 18 ml of 50 mM potassium phosphate (KPi, pH 7.8) to generate a 200  $\mu$ M stock solution. This solution was shielded from light. HRP was dissolved in 50 mM KPi (pH 7.8) to 0.02 mg/ml. To measure  $H_2O_2$ , 0.45 ml of sample was mixed with 0.25 ml of AR and 0.25 ml of HRP. Fluorescence was then measured in a Shimadzu RF Mini-150 fluorometer and converted to  $H_2O_2$  concentration using a curve obtained from standard samples. Note that a small amount of  $H_2O_2$  is generated by the dye/HRP detection system itself; this amount was accounted for by the standard curve.

**H2O2 scavenging by whole cells.** Cultures were grown aerobically for at least four generations to 0.1 to 0.3 OD. Cells were pelleted in a microcentrifuge, washed twice, and resuspended in room temperature phosphate-buffered saline (PBS) at an OD of 0.1.  $H_2O_2$  was added to the appropriate final concentration (see figure legends). At intervals, 0.45-ml aliquots were removed, diluted when necessary, and assayed immediately for  $H_2O_2$  content by the AR/HRP method.

LB medium was used for most experiments. However, in experiments designed to measure the scavenging of trace  $H_2O_2$ , cultures were grown in minimal A glucose medium containing 0.5 mM each of the 20 L-amino acids. Cells were then washed and resuspended into fresh medium containing only 0.02% glucose and 0.05 mM amino acids, so that metabolism was active and could provide Ahp with reductants, yet the amount of  $H_2O_2$  generated by the medium was minimal. When cyanide was included in the medium, fluorescence development was permitted to proceed for 10 min before a final reading was made, since the carryover of cyanide inhibits the activity of HRP.

**Measurement of H<sub>2</sub>O<sub>2</sub> accumulation in cell cultures.** To detect  $H_2O_2$  formation by drug-treated cells, we exposed log-phase cultures (0.2 OD) in LB medium to 300  $\mu$ M plumbagin, an amount that is sufficient to consume 8  $\mu$ M oxygen per min in a cyanide-resistant (nonrespiratory) fashion. After 16 min, cells were removed by centrifugation, and the residual  $H_2O_2$  in the medium was assayed by the HRP/*o*-dianisidine assay (18).

 $H<sub>2</sub>O<sub>2</sub>$  can be detected with greater sensitivity in defined medium using the

AR/HRP assay. Cells were grown anaerobically overnight in minimal A medium containing 0.2% glucose, diluted to 0.01 OD, and grown anaerobically for four generations. These log-phase cells were then subcultured to 0.05 OD in fresh aerobic minimal A medium containing 0.05% glucose. At 0.05 and 0.1 OD, the culture medium was assayed for  $H_2O_2$ . Hydrogen peroxide levels were also determined in sterile medium that was incubated at 37°C for an equivalent time. The low glucose concentration was used in order to minimize  $H_2O_2$  production by salt-catalyzed glucose autooxidation (below).

**H2O2 production rates.** In order to monitor continuously the intracellular formation of  $H_2O_2$ , extracellular  $H_2O_2$  levels were measured. Cells were first grown overnight in minimal A 0.2% glucose medium containing 0.5 mM each of the 20 amino acids. Most cultures were then diluted to  $\approx 0.001$  OD and grown aerobically to an OD of  $\approx$ 0.1; however, JI377 was grown only to an OD of  $\approx$ 0.05 in order to measure  $H_2O_2$  production before growth was significantly inhibited. Cells were then washed in fresh medium containing only 0.02% glucose and 0.05 mM amino acids, resuspended at an OD of 0.1 in the same medium, and incubated with shaking at either 25 or 37°C. Glucose was added to the minimal A salts immediately before use. Aliquots were removed at intervals, and their  $H_2O_2$  content was measured. The rate of  $H_2O_2$  production was normalized to the cytoplasmic volume of the suspended cells, using a standard ratio of  $0.47 \mu$ l of internal volume per 1 ml of 1.0 OD of *E. coli* (10).

**Total catalase activity.** Aerobic exponential-phase cells were harvested at 0.5 OD, washed in cold 50 mM KPi, resuspended in 1/4 the original volume, and lysed by passage through a French press. Extracts were centrifuged at  $13,000 \times$ *g* for 20 min to remove cell debris and then stored on ice. Catalase activity was measured in a 1-ml reaction mixture containing 50  $\mu$ l of extract, 1 mM H<sub>2</sub>O<sub>2</sub>, and PBS (pH 7.3). At various time points,  $10 \mu l$  was removed, diluted 1:500 in PBS, and assayed for  $H_2O_2$  by the AR/HRP procedure.

**HPI assay.** Cultures were grown aerobically for five to six generations in LB medium. Chloramphenicol was added, and cultures were incubated for 5 min. Cells were then washed in PBS containing chloramphenicol, and cell pellets were frozen on dry ice. Within an hour the cells were resuspended in 1/10 the culture volume of 10 mM KPi buffer (pH 6.4) and lysed by sonication. Debris was centrifuged out at 13,000  $\times$  g for 20 min. HPI activity was assayed by the *o*-dianisidine method (18).

-**-Galactosidase assay.** Studies of *katG*::*lacZ* expression used a RS45 (*katG*::*lacZ*) lysogen. Cultures were grown overnight under anaerobic conditions, diluted to an OD of 0.01 in LB, and grown anaerobically to an OD of 0.1. For measurement of anaerobic expression, chloramphenicol was added and cultures were harvested. For measurement of aerobic expression, cultures were shifted into air and grown with vigorous shaking to an OD of 0.3 to 0.4. Where indicated, 13,000 U of catalase was added every 15 min. Repeated additions were necessary because catalase rapidly loses activity in cell cultures.

At harvest, cultures were centrifuged. Pellets were washed in 50 mM cold KPi buffer (pH 7.0) with special concern to remove any residual exogenous catalase. Cells were resuspended in 50 mM KPi buffer (pH 7.0) at 1/10 the culture volume and lysed by French press. Cell debris was removed by centrifugation at  $13,000 \times$ *g* for 20 min. Extracts were incubated at 28°C for 5 min before being assayed. -Galactosidase activity was assayed in a 1.2-ml reaction mixture consisting of 0.2 ml of ONPG (*o*-nitrophenyl-β-D-galactopyranoside, 4 mg/ml), extract, and Zbuffer (19) at 28°C. Change in absorption over time was monitored at 420 nm. Protein concentrations of all extracts were determined using a Coomassie dyebased assay by Pierce. All assays were performed on duplicate samples, and the values were then averaged.

**Growth experiments.** Care was taken to determine whether the poor aerobic growth of the Ahp<sup>-</sup> Kat<sup>-</sup> strain (JI377) was due to endogenous  $H_2O_2$  or  $H_2O_2$ that was chemically generated by autooxidation of the medium. Low-peroxide medium, which contains  $\approx 0.02 \mu M H_2O_2$ , was prepared by adding filter-sterilized glucose to anaerobic minimal A salts immediately before inoculation. During aerobic incubation at 37°C, this medium accumulates  $\approx$  0.04  $\mu$ M H<sub>2</sub>O<sub>2</sub> per h. JI377 was inoculated from an anaerobic overnight culture to 0.01 OD in anaerobic, low-peroxide medium and grown anaerobically to 0.1 OD. Cells were then subcultured into the same medium and shifted into air. Growth was monitored during the first 2 h, during which the vast majority of  $H_2O_2$  present in the medium was generated by the cells.

In some experiments, anaerobic MG1655 was mixed with JI377 at a 9:1 ratio in LB, the mixed culture was diluted to 0.01 OD in aerobic LB, and the growth of both strains was monitored by intermittent dilution and plating on LB plates (to quantify total viable cells) and LB plus tetracycline (to quantify viable JI377). In other experiments JI377 was shifted into aerobic LB in pure culture, and 6,500 U of catalase was added every 15 min. JI377 growth was monitored by plating, because the absorbance of catalase interferes with measurement of biomass by optical density.

**Disk diffusion.** Standing overnight cultures in LB medium were diluted to 0.005 OD in aerobic LB and then grown to an OD of  $\approx$  0.1. Cultures were diluted 1:10, and 100  $\mu$ l was spread on LB plates. Round sterile filters (1-cm diameter) were placed in the center of the plates and spotted with  $25 \mu$  of  $3\%$  H<sub>2</sub>O<sub>2</sub>. Plates were incubated at 37°C overnight. The distance from the edge of the disk to the edge of the growth zone was measured. This experiment was performed in triplicate; mean values are reported.

## **RESULTS**

Catalase is not the primary scavenger of low-level  $H_2O_2$ . Inside bacteria, antibiotics such as paraquat, plumbagin, and juglone are cyclically reduced by redox enzymes and oxidized by molecular oxygen, thereby generating superoxide and, upon its dismutation, hydrogen peroxide. The rate at which  $H_2O_2$  is made inside drug-treated *E. coli* can be determined by measurement of nonrespiratory oxygen consumption in the presence of cyanide. However, we were unable to detect  $H_2O_2$ accumulating in the medium of plumbagin-treated wild-type cells, despite the fact that oxygen consumption measurements indicated that the amount of  $H_2O_2$  should have been easily within our detection limits (data not shown).

*E. coli* contains two catalases, and we anticipated that these were responsible for scavenging the  $H_2O_2$ . However, when the experiments were repeated with strain UM1, which has point mutations in both of the catalase structural genes (*katG* and  $k \alpha t E$ ), H<sub>2</sub>O<sub>2</sub> was again undetectable. By adding 1.5  $\mu$ M H<sub>2</sub>O<sub>2</sub> to the bacterial culture, we found that UM1 degraded  $H_2O_2$  as rapidly as did its wild-type parent (data not shown).

HPI, the catalase encoded by *katG*, exhibits weak NAD(P)H-dependent peroxidase activity in vitro, and the *katG17* mutation that is present in UM1 eliminates catalase but not peroxidase activity. Therefore, we wondered whether the residual peroxidase activity of HPI was responsible for the scavenging activity of UM1. True null mutations of both *katE* and *katG* were transduced into the wild-type strain MG1655, and the  $H_2O_2$  scavenging measurements were repeated. This catalase double mutant had no detectable catalase or HPI peroxidase activity in vitro, but it still scavenged 1.5  $\mu$ M H<sub>2</sub>O<sub>2</sub> approximately as rapidly as its wild-type parent (Fig. 1). Therefore, *E. coli* must have another means of efficiently scavenging  $H_2O_2$ .

**Ahp is the source of scavenging in a catalase null mutant.** Other mechanisms of scavenging were considered. Some  $\alpha$ -ketoacids, such as pyruvate, can be excreted into the medium of glucose-fed cells, and these can chemically reduce  $H_2O_2$  in an oxidative decarboxylation reaction (4). In separate work we have observed that both the respiratory cytochrome *o* and cytochrome *d* oxidases have weak peroxidase activities (A. Nguyen and J. A. Imlay, submitted for publication). Finally, alkyl hydroperoxide reductase (Ahp) has been shown to accept  $H<sub>2</sub>O<sub>2</sub>$  as a substrate in vitro (22).

Spent medium was examined, but it did not have significant scavenging activity. Cyanide competitively inhibits  $H_2O_2$  binding by cytochrome oxidases but did not diminish scavenging by a *katG katE* mutant; the same result was obtained when *cyo* and *cyd* mutations were placed in this background. Thus, neither growth medium nor cytochrome oxidases provided the catalase-independent scavenging activity.

Ahp is an NAD(P)H-dependent peroxidase that rapidly reduces organic hydroperoxides as diverse as cumene and *t*-



FIG. 1. Ahp scavenges  $H_2O_2$  in a Kat<sup>-</sup> strain. Cultures of MG1655 (wild type, ∇), JI367 (*katG katE*, ◇), JI370 (*ahpCF*, \*), JI377 (*ahpCF*  $k$ at*G* katE,  $\circ$ ), JI374 (*ahpCF katG*,  $\Box$ ), and JI372 (*ahpCF katE*,  $\triangle$ ) were grown aerobically in LB and resuspended in PBS at an OD of 0.1.  $H<sub>2</sub>O<sub>2</sub>$  was added at a final concentration of 1.5  $\mu$ M. At various time points after addition of  $H_2O_2$ , the  $H_2O_2$  concentration was measured as described in Materials and Methods.

butylhydroperoxide (11). Niimura et al. subsequently found some activity with hydrogen peroxide as the substrate, although we are not aware of any comparative study of turnover numbers (22). Ahp, like HPI, is a member of the OxyR regulon in diverse bacteria (15, 20, 24). OxyR is activated by organic hydroperoxides as well as by  $H_2O_2$ , and it has become accepted that the physiological role of Ahp is to scavenge organic hydroperoxides. Mutants that lack Ahp are indeed hypersensitive to growth inhibition by organic hydroperoxides, which are not substrates for catalase (32).

A null mutant lacking Ahp scavenged  $H_2O_2$  as well as did its wild-type parent (Fig. 1). However, in contrast to the *katG katE* strain, an *ahpCF katG katE* triple mutant exhibited virtually no scavenging activity. An *ahpCF katG* mutant also failed to scavenge  $H_2O_2$ , although an *ahpCF katE* mutant did so (Fig. 1). Thus, Ahp provides the scavenging activity that persists in catalase null mutants.

**HPI induction can compensate for loss of Ahp.** We wished to learn which enzyme, Ahp or HPI, was the predominant scavenger in wild-type cells. Because the *ahpCF* and *katG* single mutants each scavenged  $H_2O_2$  at the same rate as did wild-type cells (Fig. 1), we inferred that one of the enzymes might be induced to compensate for the absence of the other. Both *ahpCF* and *katG* are positively regulated by the OxyR regulon, and a mutation in the constitutive scavenger might cause intracellular  $H_2O_2$  to accumulate until OxyR activated the expression of the other, ultimately restoring wild-type levels of  $H_2O_2$  scavenging.

We observed that an *ahpCF* mutant contained sevenfold more total catalase (including both HPI and HPII) than did wild-type cells. The *o*-dianisidine peroxidase activity, which specifically reflects the HPI titer, was increased 10-fold. Similarly, the  $\beta$ -galactosidase activity of an aerobic  $\lambda$ RS45

TABLE 2. Endogenous  $H_2O_2$  activates the OxyR regulon in an *ahpCF* mutant

Strain <sup>a</sup>	Mean $\beta$ -galactosidase activity (U/mg of protein) $\pm$ SD		
	Anaerobic	Aerobic	Aerobic $+$ catalase <sup>b</sup>
Ahp <sup>+</sup> OxyR <sup>+</sup> (GSO22) Ahp <sup>-</sup> OxyR <sup>+</sup> (LC70) Ahp <sup><math>-</math></sup> OxyR $-$ (LC74)	$0.03 \pm 0.01$ $0.03 \pm 0.01$ $0.03 \pm 0.01$	$0.03 \pm 0.01$ $0.35 \pm 0.03$ $0.03 \pm 0.01$	$0.03 \pm 0.01$ $0.18 \pm 0.02$ ND <sup>c</sup>

*a* Strains were isogenic and harbored  $\lambda$ RS45 (*katG*::*lacZ*). *b* Exogenous catalase (30 µl) was added before the culture was aerated and then every 15 min until it was iced. *<sup>c</sup>* ND, not determined.

*katG*::*lacZ* lysogen was increased 10-fold when the *ahpCF* null mutation was introduced, rising from 0.04  $\pm$  0.01 to 0.39  $\pm$ 0.05  $\beta$ -galactosidase U/mg. In contrast, the introduction of *katG* (LC80), *katE* (HDO3), or both *katG* and *katE* (LC84) mutations did not increase  $\beta$ -galactosidase activity at all  $(0.04 \pm 0.01, 0.03 \pm 0.01,$  and  $0.04 \pm 0.01$  U/mg, respectively). These results agreed with the observations of Rosner and Storz (27) and Ritz et al. (25a).

Induction of *katG* in the *ahpCF* mutant was blocked by an *oxyR* mutation (Table 2). To ensure that induction of the OxyR regulon was not due to a suppressor mutation secondary to the *ahpCF* mutation, the *ahpCF* null allele was retransduced into the *katG*::*lacZ* lysogen under anaerobic conditions. Once again, the fusion was induced when the transductant was cultured in aerobic medium (data not shown). These data demonstrate that in wild-type cells Ahp rather than catalase is the primary scavenger of an endogenous inducer of the OxyR regulon.

The induction of *katG*::*lacZ* in *ahp* cultures was partially averted by the addition to the growth medium of exogenous catalase (Table 2). Since catalase scavenges only  $H_2O_2$ , we infer that  $H_2O_2$  is the species that accumulates in these cultures and activates the OxyR regulon. The extracellular catalase may have failed to completely block induction both because of its instability and because extracellular scavengers cannot fully eradicate intracellular  $H_2O_2$  accumulation (30). Alternatively, it is possible that an additional inducer was present.

Ahp and HPI have discrete roles in scavenging  $H_2O_2$ . Mutants that contain only one or the other scavenger exhibited very different dose-response curves (Fig. 2). Whole cells that contained only Ahp scavenged low concentrations of  $H_2O_2$ very effectively, exhibiting a half-maximal rate when the extracellular concentration of  $H_2O_2$  was about 5  $\mu$ M. The activity became saturated when extracellular  $H_2O_2$  exceeded 20  $\mu$ M (Fig. 2). In contrast, HPI-expressing cells were not saturated by even millimolar concentrations of  $H_2O_2$ , consistent with its  $K_m$ of 5.9 mM (9). The dose-response curves suggest that the primary role of HPI might be to scavenge higher concentrations of  $H_2O_2$ , against which Ahp is ineffective.

To directly contrast the kinetic behaviors of Ahp and HPI, we measured the rates at which cells decomposed low  $(0.1 \mu M)$ and high (150  $\mu$ M) concentrations of H<sub>2</sub>O<sub>2</sub>. (To simplify interpretation, the strains used in these experiments lacked HPII. However, control experiments demonstrated that HPII



FIG. 2. Dependence of scavenging rate on  $H_2O_2$  concentration. Rates of  $H_2O_2$  decomposition were measured in dilute suspensions of JI370 (ahp $\overline{CF}$ ,  $\Diamond$ ) and JI367 (katG katE,  $\Box$ ). Rates were normalized to a value of 1.0 OD.

provides significant scavenging activity only in stationary phase, when it is induced by RpoS [28].) An  $HPI^{-}$  Ahp<sup>+</sup> mutant scavenged 0.1  $\mu$ M H<sub>2</sub>O<sub>2</sub> slightly more rapidly than did an  $HPI^{+}$  Ahp<sup>-</sup> mutant, despite the fact that the latter strain has 10-fold-induced levels of HPI (Fig. 3, left panel). Thus, in wild-type *E. coli* most scavenging of low-dose  $H_2O_2$  must be done by Ahp. Conversely, the HPI<sup>-</sup> Ahp<sup>+</sup> strain was poor at degrading 150  $\mu$ M H<sub>2</sub>O<sub>2</sub>, unlike the HPI<sup>+</sup> Ahp<sup>-</sup> and HPI<sup>+</sup>  $A$ hp<sup>+</sup> strains (Fig. 3, right panel). Although Ahp can be inactivated by  $H_2O_2$  in vitro (11), its poor activity in vivo seemed not

TABLE 3. HPI (*katG*) is the primary scavenger of supranormal H2O2 concentrations*<sup>a</sup>*

Strain	Zone of inhibition $^b$ (cm)

<sup>*a*</sup> Assay was done by disk diffusion of 25  $\mu$ l of 3% (880 mM) H<sub>2</sub>O<sub>2</sub>. Assay was done in triplicate, and trends were always consistent. One data set is shown. Strains used were MG1655 (wild type), JI364 (*katG*), JI370 (*ahpCF*), JI367 (*katG*

*k*<sub>*b*</sub> Distance of clearing from the edge of the disk.

to stem from this problem, since normal scavenging activity was observed when the 150  $\mu$ M H<sub>2</sub>O<sub>2</sub> was washed away and the cells were exposed to 1.5  $\mu$ M H<sub>2</sub>O<sub>2</sub> (data not shown). The Ahp<sup>-</sup> mutant scavenged 150  $\mu$ M H<sub>2</sub>O<sub>2</sub> twice as rapidly as did the wild type, because  $HPI^+$  was induced (30).

These data indicate that Ahp and HPI have distinct roles: Ahp is more effective at scavenging very low concentrations of  $H<sub>2</sub>O<sub>2</sub>$ , while HPI is the more effective enzyme at higher concentrations. The dominance of HPI at supranormal levels of  $H<sub>2</sub>O<sub>2</sub>$  was also apparent in disk diffusion assays, which test the ability of strains to degrade high concentrations of  $H_2O_2$  to a level that permits growth. The *katG* mutant was hypersensitive, while the *ahpCF* mutant showed wild-type resistance (Table 3). The *ahpCF* mutation debilitated only strains that lacked HPI. As before, the addition of a *katE* mutation did not affect the sensitivity of any strain.

Calculation of endogenous  $H_2O_2$  production during aerobic **growth.** Substantial  $H_2O_2$  accumulated in the medium of  $Ahp^ \text{Kat}^-$  cultures. The extracellular concentration of  $\text{H}_2\text{O}_2$  rose to



FIG. 3. Distinct efficiencies of Ahp and HPI at different H<sub>2</sub>O<sub>2</sub> concentrations. H<sub>2</sub>O<sub>2</sub> was added at a final concentration of 0.1  $\mu$ M (right panel) and 150  $\mu$ M (left panel) to cultures of JI372 (*ahpCF katE*), JI367 (*katG katE*), and JI362 (*katE*). Two minutes after addition of H<sub>2</sub>O<sub>2</sub>, the H<sub>2</sub>O<sub>2</sub> concentration was measured as described in Materials and Methods.



FIG. 4.  $H_2O_2$  production by Ahp<sup>-</sup> Kat<sup>-</sup> cells. MG1655 (wild type), JI370 (*ahpCF*), JI367 (*katG katE*), and JI377 (*ahpCF katG katE*) were grown aerobically in LB and resuspended in 37°C minimal A salts containing 0.2% glucose. At various time points after resuspension, the  $H<sub>2</sub>O<sub>2</sub>$  concentration of the medium was measured. (The  $H<sub>2</sub>O<sub>2</sub>$  levels drop for the three scavenger-proficient strains because these strains degrade the 0.05  $\mu$ M H<sub>2</sub>O<sub>2</sub> that is present in the initial medium.)

1.8  $\mu$ M during aerobic growth in minimal 0.05% glucose medium when cells were cultured for a single generation, from 0.05 to 0.10  $OD<sub>600</sub>$ . In contrast, the H<sub>2</sub>O<sub>2</sub> concentrations of wild-type (MG1655),  $Kat^-$  (JI367), and  $Ahp^-$  (JI370) cultures were below our detection limit of  $\approx 0.04 \mu M H_2O_2$ . The fact that the  $H_2O_2$  concentration was so low in Ahp<sup>-</sup> cultures was initially surprising, given that sufficient  $H_2O_2$  was present to activate the OxyR protein. However, subsequent work indicated that the concentration of  $H_2O_2$  inside these cells may have been substantially higher than that outside them (30).

Kat<sup>-</sup> Ahp<sup>-</sup> strains have <5% of the scavenging activity of wild-type cells, so that virtually all of the  $H_2O_2$  that enters or is formed within these cells diffuses out without being scavenged (30). For this reason the measurement of excreted  $H_2O_2$ is a valid proxy for measurement of endogenous  $H_2O_2$  formation. Using this strain, we quantitated the rate at which *E. coli* generates  $H_2O_2$  (Fig. 4). Measurements were made after exponentially growing cells were suspended in fresh glucose/ amino acids medium at 37°C (and do not necessarily apply in other media). The rate of  $H_2O_2$  formation, normalized to cell volume, was 14  $\mu$ M H<sub>2</sub>O<sub>2</sub>/s at 37°C. This result is consistent with earlier predictions (Discussion).

**Inability to scavenge endogenous**  $H_2O_2$  **causes a growth defect.** It has long been suspected that aerobic metabolism generates enough  $H_2O_2$  to damage cells that cannot scavenge it. No growth defects were apparent in the *ahpCF* mutant; we presume this is because the cell compensates sufficiently for the loss of *ahpCF* by inducing  $k \cdot \text{ad}$ . However, the Ahp<sup>-</sup> Kat<sup>-</sup> strain grew poorly in all aerobic media that were tested (Fig. 5). Growth slowed progressively over time, and when cells were repeatedly subcultured, growth often stopped entirely (Fig. 5B). This may reflect the continual accumulation of damage in the cell. Growth was particularly poor on defined media that

lacked amino acids (data not shown). Wild-type growth was restored when catalase was added to the medium (Fig. 5C) or when the mutant was cultured anaerobically (not shown).

Special efforts were undertaken to confirm that the growth defect was due to endogenous  $H_2O_2$  rather than  $H_2O_2$  made by autooxidation of the glucose medium. Cells were cultured anaerobically in glucose medium to log phase and then diluted into fresh aerobic "peroxide-free" medium (Fig. 5D). Within the first half-hour the Ahp<sup>-</sup> Kat<sup>-</sup> strain grew more poorly than did its wild-type parent. Our measurements over this period confirmed that the  $H_2O_2$  found in this culture was generated primarily by the cells rather than by glucose oxidation. Thus, aerobic *E. coli* generates enough  $H_2O_2$  to debilitate the cell. In previous studies of catalase mutants, the toxicity of endogenous  $H_2O_2$  was obscured by the scavenging action of Ahp.

The necessity for care in these experiments was underscored by the observation that medium that had been stored on the bench often contained micromolar amounts of  $H_2O_2$ , and dilution of even wild-type cells into stored medium was sufficient to transiently induce the OxyR regulon (data not shown).

# **DISCUSSION**

**H2O2 may be the sole physiological substrate of Ahp.** The results of this study indicate that in exponentially growing *E. coli* Ahp is responsible for the degradation of low concentrations of hydrogen peroxide. Previously, Ahp was primarily associated with the detoxification of organic hydroperoxides. However, the range of organic peroxides that are good substrates for the enzyme indicates that its active site can accommodate virtually any ROOH, so it is not surprising that it reacts with HOOH as well. Earlier genetic studies confirmed that Ahp provides cellular resistance to organic hydroperoxides but found little role in resistance to  $H_2O_2$ . This result is now understandable: because catalases cannot degrade organic hydroperoxides, Ahp is likely to be the only enzyme in *E. coli* with that catalytic ability. At the same time, the role of Ahp in  $H_2O_2$ decomposition was obscured because the widely employed disk inhibition assays confront cells with high  $H_2O_2$  concentrations that are more efficiently decomposed by catalases.

It is not clear whether *E. coli* must ever detoxify organic hydroperoxides in nature. Lipid hydroperoxides were suggested to be the physiological substrates for Ahp, but *E. coli* lacks the polyunsaturated fatty acids that appear to be necessary for lipid peroxidation (1), and we are not aware of any study that has documented the recovery of peroxidized lipids from this bacterium. It is possible that organic hydroperoxides are pseudosubstrates for an enzyme whose only role in nature is the decomposition of hydrogen peroxide.

**Why does** *E. coli* **use multiple scavengers?** In general, all peroxidases will be inferior to catalases at scavenging high concentrations of  $H_2O_2$  because peroxidases can turn over only as quickly as the cell can provide a reductant to them. Catalases escape this restriction. For example, the HPI catalase of Ahp<sup>-</sup> *E. coli* degraded 500  $\mu$ M H<sub>2</sub>O<sub>2</sub> at a rate of 7  $\times$  10<sup>7</sup> molecules cell<sup>-1</sup> s<sup>-1</sup>. To achieve the same rate using Ahp would require an equivalent amount of NADH, which substantially exceeds the rate at which metabolism generates NADH. (Glucose-saturated cells can generate enough NADH to respire  $6 \times 10^6$  molecules of oxygen cell<sup>-1</sup> s<sup>-1</sup>.) The disparity is



FIG. 5. An Ahp<sup>-</sup> Kat<sup>-</sup> strain has an aerobic growth defect. (A) Growth of MG1655 (wild type,  $\blacksquare$ ), JI370 (*ahpCF*,  $\bigcirc$ ), JI367 (*katG katE*,  $\Diamond$ ), JI374 (*katG ahpCF*, \*), and JI377 (*katG katE ahpCF*,  $\nabla$ ) in aerobic LB medium. (B) MG1655 and JI377 were grown aerobically in LB from 0.001 OD to mid-log phase, as for panel A. Cells were then subcultured into fresh LB at an OD of 0.01, and residual growth was observed. (C) Exogenous catalase protects against an aerobic growth defect in LB. MG1655 and JI377 were grown aerobically in fresh LB. Cultures were then subcultured to 0.01 OD in LB. Exogenous catalase was added to one culture of JI377 every 15 min to maintain catalase activity. At various time points, aliquots were removed from each culture and plated in selective top agar. Growth rate was determined the next day. (D) Endogenous  $H_2O_2$  can be toxic to cells. Exponential anaerobic MG1655 and JI377 were subcultured into fresh aerobic peroxide-free minimal A glucose (0.2%) medium, and growth was monitored.

even greater when one considers decomposition of higher concentrations of  $H_2O_2$ . Therefore, catalases allow cells to degrade high concentrations of  $H_2O_2$  far more quickly than would peroxidases alone.

The preference for Ahp at low  $H_2O_2$  concentrations may derive from its greater catalytic efficiency. Since normal intracellular concentrations of  $H_2O_2$  are well below the  $K_m$  of both enzymes,  $k_{\text{cat}}/K_m$  is the relevant kinetic parameter. The  $k_{\text{cat}}/K_m$ of the catalase activity of HPI is  $9 \times 10^5$  M<sup>-1</sup> s<sup>-1</sup> (9); that of Ahp has not been reported, but the data of Niimura et al. (22) imply that it is at least  $8 \times 10^6$  M<sup>-1</sup> s<sup>-1</sup>. The implication is that at least 10-fold more molecules of HPI need to be synthesized to provide the scavenging activity provided by Ahp. A second disadvantage to catalase may stem from the fact that enzymes that require reactions with two molecules of substrate to complete a catalytic cycle can have difficulty at low substrate concentrations, when an intermediate state is long-lived. Compound I, the divalently oxidized catalase intermediate, can be unstable and in some circumstances may reversibly deactivate (9), which would further diminish the ability of catalase to scavenge trace amounts of  $H_2O_2$ . Peroxidases may comprise an evolutionary solution to this problem.

The kinetic efficiency of HPII ( $k_{\text{cat}}/K_m = 1.3 \times 10^6 \text{ M}^{-1} \text{ s}^{-1}$ ) (23), the stationary-phase catalase, is similar to that of HPI. It seems reasonable that *E. coli* would increase its catalase titer in stationary phase, when there may not be sufficient NADH for Ahp to remain an effective scavenger. Gonzalez-Flecha and Demple reported that both catalases are induced as growth slows (8). It is not clear why scavenging efficiency could not be maintained by induction of HPI alone. For now, the benefit to *E. coli* of having two catalases instead of one remains obscure.

The compensatory interactions that we observed between catalase and Ahp synthesis have been observed previously in a wide range of bacteria. In *Xanthomonas campestris, Bacteroides fragilis,* and *Pseudomonas aeruginosa*, mutations in *ahp* cause catalase induction (20, 24, 26); this may be true in *Bacillus subtilis* as well (2). Conversely, *katG* mutations in *Mycobacterium tuberculosis* select for promoter upmutations in *ahp* (31). It seems likely that the division of labor found in *E. coli*, wherein Ahp scavenges low levels of  $H_2O_2$  and catalase scavenges high levels, is widespread.

**Mechanisms of**  $H_2O_2$  **formation and cell damage.** The existence of a strain unable to scavenge the  $H_2O_2$  it produces has enabled us to measure the rate at which  $H_2O_2$  is formed. In earlier studies in vitro, we determined that the respiratory chain was likely to be the primary source of  $H_2O_2$  during aerobic growth on glucose, largely because of the autooxidizability of the NADH dehydrogenase II (18). This enzyme adventitiously transfers electrons to oxygen from its reduced flavin in a direct second-order chemical reaction. Other flavoenzymes can do so as well, and in circumstances where these other enzymes are especially abundant, they may be the predominant sources of superoxide and  $H_2O_2$ .

The rate at which  $H_2O_2$  is likely to be formed in vivo can be predicted from the in vitro data. We found that 9 molecules of  $H<sub>2</sub>O<sub>2</sub>$  were formed per 10,000 electrons flowing through the NADH dehydrogenase II (18). We do not know the fraction of the respiratory flux that flows through this enzyme during growth on glucose, but given that exponentially growing cells consume about 3.2 mM  $O_2/s$ , one extrapolates that the rate of  $H_2O_2$  formation could be up to 12  $\mu$ M/s. This is close to the rate determined in vivo in this study (14  $\mu$ M/s). Efforts to test the sources of  $H_2O_2$  production in vivo are under way.

Gonzalez-Flecha and Demple inferred rates of  $H_2O_2$  production by intact, wild-type cells (ca. 1 to 2  $\mu$ M/s) that are lower than those that we have reported (8). However, we observed rates similar to theirs when we suspended cells in room temperature buffer, as they did  $(3 \mu M/s)$ . It is not surprising that  $H_2O_2$  is most rapidly formed when metabolism is active and temperatures are high enough to overcome the activation energy (18). There was a procedural difference, however. Their calculation was based on measurements of steadystate concentrations of extracellular  $H_2O_2$  in suspensions of cells: by assuming that catalase was the predominant scavenger of  $H_2O_2$ , they used measurements of its activity to calculate the rate of  $H_2O_2$  production. In our experiments we found that  $H_2O_2$  did not accumulate extracellularly due to the action of Ahp. These differences may reflect the different conditions under which the experiments were performed.

Inside growing cells, the steady-state concentration of  $H_2O_2$ depends on the rates of its formation and of its dissipation. In this study we quantified the rate at which  $H_2O_2$  is formed when *E. coli* is cultured under a particular set of growth conditions. In the accompanying work (30), we measured the processes that consume  $H_2O_2$  and estimated the internal  $H_2O_2$  concentration.

### **ACKNOWLEDGMENTS**

We are grateful to Gigi Storz, Peter Loewen, Bruce Demple, Bob Gennis, and Jim Slauch for providing strains and discussion that helped us in this study, and we thank Holly Oliver for assistance with strain constructions.

This work was supported by grant GM49640 from the National Institutes of Health.

#### **REFERENCES**

- 1. **Bielski, B. H. J., R. L. Arudi, and M. W. Sutherland.** 1983. A study of the reactivity of  $HO_2^-/O_2^-$  with unsaturated fatty acids. J. Biol. Chem. 258: 4759–4761.
- 2. **Bsat, N., L. Chen, and J. Helmann.** 1996. Mutation of the *Bacillus subtilis* alkyl hydroperoxide reductase (*ahpCF*) operon reveals compensatory interactions among hydrogen peroxide stress genes. J. Bacteriol. **178:**6579–6586.
- 3. **Carlioz, A., and D. Touati.** 1986. Isolation of superoxide dismutase mutants in *Escherichia coli*: is superoxide dismutase necessary for aerobic life? EMBO J. **5:**623–630.
- 4. **Constantopoulos, G., and J. Barranger.** 1984. Nonenzymatic decarboxylation of pyruvate. Anal. Biochem. **139:**353–358.
- 5. Dukan, S., and T. Nyström. 1999. Oxidative stress defense and deterioration of growth-arrested *Escherichia coli* cells. J. Biol. Chem. **274:**26027–26032.
- 6. **Gaudu, P., D. Touati, V. Niviere, and M. Fontecave.** 1994. The NAD(P)Hflavin oxidoreductase from *Escherichia coli* as a source of superoxide radicals. J. Biol. Chem. **269:**8182–8188.
- 7. **Gonzalez-Flecha, B., and B. Demple.** 1995. Metabolic sources of hydrogen peroxide in aerobically growing *Escherichia coli*. J. Biol. Chem. **1995:**13681– 13687.
- 8. **Gonzalez-Flecha, B., and B. Demple.** 1997. Homeostatic regulation of intracellular hydrogen peroxide concentration in aerobically growing *Escherichia coli*. J. Bacteriol. **179:**382–388.
- 9. **Hillar, A., B. Peters, R. Pauls, A. Loboda, H. Zhang, A. G. Mauk, and P. C. Loewen.** 2000. Modulation of the activities of catalase-peroxidase HPI of *Escherichia coli* by site-directed mutagenesis. Biochemistry **59:**5868–5875.
- 10. **Imlay, J. A., and I. Fridovich.** 1991. Assay of metabolic superoxide production in Escherichia coli. J. Biol. Chem. **266:**6957–6965.
- 11. **Jacobson, F. S., R. W. Morgan, M. F. Christman, and B. N. Ames.** 1989. An alkyl hydroperoxide reductase from *Salmonella typhimurium* involved in the defense of DNA against oxidative damage: purification and properties. J. Biol. Chem. **264:**1488–1496.
- 12. **Kaysser, T. M., J. B. Ghaim, C. Georgiou, and R. B. Gennis.** 1995. Methionine-393 is an axial ligand of the heme b558 component of the cytochrome *bd* ubiquinol oxidase from *Escherichia coli*. Biochemistry **34:**13491–13501.
- 13. **Lawley, P. D., and C. J. Thatcher.** 1970. Methylation of deoxyribonucleic acid in cultured mammalian cells by *N*-methyl-*N*-nitro-*N*-nitrosoguanidine: the influence of cellular thiol concentrations. Biochem. J. **116:**693–707.
- 14. **Loewen, P. C.** 1984. Isolation of catalase-deficient *Escherichia coli* mutants and genetic mapping of *katE*, a locus that affects catalase activity. J. Bacteriol. **157:**622–626.
- 15. **Loprasert, S., M. Fuangthong, W. Whangsuk, A. Atichartpongkul, and S. Mongkolsuk.** 2000. Molecular and physiological analysis of an OxyR-regulated *ahpC* promoter in *Xanthomonas campestris* pv. phaseoli. Mol. Microbiol. **37:**1504–1514.
- 16. **Maloy, S. R., and W. D. Nunn.** 1981. Selection for loss of tetracycline resistance by *Escherichia coli*. J. Bacteriol. **145:**1110–1112.
- 17. **McCord, J. M., B. B. Keele, Jr., and I. Fridovich.** 1971. An enzyme-based theory of obligate anaerobiosis: the physiological function of superoxide dismutase. Proc. Natl. Acad. Sci. USA **68:**1024–1027.
- 18. **Messner, K. R., and J. A. Imlay.** 1999. The identification of primary sites of superoxide and hydrogen peroxide formation in the aerobic respiratory chain and sulfite reductase complex of *Escherichia coli*. J. Biol. Chem. **274:**10119– 10128.
- 19. **Miller, J. H.** 1972. Experiments in molecular genetics*.* Cold Spring Harbor Laboratory, Cold Spring Harbor, N.Y.
- 20. **Mongkolsuk, S., W. Whangsuk, P. Vattanaviboon, S. Loprasert, and M. Fuangthong.** 2000. A *Xanthomonas* alkyl hydroperoxide reductase subunit C (*ahpC*) mutant showed an altered peroxide stress response and complex regulation of the compensatory response of peroxide detoxification enzymes. J. Bacteriol. **182:**6845–6849.
- 21. **Moskovitz, J., M. A. Rahman, J. Strassman, S. O. Yancey, S. R. Kushner, N. Brot, and H. Weissbach.** 1995. *Escherichia coli* peptide methionine sulfoxide reductase gene: regulation of expression and role in protecting against oxidative damage. J. Bacteriol. **177:**502–507.
- 22. **Niimura, Y., L. B. Poole, and V. Massey.** 1995. *Amphibacillus xylanus* NADH oxidase and *Salmonella typhimurium* alkyl-hydroperoxide reductase flavoprotein components show extremely high scavenging activity for both alkyl

hydroperoxide and hydrogen peroxide in the presence of *S. typhimurium* alkyl-hydroperoxide reductase 22-kDa protein component. J. Biol. Chem. **270:**25645–25650.

- 23. **Obinger, C., M. Maj, P. Nicholls, and P. Loewen.** 1997. Activity, peroxide compound formation, and heme d synthesis in *Escherichia coli* HPII catalase. Arch. Biochem. Biophys. **342:**58–67.
- 24. **Ochsner, U. A., M. L. Vasil, E. Alsabbagh, K. Parvatiyar, and D. J. Hassett.** 2000. Role of the *Pseudomonas aeruginosa oxyR-recG* operon in oxidative stress defense and DNA repair: OxyR-dependent regulation of the *katBankB*. ahpB, and *ahpC-ahpF*. J. Bacteriol. **182:**4533–4544.
- 25. **Reynolds, C. M., and L. B. Poole.** 2001. Activity of one of two engineered heterodimers of AhpF, the NADH:peroxiredoxin oxioreductase from *Salmonella typhimurium*, reveals intrasubunit electron transfer between domains. Biochemistry **40:**3912–3919.
- 25a.**Ritz, D., H. Patel, B. Doan, M. Zheng, F. Aslund, G. Storz, and J. Beckwith.** 2000. Thioredoxin 2 is involved in the oxidative stress response in *Escherichia coli*. J. Biol. Chem. **275:**2505–2512.
- 26. **Rocha, E. R., and C. J. Smith.** 1999. Role of the alkyl hydroperoxide reductase (*ahpCF*) gene in oxidative stress defense of the obligate anaerobe

*Bacteroides fragilis*. J. Bacteriol. **181:**5701–5710.

- 27. **Rosner, J. L., and G. Storz.** 1994. Effects of peroxides on susceptibilities of *Escherichia coli* and *Mycobacterium smegmatis* to isoniazid. Antimicrob. Agents Chemother. **38:**1829–1833.
- 28. **Sak, B. D., A. Eisenstark, and D. Touati.** 1989. Exonuclease III and the catalase hydroperoxidase II in *Escherichia coli* are both regulated by the katF gene product. Proc. Natl. Acad. Sci. USA **86:**3271–3275.
- 29. **Schellhorn, H. E., and H. M. Hassan.** 1988. Response of hydroperoxidase and superoxide dismutase deficient mutants of *Escherichia coli* K-12 to oxidative stress. Can J. Microbiol **34:**1171–1176.
- 30. **Seaver, L. C., and J. A. Imlay.** 2001. Hydrogen peroxide fluxes and compartmentalization inside growing *Escherichia coli*. J. Bacteriol. **183:**7182–7189.
- 31. **Sherman, D. R., K. Mdluli, M. J. Hickey, T. M. Arain, S. L. Morris, C. E. Barry III, and C. K. Stover.** 1996. Compensatory *ahpC* gene expression in isoniazid-resistant *Mycobacterium tuberculosis*. Science **272:**1641–1643.
- 32. **Storz, G., F. S. Jacobson, L. A. Tartaglia, R. W. Morgan, L. A. Silveira, and B. N. Ames.** 1989. An alkyl hydroperoxide reductase induced by oxidative stress in *Salmonella typhimurium* and *Escherichia coli*: genetic characterization and cloning of *ahp*. J. Bacteriol. **171:**2049–2055.