Trade-Off between Iron Uptake and Protection against Oxidative Stress: Deletion of cueO Promotes Uropathogenic Escherichia coli Virulence in a Mouse Model of Urinary Tract Infection

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The periplasmic multicopper oxidase (CueO) is involved in copper homeostasis and protection against oxidative stress. Here, we show that the deletion of cueO in uropathogenic Escherichia coli increases its colonization of the urinary tract despite its increased sensitivity to hydrogen peroxide. The cueO deletion mutant accumulated iron with increased efficiency compared to its parent strain; this may account for its advantage in the iron-limited environment of the urinary tract.

Urinary tract infections (UTI) are among the most common infectious diseases of humans and a major cause of morbidity. It is estimated that 40 to 50% of healthy adult women have experienced at least one UTI episode in their lifetime (6). Uropathogenic Escherichia coli (UPEC) is the cause of the majority (>80%) of UTI in humans. UPEC isolates exhibit a high degree of genetic diversity due to the possession of specialized virulence genes located on pathogenicity islands (21). Although no single virulence factor is uniquely definitive of UPEC, its ability to cause symptomatic UTI is enhanced by adhesins (e.g., type 1 and P fimbriae) and toxins (e.g., hemolysin) (16, 38). Another important virulence property of UPEC is its ability to sequester iron. The concentration of soluble iron is very low in urine and represents an important growth-limiting factor for bacteria. UPEC possesses multiple mechanisms to acquire iron, including the production of siderophores, such as aerobactin and enterobactin (and the glycosylated enterobactin derivative, salmochelin), and the direct utilization of host iron compounds (particularly heme or hemoglobin) (1, 3, 10, 23, 25, 26). UPEC mutants deleted in these processes display reduced virulence in the mouse urinary tract (32).

The global oxidative stress response regulator OxyR is required for virulence in a mouse model of UTI (14). This indicates that UPEC responds to oxidative stress during infection, consistent with evidence that the attachment of UPEC to the uroepithelium leads to neutrophil recruitment (11, 14). OxyR also acts in concert with Dam methyltransferase to regulate the expression of the antigen 43-encoding flu gene (9, 12, 28). Antigen 43 is an autotransporter protein that promotes aggregation, biofilm formation, and long-term persistence of UPEC in the urinary bladder (15, 36).

The cueO gene encodes a periplasmic multicopper oxidase which is known to be involved in copper homeostasis and protection against oxidative stress. CueO possesses ferrooxidase (Fe2+→Fe3+), cuprous oxidase (Cu+→Cu2+), and polyphenol oxidase (oxidation of phenolic compounds, including enterobactin) activities (8, 24, 30). Both Cu2+ and Fe2+ generate toxic hydroxyl radicals via the Fenton reaction (Fe2+ or Cu2++H2O2→Fe3++OH+ + OH−). Oxidation of enterobactin by CueO prevents Cu2+ reduction by the reactive catechol groups on enterobactin and has been proposed to form a 2-carboxymuconate derivative in the periplasm that may sequester both copper and iron ions (8) and thereby protect bacteria against metal ion-promoted oxidative stress. In view of the role of CueO in E. coli K-12, we were interested in determining whether it contributes to the pathogenesis of the prototypical UPEC strain CFT073 (19, 37).

CueO is required for copper resistance in UPEC CFT073. The deletion of cueO renders E. coli K-12 sensitive to CuSO4, a phenotype that is enhanced under conditions of iron limitation in which high concentrations of enterobactin are produced. To assess the role of CueO from E. coli CFT073 in copper tolerance, a cueO deletion strain (CFT073cueO) was constructed by λ red-mediated homologous recombination, as previously described (5). Copper sensitivity was assessed by growing CFT073 and CFT073cueO on Tris-buffered mineral salts agar supplemented with 0.2% glycerol and 0.3% CAS amino acids in the presence of filter discs impregnated with 5 μl of 1 M CuSO4. In this assay, CFT073 was resistant to copper and produced a distinct brown pigment in the region of growth at the periphery of the clearing zone (Fig. 1). In contrast, CFT073cueO was highly sensitive to copper. The copper resistance phenotype of CFT073cueO could be restored by complementation with a plasmid containing the cueO gene (pCueO) (Fig. 1).

Deletion of cueO promotes colonization of the mouse bladder and shedding in urine. An established mouse model of UTI was employed to examine the role of cueO in UPEC virulence (36). Briefly, female C57BL/6 mice (8 to 10 weeks old) were catheterized using a sterile Teflon catheter by inserting the device directly into the bladder through the urethra. An inoculum of 25 μl, containing 5 × 107 CFU of CFT073 or
CFT073cueO in phosphate-buffered saline (PBS), was instilled directly into the bladder by using a 1-ml tuberculin syringe attached to the catheter. Mice were euthanized at 18 h after challenge by cervical dislocation; bladders and kidneys were then excised aseptically, weighed, and homogenized in PBS for colony counts. Urine samples were also collected from each mouse prior to euthanasia for quantitative colony counts. Compared to CFT073, CFT073cueO colonized the mouse bladder in significantly higher numbers in this infection model (Fig. 2A). This also correlated with increased shedding of CFT073cueO in urine compared to that of CFT073 (Fig. 2B). No colonization of the kidneys was observed for CFT073 or for CFT073cueO; this is consistent with previous data from our laboratory using C57BL/6 mice (36).

Deletion of cueO does not affect type 1 fimbia expression, adhesive capacity, or growth in urine. Expression of type 1 fimbriae significantly enhances the attachment of UPEC to uroepithelial cells and the subsequent colonization of the mouse bladder (2, 4, 20). We compared the levels of type 1 fimbia expression in CFT073 and in CFT073cueO by using a combination of standard techniques. First, the abilities of CFT073 and CFT073cueO to cause mannose-sensitive agglutination of yeast cells were examined. There was no difference in the agglutination titers for the two strains (data not shown). Second, the amounts of FimA produced by the two strains were compared by Western blot analysis using a polyclonal serum raised against purified type 1 fimbriae. There was no difference in the amounts of FimA produced by the two strains (Fig. 3A). Third, a PCR-based assay was employed to determine the orientation of the phase-variable fimA promoter (7, 29). There was no difference in the amounts of “on” and “off” fragments amplified from both strains (Fig. 3B). Finally, we compared the abilities of CFT073 and CFT073cueO to adhere to HeLa epithelial cells as previously described (35). In these assays, CFT073 and CFT073cueO displayed equivalent adherence levels (Fig. 3C). Taken together, the data suggest that the expression levels of type 1 fimbriae were the same in CFT073 and CFT073cueO.

In E. coli K-12, deletion of cueO results in increased aggregation, and this phenotype correlates with the enhanced ex-
performed 55Fe2+ uptake experiments as previously described (18). In these assays, CFT073 cueO accumulated 5.5-fold more 55Fe2+ uptake than did CFT073 (Fig. 5). The introduction of plasmid pCueO into CFT073 cueO restored the level of 55Fe2+ accumulation to approximately wild-type levels (Fig. 5).

Deletion of cueO does not confer a growth advantage in a systemic infection model. To examine if the increased colonization observed for CFT073 cueO in the mouse bladder was also reflected in colonization of other sites, we tested CFT073 and CFT073 cueO in a mouse systemic infection model. Mice were infected with 2 × 10⁷ cells of CFT073 (n = 15) or CFT073 cueO (n = 15) by a subcutaneous injection into the abdomen and monitored to assess the clinical effects of infection, as previously described (22). There was no difference in the virulence levels of the two strains (the time taken to kill the mice was approximately 24 h for both strains). Bacterial colony counts were also performed from the liver, spleen, and kidneys of each mouse; there was no significant difference in the abilities of the strains to colonize these organs.

Conclusions. The results presented herein show that while deletion of cueO in E. coli CFT073 renders the cell sensitive to hydrogen peroxide stress, it also promotes increased uptake of iron. Under the iron-limited conditions encountered in the urinary tract (25, 27, 31, 32), this may represent a competitive advantage, though the consequence may be that it leaves the cell vulnerable to iron-promoted oxidative stress. Recent work by Grass and coworkers (8) has shown that CueO in E. coli K-12 has a high affinity for Fe-enterobactin (Kₘ of 1.5 μM) and that oxidation of the enterobactin precursor 2,3-dihydroxybenzoic acid leads to formation of a polymer capable of chelating copper and iron in the periplasm (8). Production of an iron-chelating polymer in the periplasm of CFT073 under iron-limited conditions may account for the restricted iron uptake observed in this strain; loss of this biological function in the cueO mutant may lead to increased uptake of ferrous iron.

Although the CFT073 cueO strain displayed enhanced bladder colonization in comparison to CFT073, this did not translate into increased growth in human urine. Recent transcriptional profiling of UPEC during the formation of intracellular bacterial communities within epithelial cells revealed that the intracellular environment is iron limited and aerobic (23). Increased iron acquisition from damaged epithelial cells could explain the enhanced colonization of CFT073 cueO in the mouse bladder. Our results lead us to conclude that CueO is critical for iron and copper homeostasis in UPEC and that it has a key role in maintaining a tightly controlled flux of iron into the cell to avoid oxidative stress. The attachment of UPEC to the uroepithelium induces neutrophil recruitment with a...
concomitant oxidative burst that would expose UPEC to reactive oxygen species (11). In view of this, the superior colonization of the bladder by the cue mutant is surprising. However, we note that we employed a short-term infection model, and thus, it is possible that the long-term fitness of CFT073cueO would be diminished in inflamed tissues in which reactive oxygen species cause oxidative stress. This is consistent with our observations that the CFT073cueO mutant is not attenuated in a systemic infection model.

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REFERENCES


