

Genetic Engineering of *Escherichia coli* for Enhanced Uptake and Bioaccumulation of Mercury

WEON BAE,^{1,2} RAJESH K. MEHRA,² ASHOK MULCHANDANI,^{1*} AND WILFRED CHEN^{1*}

*Department of Chemical and Environmental Engineering¹ and Environmental Toxicology Program,²
University of California, Riverside, California 92521*

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Synthetic phytochelatin (ECs) are a new class of metal-binding peptides with a repetitive metal-binding motif, (Glu-Cys)_nGly, which were shown to bind heavy metals more effectively than metallothioneins. However, the limited uptake across the cell membrane is often the rate-limiting factor for the intracellular bioaccumulation of heavy metals by genetically engineered organisms expressing these metal-binding peptides. In this paper, two potential solutions were investigated to overcome this uptake limitation either by coexpressing an Hg²⁺ transport system with (Glu-Cys)₂₀Gly (EC20) or by directly expressing EC20 on the cell surface. Both approaches were equally effective in increasing the bioaccumulation of Hg²⁺. Since the available transport systems are presently limited to only a few heavy metals, our results suggest that bioaccumulation by bacterial sorbents with surface-expressed metal-binding peptides may be useful as a universal strategy for the cleanup of heavy metal contamination.

Mercury is one of the most toxic heavy metals in the environment. The principal sources of contamination in wastewater are chloralkali plants, battery facilities, mercury switches, and medical wastes (12). In an aqueous environment, Hg²⁺ in sediment is subject to methylation, forming more toxic methylmercury (4). Bioaccumulation of methylmercury through the food chains is a potential risk to consumers of contaminated fish or shellfish (7). One of the most severe cases of mercury poisoning occurred in Minamata Bay, Japan, in which hundreds of people died and thousands were affected by consuming contaminated fish.

Common treatments to remove Hg²⁺ from contaminated sources are based on adsorption with ion-exchange resins (14). These technologies, however, are inadequate to reduce Hg²⁺ concentrations to acceptable regulatory standards. Another emerging technology that is receiving more attention is the use of biosorbents. The first commercial biosorbents developed (MRA and Algasorb) were based on sequestration of toxic metals by cell-surface moieties (8). These biosorbents, however, generally lack the required affinity and specificity.

The availability of genetic engineering technology provides the possibility of specially tailoring microbial biosorbents with the required selectivity and affinity for Hg²⁺. One emerging strategy that is receiving more attention is the use of metal-binding peptides. Naturally occurring metal-binding peptides, such as metallothioneins (MTs) and phytochelatin (17), are the main metal-sequestering molecules used by cells to immobilize metal ions, offering selective, high-affinity binding sites. However, the de novo design of metal-binding peptides is an attractive alternative to MTs, as they offer the potential of enhanced affinity and selectivity for heavy metals. Recently, a new class of metal-binding peptides known as synthetic phy-

tochelatin (ECs) with the repetitive metal-binding motif (Glu-Cys)_nGly were shown to have improved Cd²⁺ binding capability over that of MTs (1).

Overexpression of metal-binding proteins such as MTs in bacterial cells resulted in enhanced Hg²⁺ accumulation and thus offers a promising strategy for the development of microbe-based biosorbents (13, 15) for the removal and recovery of Hg²⁺ from contaminated water or soil. However, Hg²⁺ removal by intracellular accumulation has been problematic because of the limited metal uptake (2). This uptake limitation could be potentially overcome either by coexpressing an Hg²⁺ transport system (3) or by anchoring the metal-binding proteins directly on the cell surface (1). In this paper, we describe the characterization of recombinant *Escherichia coli* strains with EC20 either anchored on the cell surface or coexpressed intracellularly with the mercury transport proteins MerP and MerT (2, 10). The ability of these strains to accumulate Hg²⁺ was investigated.

Expression of ECs. Synthetic genes coding for EC20 were synthesized as described previously (1). To express EC20 intracellularly, plasmid pM20 (1) was digested with *Bam*HI and *Hind*III and the DNA fragment coding for EC20 was inserted into pMAL-c2x (New England BioLabs), resulting in pMC20. This construct allows the cytoplasmic expression of EC20 as a fusion to the maltose-binding protein (MBP).

To facilitate the transport of Hg²⁺ across the cell membrane, the Hg²⁺ transport proteins MerP and MerT were coexpressed with MBP-EC20. Plasmid pCLTP (2), containing the *merT* and *merP* genes, was cotransformed with pMC20. Transformed cells were selected on Luria-Bertani (LB) plates containing ampicillin and spectinomycin. For comparison, *E. coli* strain JM109, carrying only pMC20, was also used. An alternate strategy to bypass Hg²⁺ uptake is to directly anchor EC20 on the cell surface. We have successfully demonstrated this possibility using the Lpp-OmpA fusion system (1). Plasmid pLO20, expressing the Lpp-OmpA-EC20 fusion, was used in

* Corresponding author. Mailing address: Department of Chemical and Environmental Engineering, University of California, Riverside, CA 92521. Phone: (909) 787-2473. Fax: (909) 787-2425. E-mail: wilfred@enr.ucr.edu.

TABLE 1. Plasmids and strains

Strain or plasmid	Description	Reference or source
Strain		
JM109	<i>recA1 supE44 endA1 hsdR17 gyrA96 thi Δ(lac-proAB) F' (traD36, proAB, lacI^q ZΔM15)</i>	18
Plasmids		
pUC18	Cloning vector	18
pLO20	A pUC18 derivative containing the <i>lpp-ompA-ec20</i> fusion	1
pMAL-c2x	MBP gene fusion vector	New England BioLabs
pMC20	A pMAL-c2x derivative containing the <i>malE-ec20</i> fusion	This work
pCLTP	A pCL1921 derivative containing the <i>merT</i> and <i>merP</i> genes	2

this study. Bacterial strains and vectors used are listed in Table 1.

To confirm the production of Lpp-OmpA-EC20 and MBP-EC20, cultures were induced with 1 mM isopropyl-β-D-thiogalactopyranoside (IPTG) and radiolabeled cysteine (³⁵S, 1,075 Ci/mmol; ICN) was added at the time of induction. After 15 h, total cell lysates were separated by a sodium dodecyl sulfate (SDS)-12.5% polyacrylamide gel (6). The gel was then dried and exposed to an X-ray film. The high cysteine content of EC20 enables detection of these proteins by autoradiography. Synthesis of full-length Lpp-OmpA-EC20 (21 kDa) and MBP-EC20 (47.5 kDa) fusions was detected at the expected molecular weight (Fig. 1). The intensity of the protein bands was quantified using a Bio-Rad Gel Doc 2000 Gel Documentation System and Quantity One software. Intracellular expression of MBP-EC20 was approximately 12 times higher than expression

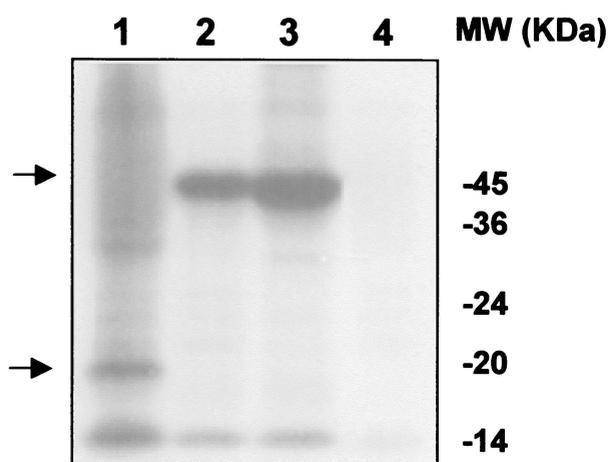


FIG. 1. Expression of EC fusion proteins. [³⁵S]cysteine was added to the cultures at an OD₆₀₀ of 0.3. The cultures were further grown for 15 h. Total cell proteins were separated on SDS-12.5% polyacrylamide gel electrophoresis. The gel was dried and autoradiographed. Expression from induced cultures harboring pLO20 (lane 1), pMC20/pCLTP (lane 2), pMC20 (lane 3), and pMAL-c2x (lane 4), respectively, is shown. The desired fusion proteins are marked with arrows. Molecular mass is shown in kilodaltons on right.

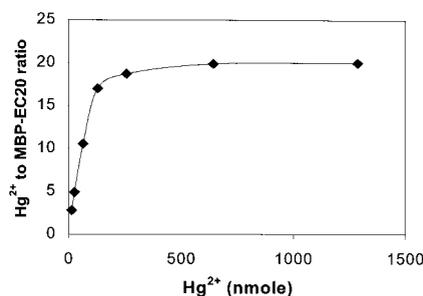


FIG. 2. The Hg²⁺-to-MBP-EC20 stoichiometry expressed as the plot of initial Hg²⁺ concentration against the complexed Hg²⁺-to-peptide ratio. Five nanomoles of purified MBP-EC20 was incubated with 1 to 1,200 nmol of Hg²⁺ in 5 mM dithiothreitol for 1 h. The portion of bound and unbound Hg²⁺ was determined by a mercury analyzer.

on the cell surface, and coexpression of the MerT-MerP transporters reduced MBP-EC20 production by about twofold.

Hg²⁺ binding to EC20. To investigate the Hg²⁺ binding stoichiometry of EC20, MBP-EC20 fusion proteins were purified from cultures of JM109/pM20 using an amylose resin affinity column (New England BioLabs). The purity of the protein was confirmed through SDS-12.5% polyacrylamide gel electrophoresis. Five nanomoles of the purified fusion protein was resuspended in 50 mM Tris-Cl buffer (pH 7.4) supplemented with 5 mM dithiothreitol and incubated with 1 to 1,200 nmol of Hg²⁺ for 2 h. Hg(II)-glutathione complexes were used instead of HgCl₂ in order to prevent precipitation as reported previously (9). The protein-Hg²⁺ complex was recovered using a Microcon centrifugal filter membrane (Millipore), and the amount of bound Hg²⁺ was measured by cold-vapor atomic absorption spectroscopy (Coleman Model 5B Mercury Analyzer System). The Hg²⁺-to-MBP-EC20 stoichiometry was determined by plotting the initial Hg²⁺ concentration against the molar ratio of bound Hg²⁺ to MBP-EC20 (Fig. 2). A saturating ratio of 20 Hg²⁺ per MBP-EC20 was obtained, a value much higher than the typical ratio of 7 reported for MTs (11). A similar binding experiment was conducted with purified MBP, with no significant binding of Hg²⁺ observed.

Bioaccumulation of Hg²⁺. To investigate the effect of uptake on bioaccumulation of Hg²⁺, the binding capabilities of various *E. coli* strains were compared. Overnight cultures grown in LB medium at 37°C were harvested, washed with distilled water twice, and resuspended to a final optical density at 600 nm (OD₆₀₀) of 1.0 in LB medium containing 5 μM Hg²⁺. The Hg²⁺ contents were determined after 1 h. As shown in Fig. 3, *E. coli* strain JM109/pUC18 accumulated a very low level of Hg²⁺. The intracellular accumulation of Hg²⁺ increased by sixfold for cells overexpressing MBP-EC20 (JM109/pMC20). By elimination of Hg²⁺ uptake, cells with EC20 anchored on the cell surface (JM109/pLO20) accumulated about threefold more Hg²⁺ than did cells with EC20 expressed in the cytoplasm. This threefold improvement is in good agreement with our earlier observation of Cd²⁺ accumulation using cells with surface-expressed EC20 (1). In the presence of the Hg²⁺ transporters (JM109/pCLTP/pMC20), intracellular accumulation of Hg²⁺ also increased significantly. The level of Hg²⁺ accumulation was similar to that for cells expressing EC20 on the

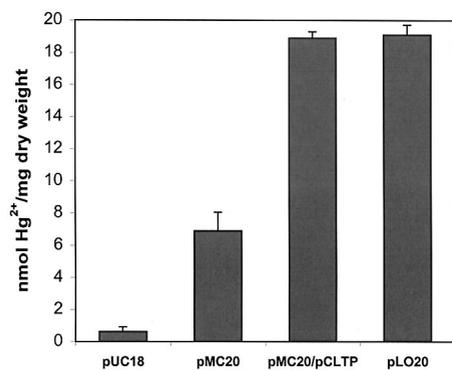


FIG. 3. Bioaccumulation of Hg²⁺ by resuspended cultures harboring various plasmids from LB medium containing 5 μ M Hg²⁺. Data were obtained from three independent experiments.

surface. In both cases, 100% of the added Hg²⁺ was removed after 1 h. These results indicate that uptake is indeed the rate-limiting step for the intracellular accumulation of Hg²⁺.

Localization of the accumulated Hg²⁺ was determined by separating cells into cytoplasmic and membrane fractions as described before (1). Consistent with the localization of EC20, 80% of the accumulated Hg²⁺ was associated with the cytoplasmic fraction for both JM109/pMC20 and JM109/pCLTP/pMC20 cells, while over 90% of the accumulated Hg²⁺ was found in the membrane fraction of JM109/pLO20 cells. These results demonstrate that bioaccumulation proceeds in the virtual absence of Hg²⁺ uptake for cells with EC20 displayed on the surface. Such an approach should be beneficial not only to the overall capacity but also to the kinetics of the bioaccumulation.

To determine the benefits on the rate of Hg²⁺ bioaccumulation, a time course assay was carried out. Overnight cultures were harvested, washed with distilled water twice, and resuspended to a final OD₆₀₀ of 1.0 in LB medium containing 5 μ M Hg²⁺. As shown in Fig. 4A, cells expressing only MBP-EC20 (JM109/pMC20) accumulated Hg²⁺ at a very low rate, with less than 20% removed after 20 min. Coexpression of the Hg²⁺ transporters and MBP-EC20 (JM109/pCLTP/pMC20) improved the bioaccumulation rate significantly, with 95% of the added Hg²⁺ removed within 20 min. These results again confirmed that Hg²⁺ uptake is the rate-limiting step in the bioaccumulation of Hg²⁺. However, the rate of bioaccumulation was further improved for JM109/pLO20 cells with EC20 expressed on the surface; over 95% of the added Hg²⁺ was removed within 1 min. It appears that the introduction of EC20 on the cell surface is even more effective in eliminating the uptake limitation, resulting in virtually instantaneous removal of Hg²⁺.

Evaluation of bioaccumulation. The maximum bioaccumulation capacity of two best cell lines overexpressing EC20 (JM109/pLO20 and JM109/pMC20/pCLTP) was determined over a range of Hg²⁺ concentrations (Fig. 4B). At the lower levels (<20 nmol), 100% of the added Hg²⁺ was removed within 1 h. Although the level of EC20 expressed on the surface is approximately fivefold lower than the expression of MBP-EC20 from JM109/pCLTP/pMC20, the highest level of accumulation was around 230 nmol/mg (dry weight) for both

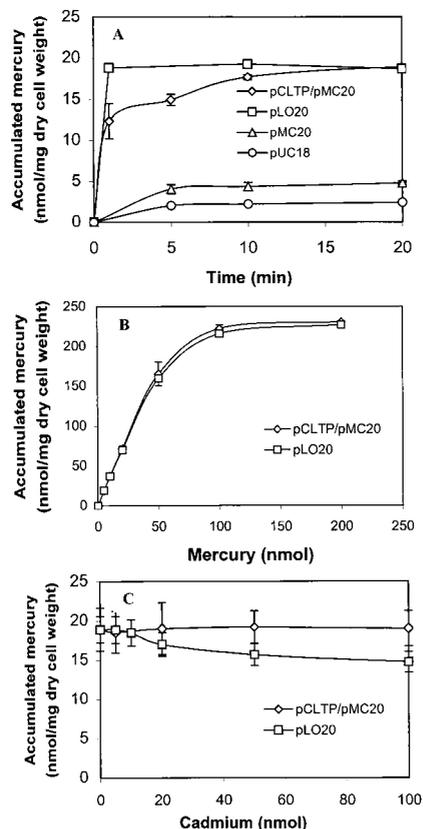


FIG. 4. (A) Time course of mercury uptake by resting cultures harboring various plasmids. Resting cultures were resuspended in LB medium containing 5 μ M Hg²⁺ and were incubated for 20 min. Cells were harvested at various times, and the supernatant was removed by centrifugation for 30 s. (B) Hg²⁺ bioaccumulation capacity of JM109 cells (0.265 mg [dry weight]) harboring either pLO20 or pCLTP/pMC20. Bioaccumulation of Hg²⁺ was measured at various concentrations after 1 h of incubation. (C) Effect of cadmium on bioaccumulation of Hg²⁺. JM109 cells harboring either pLO20 or pCLTP/pMC20 were incubated with 5 nmol of Hg²⁺ and various concentrations of Cd²⁺. The amount of Hg²⁺ accumulated by cells after 1 h was determined.

cell lines. It has been reported previously that the bioaccumulation of Cd²⁺ by cells expressing MT on the surface exceeds by at least 1 order of magnitude the theoretical amount contributed by the surface-exposed MT moiety (16). It appears that the surface-exposed MT helps to increase the local metal concentration around the cells and facilitates interactions of the metal ions with other cell wall components. A similar situation may have occurred here for cells with surface-exposed EC20. It should be noted that the maximum bioaccumulation observed in this study is twofold higher than that reported for cells overexpressing MT and the mercury transporters (3) and falls within the higher range reported for other microorganisms (15 to 290 μ mol/g of cells [dry weight]). This increase in capacity may also reflect the improved Hg²⁺ binding stoichiometry offered by EC20 over that of MT.

The selectivity of the different cell lines for Hg²⁺ bioaccumulation was investigated by performing the Hg²⁺ accumulation experiments in the presence of various amounts of cadmium (Fig. 4C). Cd²⁺ was selected because it is commonly

found in sites contaminated with Hg^{2+} and is also one of the most toxic heavy metals. Because of the specificity of the Hg^{2+} transporters (3), no effect on Hg^{2+} bioaccumulation was observed with JM109/pCLTP/pMC20 cells even in the presence of a 20-fold excess of Cd^{2+} . JM109/pLO20 cells with EC20 displayed on the surface were slightly less selective for Hg^{2+} ; the amount of accumulated Hg^{2+} declined gradually with an increasing excess of Cd^{2+} . However, even in the presence of a 20-fold excess of Cd^{2+} , JM109/pLO20 cells retained about 80% of their Hg^{2+} bioaccumulation activity. Since the binding affinity of Hg^{2+} to MTs and phytochelators is reported to be much stronger than that of Cd^{2+} (11, 17), this slight decrease in selectivity may be due to the nonspecific binding of Cd^{2+} to the other cell wall components, which contributes greatly to the overall bioaccumulation of Hg^{2+} .

The effects of ionic strength and metal chelators on Hg^{2+} bioaccumulation were investigated. The addition of up to 200 mM NaCl and 1 mM EDTA did not change the Hg^{2+} bioaccumulation levels for either JM109/pCLTP/pMC20 or JM109/pLO20 cells. The resistance of both systems to the presence of EDTA and NaCl makes them ideal for the removal of Hg^{2+} in contaminated wastewaters.

Conclusions. Two different strategies were used to enhance the uptake and bioaccumulation of Hg^{2+} by cells overexpressing EC20. Our results indicate that the expression of EC20 on the cell surface is as efficient as the coexpression of Hg^{2+} transporters in alleviating the uptake limitation, resulting in rapid, selective, and high-level bioaccumulation of Hg^{2+} . Since specific transporters have been identified only for a few heavy metals such as mercury and nickel (5, 10), surface expression of metal-binding peptides may be useful as a common strategy to bypass the uptake of any heavy metal of interest, a highly desirable property not associated with the metal transporter systems.

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