The Mtr Respiratory Pathway Is Essential for Reducing Flavins and Electrodes in *Shewanella oneidensis*⁷†

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The Mtr respiratory pathway of Shewanella oneidensis strain MR-1 is required to effectively respire both soluble and insoluble forms of oxidized iron. Flavins (riboflavin and flavin mononucleotide) recently have been shown to be excreted by MR-1 and facilitate the reduction of insoluble substrates. Other Shewanella species tested accumulated flavins in supernatants to an extent similar to that of MR-1, suggesting that flavin secretion is a general trait of the species. External flavins have been proposed to act as both a soluble electron shuttle and a metal chelator; however, at biologically relevant concentrations, our results suggest that external flavins primarily act as electron shuttles for MR-1. Using deletion mutants lacking various Mtr-associated proteins, we demonstrate that the Mtr extracellular respiratory pathway is essential for the reduction of flavins and that decaheme cytochromes found on the outer surface of the cell (MtrC and OmcA) are required for the majority of this activity. Given the involvement of external flavins in the reduction of electrodes, we monitored current production by Mtr respiratory pathway mutants in three-electrode bioreactors under controlled flavin concentrations. While mutants lacking MtrC were able to reduce flavins at 50% of the rate of the wild type in cell suspension assays, these strains were unable to grow into productive electrode-reducing biofilms. The analysis of mutants lacking OmcA suggests a role for this protein in both electron transfer to electrodes and attachment to surfaces. The parallel phenotypes of Mtr mutants in flavin and electrode reduction blur the distinction between direct contact and the redox shuttling strategies of insoluble substrate reduction by MR-1.

Shewanella oneidensis strain MR-1 (MR-1) is a facultative anaerobe capable of respiring a variety of substrates, including various metals and metal oxides, a phenotype that is important for bioremediation and metal cycling in natural environments (22, 53). At near-neutral pH, Fe(III) and Mn(IV) often are present as insoluble oxide minerals. Dissimilatory metal-reducing bacteria such as MR-1 have developed pathways to transfer electrons from the interior of the cell to these external terminal electron acceptors. In some bacteria, these pathways also can transfer electrons to electrodes, which can be harnessed for renewable energy and remote biosensor applications (23, 26, 27). Beyond increasing our understanding of this unusual process, applying anaerobic microbial extracellular respiration to new technologies requires a thorough understanding of the molecular dynamics and cellular physiology of electron source utilization (substrate oxidation) and the reduction of insoluble terminal electron acceptor(s). There are four proposed mechanisms to explain how insoluble substrates are reduced by Shewanella: (i) direct contact, (ii) electron shuttling, (iii) chelation, and (iv) electrically conductive appendages (reviewed in reference 18). We will focus on the first three strategies here.

Flavins recently have been discovered to accelerate the reduction of both iron oxide minerals (51) and electrodes (30) by MR-1. Riboflavin (vitamin B_2) is a precursor for the biosynthesis of flavin mononucleotide (FMN) and flavin adenine dinucleotide (FAD) (13). Riboflavin and FMN both can be observed to build up in the supernatant of anaerobic and aerobically grown cultures of MR-1 (30, 51). However, the mechanism by which flavins enhance the rate of iron oxide mineral or electrode reduction is unknown, although recent work is consistent with a critical role for these compounds in mediating solid Fe(III) reduction by MR-1 (42). Since soluble (chelated) Fe(III) is reduced faster than insoluble Fe(III) by MR-1 (6), one possible explanation for the enhancement of insoluble iron reduction by flavins is increased available soluble iron via chelation (1, 2, 30). Flavins also may be utilized as redox-active compounds to traffic electrons between extracellular reductases on the surface of the cell and insoluble substrates (30, 51), a process termed electron shuttling (18, 39, 41). The chelation of the terminal electron acceptor during electrode reduction is not relevant when the anode is composed of graphite. Therefore, electron shuttling likely is responsible for the flavin enhancement of current production on poised-potential electrodes (30). However, it is unclear if the chelation of metals by flavins influences insoluble metal reduction by S. oneidensis (30).

The Mtr pathway is required for the reduction of metals and electrodes (5, 6, 9, 17). Five primary protein components have been identified in this pathway: OmcA, MtrC, MtrA, MtrB, and CymA (47). Current models of electron transfer in MR-1 assume that electrons from carbon source oxidation are passed via the menaquinone pool to the inner membrane-anchored c-type cytochrome CymA (19, 31). These electrons then are transferred to a periplasmic c-type cytochrome, MtrA, and eventually to outer membrane (OM)-anchored c-type cytochromes MtrC and OmcA, which interact with an integral OM scaffolding protein, MtrB (32, 33, 43). These OM cytochromes

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then can reduce various substrates, including iron oxides and electrodes (8, 9, 12, 36, 47). Since the Mtr system is required by MR-1 to reduce many different substrates (18), it also could be capable of reducing extracellular flavins. Indeed, electron transfer to carbon electrodes is impaired in strains lacking Mtr pathway components (9, 17), which may be explained simply by a decreased ability to reduce extracellular flavins. The observation that Mtr mutants produce less current on electrodes than the wild type could be due to (i) less current generated per cell (either direct reduction or flavin mediated), (ii) decreased attachment to the electrode surface, (iii) differences in external flavin concentrations, or (iv) a combination of these three possibilities. Determining the specific activity (current produced per unit of attached biomass) of Mtr mutants on electrodes under conditions where flavin levels were controlled would allow for differentiation between these possibilities. To date, this kind of analysis has not been reported.

The results presented here extend our knowledge of how S. oneidensis catalyzes the reduction of insoluble substrates. Experiments using a model iron chelator and electron shuttle are consistent with electron shuttling being the primary mechanism by which flavins enhance insoluble iron oxide reduction rates. Moreover, we demonstrate that MR-1 reduces extracellular flavins at physiologically relevant rates and that the Mtr pathway accounts for at least 95% of this activity. The specific activities of various mutant strains lacking Mtr pathway components on poised-potential electrodes also are reported. Our data suggest that MtrC is responsible for most of the electron transfer to carbon electrodes, while OmcA is involved in attachment and has a lesser role in electron transfer. These observations could have broader implications regarding the role of OmcA in the reduction of soluble and insoluble substrates (8, 9, 36).

MATERIALS AND METHODS

Bacterial strains and media used. *S. oneidensis* strain MR-1 originally was isolated from Oneida Lake in New York (34, 50). Overnight cultures were made from single colonies freshly streaked from a frozen stock culture inoculated into 2 ml of Luria-Bertani (LB) medium and aerobically grown for 16 h. The minimal medium used for experiments was *Shewanella* basal medium (SBM) as previously described (21). Anaerobic cultures were flushed with nitrogen gas for 15 min in Balch anaerobic tubes sealed with a butyl rubber stopper (3). All anaerobic cultures were supplemented with 0.05% Casamino Acids and contained 40 mM fumarate or 5 mM iron oxide (ferrihydrite) prepared as previously described (45) as the electron acceptor, where indicated. *S. baltica* strain OS155, *S. putrefaciens* strain CN-32, *S. loihica* strain PV-4, *S. frigidimarina* strain NCIMB 400, *S. denitrificans* strain OS217T, and *S. amazonensis* strain SB2B were obtained from Jim Fredrickson (Pacific Northwest National Laboratories).

Total flavin quantification. Overnight cultures of different strains of *Shewanella* were grown aerobically for 16 h in LB medium, and the optical density (OD) was read at 600 nm. Cultures were centrifuged, and oxidized supernatants were analyzed in a 96-well plate in a Molecular Devices Spectra-Max Pro reader with an excitation wavelength of 440 nm and an emission wavelength of 525 nm. Standard curves were made from stock solutions of oxidized riboflavin or FMN in LB medium for comparison. High-performance liquid chromatography (HPLC) was performed as previously described (30) on the same supernatants to confirm that fluorescence was due specifically to riboflavin.

Flavin reduction assay. OD readings of overnight cultures were taken at 600 nm, and cells were washed once with SBM and resuspended to an OD of 4.0 (verified by monitoring diluted samples). For cuvette experiments, 100 μ l of this suspension was injected into a sealed cuvette containing 1 ml of SBM with vitamins and minerals to obtain a final concentration of 12 μ M riboflavin and either 18 mM lactate or no electron donor. The sealed cuvette was purged with nitrogen gas for 5 min prior to cell addition. For reoxidation experiments the

same protocol was followed, except that a 1:10 dilution of cells was used (for a final OD_{600} of 0.4). After 1 h, the cuvette was removed from the spectrophotometer and vortexed thoroughly without the stopper for ~ 10 s and resealed, and then readings were resumed.

For 96-well plate assays, 20 μ l of mutant or wild-type cultures prepared as described above were aliquoted into a black 96-well plate in a Coy anaerobic chamber containing an atmosphere of 5% hydrogen, 20% carbon dioxide, and 75% nitrogen. Reactions were initiated with 280 μ l SBM to obtain a final concentration of 120 μ M riboflavin and 18 mM lactate, and fluorescence (excitation, 440 nm; emission, 525 nm) was monitored over time. High concentrations of flavins were used in these experiments to obtain accurate and reproducible reduction rate measurements for the various strains tested.

Iron reduction. MR-1 was grown overnight aerobically in LB medium, washed once, and resuspended in SBM to an OD of 1.30. Thirty microliters of this suspension was added to 270 μ l of SBM containing vitamins; minerals; final concentrations of 18 mM lactate, 4.5 mM ferrihydrite, and 12 μ M of 9,10-anthraquinone-2,6-disulfonic acid (AQDS); 12 mM or 12 μ M sodium citrate; or 12 μ M riboflavin in 96-well plates. Plates were kept in a GasPak System anaerobic petri dish holder that was flushed for 15 min with nitrogen gas between time points. At each time point, the oxidation of Fe(II) was prevented by adding 50 μ l of 5 M hydrochloric acid (HCl). Thirty microliters was taken from this well and diluted in 270 μ l of 0.5 M HCl to yield concentrations within the range of standard curves. Fifty microliters of the diluted sample was mixed with 300 μ l ferrozine reagent (48) and read at 562 nm. Standard curves were made from ferrous sulfate dissolved in 0.5 N HCl.

Mutant construction/verification. Mutant strains lacking *mtrC*, *mtrB*, *mtrA*, and *omcA* have been described previously (21) and were generated by the ligation of regions flanking target genes into the suicide vector pSMV3. For the strain lacking both *omcA* and *mtrC*, the upstream and downstream primers were as follows: DCdelUP1, NNNGAGCTCTTATGCTAACTGGGGAACC; DCdelUP2, NNNGAATTCACAGCATCATCTGAGTTCCT; DCdelDOWN1, NNNGGTCTCGTAATTTGCCCAAGCAGG; and DCdelDOWN2, NNNACG AGTCATTCACACTACCGTGAG. PCR using primers flanking target genes confirmed that deletions had occurred, and all single deletion strains were complemented using the expression vector pBBR1MCS-2 (24). Quantitative reverse transcription-PCR (RT-PCR) was performed on all mutant strains using a DyNAmo Sybr green kit according to the manufacturer's protocol and read using an Applied Biosystems 7900 HT sequence detection system.

Operation of the three-electrode bioreactor. Bioreactors were constructed as previously described (30). Cells inoculated from a fresh LB medium overnight culture were grown aerobically for 24 h in SBM with 0.05% Casamino Acids, 30 mM lactate, and 40 mM fumarate. A 10% transfer into anaerobic SBM with vitamins, minerals, 0.05% Casamino Acids, 20 mM lactate, and 40 mM fumarate was performed, and the cells were allowed to grow until an OD between 0.3 and 0.4 was achieved. Ten milliliters of this anaerobic culture was used to inoculate sterile and nitrogen-degassed electrochemical cells. The chamber was flushed continuously with oxygen-free nitrogen gas, and the electrodes were maintained at oxidizing potential (+0.242 V versus SHE [standard hydrogen electrode]) using a 16-channel VMP potentiostat (Bio-Logic SA). To ensure that all strains were exposed to similar initial flavin concentrations, the bioreactor medium was supplemented with 250 nM riboflavin. Additional lactate at 10 mM also was added at the start of the experiment to prevent carbon source limitation during growth.

CA measurements. The electron transfer rate at the poised (+0.242 V versus SHE) working electrode was measured during a period of 96 h by recording amperometry data points every 900 s. Chronoamperometry (CA) growth curves were collected four times for each strain to demonstrate the repeatability of the procedure and increase statistical significance. To obtain data from fully colonized electrodes under identical flavin and lactate concentrations, the medium was replaced when strains reached a phase of maximum current production. Within 6 h of this medium replacement (containing 20 mM lactate and 0.5 μ M riboflavin), current production stabilized, and electrodes were harvested for protein determination.

BCA protein measurement. Attached biomass was quantified using a bicinchoninic acid (BCA) assay (Pierce). Electrodes were detached from bioreactors and placed in a 1.5-ml tube containing 1 ml of 0.2 N NaOH. The tubes were heated at 96°C in a heat block for 20 min to lyse and detach cells attached to the electrode surface. The tubes were cooled to room temperature and assayed according to the manufacturer's protocol at 562 nm using a SpectraMax Pro (Molecular Devices). Standard curves were generated using bovine serum albumin.

Confocal microscopy. Biofilm coverage and thickness were analyzed by confocal imaging. Carbon working electrodes prepared as described above display-

 TABLE 1. Flavin production in rich medium by various Shewanella species^a

Strain	Flavin concn. measured as:	
	μM (in culture)	μmol/mg protein
S. oneidensis MR-1	1.31 ± 0.20	2.8 ± 0.4
S. frigidimarina NCIMB 400	1.01 ± 0.10	2.4 ± 0.2
S. loihica PV-4	1.05 ± 0.16	5.3 ± 0.8
S. baltica OS155	0.56 ± 0.04	2.8 ± 0.2
S. amazonensis SB2B	0.87 ± 0.28	1.9 ± 0.6
S. denitrificans OS217T	0.54 ± 0.03	3.0 ± 0.2

^{*a*} Strains were grown for 16 h aerobically, and supernatant fluorescence was quantified to determine total flavin excreted. PV-4, OS155, NCIMB 400, and OS217T were grown for 30 h to reach optical densities similar to those of other strains.

ing steady-state current production were harvested and rapidly placed in amber 1.5-ml centrifuge tubes containing 1 ml SBM with 60 µM propidium iodide and 10 µM Syto 9 from a Live/Dead BacLight bacterial viability kit (Invitrogen, Carlsbad, CA). The sample was stained for 20 min, rinsed, and placed in the center of a microscope slide surrounded by a gasket. The basin was filled with 50 μl SBM, and a large coverslip was sealed in place. The stained biofilm was viewed with a Nikon Eclipse C1 confocal microscope using 488- and 561-nm filters. The morphology and coverage of the biofilm were evaluated at several locations, and representative sites were chosen for the collection of three-dimensional images using 0.5-µm intervals, beginning approximately 20 µm above the top of the film and continuing until 20 µm beyond the electrode/biofilm interface. The resulting slices were rendered using EZ-C1 3.20 FreeViewer software. Confocal images were picked as the best representatives of at least n = 3 experiments for each strain, with z projections performed at multiple locations on each electrode to ensure accurate results. All averaged protein data are the results of at least n =5 measurements for each strain.

RESULTS

Flavin production by Shewanella strains. MR-1, Shewanella sp. strain MR-4, and Shewanella strains isolated from textile waste facilities have been shown previously to produce and excrete flavins (30, 51). We tested other Shewanella strains closely related to MR-1, and several much more distantly related strains (as determined by 16S rRNA gene phylogeny [22]), to determine if flavin accumulation in supernatants is a common feature of these bacteria. The strains tested were S. baltica strain OS155, S. putrefaciens strain CN-32, S. loihica strain PV-4, S. frigidimarina strain NCIMB 400, S. denitrificans strain OS217T, and S. amazonensis strain SB2B. Of the strains tested, all have been reported to reduce iron oxide, except S. denitrificans (10, 22), and all accumulated flavins at levels near the value observed for MR-1 (Table 1). Flavins accumulated to $\sim 1.0 \ \mu$ M in cultures grown in rich medium, and this accumulation appears to be a general trait among the shewanellae (as previously suggested by von Canstein et al. [51]), and it may have additional functionality beyond accelerating the rates of insoluble substrate reduction in certain species such as S. denitrificans.

Chelation and electron shuttling. We previously hypothesized that flavins have two distinct mechanisms to accelerate rates of insoluble iron reduction by MR-1: chelation and electron shuttling (30). To observe the effects of extracellular chelators and electron shuttles on insoluble iron reduction in MR-1, we conducted experiments using AQDS (a synthetic analogue of the redox active moieties in humic acids) as a model electron shuttle and citrate as a model iron chelator. We quantified the degree to which these compounds increased iron oxide reduction rates when added exogenously to MR-1 cultures. With no addition, MR-1 reduced iron oxide at an initial rate of 75.7 μ M/h under the conditions tested. AQDS at $12 \,\mu\text{M}$ increased this rate to $183 \,\mu\text{M/h}$ (Fig. 1), while the same amount of riboflavin resulted in a rate of 317 µM/h. Using FMN instead of riboflavin yielded identical results (data not shown). Citrate addition did not have a significant effect at the same concentration (12 µM), but at concentrations 1,000 times higher (12 mM) the initial rate of iron reduction was increased to 613 µM/h. The binding constant of citrate for iron is four orders of magnitude higher than that of riboflavin (2, 15), suggesting that at μM concentrations, the chelation of iron by flavins does not significantly contribute to the increased rate of iron reduction. These data are consistent with the electron shuttling activity of flavins as the primary mechanism for the enhancement of insoluble iron reduction by MR-1.

Flavin reduction by wild-type cells. If flavins shuttle electrons between cells and solid surfaces, extracellular flavins must be reduced directly by MR-1. Flavins fluoresce when oxidized but not when reduced (16), making it possible to monitor the oxidation state of riboflavin over time. When wildtype cells were anaerobically incubated with riboflavin and lactate, fluorescence decreased over time, which is consistent with the reduction of flavins (Fig. 2A). When lactate was omitted from the reaction, fluorescence did not change over time, demonstrating that this loss of fluorescence was dependent on electron donor availability (Fig. 2A). The fluorescence change of cultures incubated with FMN occurred at the same rate as that of riboflavin (data not shown), indicating that FMN is reduced at the same rate as that of riboflavin. After fluorescence readings had reached a minimum value, samples were exposed to oxygen. Fluorescence readings returned to the



FIG. 1. Acceleration of *S. oneidensis* iron oxide reduction by electron shuttles versus that by a strong chelator. MR-1 cells were inoculated into SBM containing lactate, iron oxide, and 12 μ M AQDS (\blacktriangle), 12 μ M riboflavin (\bigcirc), 12 μ M sodium citrate (\bigcirc), 12 mM sodium citr



FIG. 2. Development of flavin reduction assay by *S. oneidensis*. (A) MR-1 cells were washed in minimal medium and injected to an anaerobic cuvette containing minimal medium and riboflavin as the electron acceptor and given either 18 mM lactate (\Box) or no carbon source (\bigcirc). Relative fluorescence units (RFU) were measured over time as described in the text. Error bars represent standard deviations from triplicate experiments. (B) MR-1 cells were examined as described for panel A, only a lower cell density was used. The arrow indicates when the culture was exposed to the atmosphere and vortexed for ~10 s. Culture then was resealed and readings were resumed. This figure is representative of three independent experiments.

same value as that at the beginning of the experiment (Fig. 2B) and could be reduced again when returned to anaerobic conditions. These experiments confirmed that flavins were not consumed or destroyed and could be recycled. Therefore, MR-1 is able to couple the reduction of flavins to the oxidation of lactate in a manner consistent with its use as a continuously regenerated electron shuttle.

Flavin reduction by Mtr mutants. To determine if the Mtr pathway is used in the reduction of flavins, gene deletion mutants were constructed. Strains lacking the genes omcA $(\Delta omcA)$, mtrC $(\Delta mtrC)$, mtrA $(\Delta mtrA)$, and mtrB $(\Delta mtrB)$, as well as a strain lacking both *omcA* and *mtrC* ($\Delta omcA \Delta mtrC$), were tested for their ability to reduce both riboflavin and FMN. Both flavins were tested here, because they have been detected previously in wild-type cultures of electrode (28) and iron-reducing cultures (49) of S. oneidensis strain MR-1. Quantitative RT-PCR confirmed that in all constructed mutants, the expression of the remaining mtr genes was not altered more than twofold compared to that of the wild type (see Table S1 in the supplemental material). Mutants in the Mtr pathway did not reduce flavins at the same rate as the wild type, indicating that this pathway is involved in the reduction of riboflavin and FMN by MR-1 (Fig. 3). All single gene deletion strains were



FIG. 3. Reduction rates of riboflavin and FMN by mutants in the Mtr pathway. Rates were calculated over the linear portion of the initial reduction curves, such that all r^2 values were greater than 0.95. Error bars indicate standard errors of reduction rates from three independent cultures. Under these experimental conditions, 1 μ M flavin is equal to approximately 166 relative fluorescence units (RFU).

complemented, and flavin reduction activity was restored to near-wild-type rates (data not shown). Both MtrA and MtrB were essential for this process, as $\Delta mtrA$ and $\Delta mtrB$ reduced flavins at less than 5% of the rate of MR-1. Mutants lacking OmcA reduced flavins at nearly the same rate as the wild type, while $\Delta mtrC$ reduced flavins at about half the rate of the wild type. $\Delta omcA \ \Delta mtrC$ reduced flavins at only ~10% of the rate of MR-1 (Fig. 3). This suggests that OmcA can reduce riboflavin and FMN in the absence of MtrC, but it does not significantly contribute to the observed rate in wild-type cells. We conclude that MR-1 transfers electrons from the interior of the cell to flavins using the Mtr pathway, and approximately 90% of this activity occurs at the outer membrane using the OM proteins OmcA and MtrC.

Specific activity of Mtr mutants in three-electrode bioreactors. The presence of flavins has been shown to significantly accelerate current production when MR-1 utilizes carbon electrodes as a terminal electron acceptor (30). If the Mtr pathway is responsible for flavin-mediated electrode reduction as well, the ability of strains to reduce flavins should mirror rates of current production, especially in a poised-electrode system where the electron acceptor is not limiting. To address this, current production was monitored over time for all mutant strains when grown in a three-electrode bioreactor until a maximum current was obtained (Fig. 4A). Consistently with defects in flavin reduction, $\Delta mtrC$, $\Delta mtrA$, $\Delta mtrB$, and $\Delta omcA$ $\Delta mtrC$ all grew slower and produced less maximum current than the wild type. Strikingly, $\Delta omcA$ produced only 50% of the maximum current of the wild type, although it had only a slight defect in flavin reduction (Fig. 3).

As all of these observations were collected at identical levels of extracellular flavins, differences in flavin concentrations could not explain the observed differences between strains. In addition, measurements also were collected for electrodeattached cells that were washed free of extracellular flavins. The depletion of flavins decreased current production by all wild-type and mutant cultures by more than 70% (Fig. 4B), demonstrating that flavin shuttling was the dominant route of electron transfer in all of these incubations.

It was unknown if Mtr mutant strains produced lower cur-



FIG. 4. Analysis of electrode reduction capacity of Mtr pathway mutants. (A) Example current production over time by Mtr mutants, using controlled-potential 2-cm² graphite electrodes. The single curves shown are representative of four replicates. Experiments were performed with MR-1 (\blacksquare), $\Delta omcA$ (\blacktriangle), $\Delta mtrC$ (\blacklozenge), $\Delta mtrA$ (\Box), $\Delta mtrB$ (\triangle), and $\triangle omcA \ \Delta mtrC$ (\Diamond). (B) Flavin acceleration of electrode reduction by S. oneidensis requires the Mtr pathway. Cultures were grown on electrodes to a maximum current production rate (72 to 96 h) and were washed free of external flavins. Current produced at electrodes was recorded after a 6-h equilibration period (white bars). Riboflavin at 0.5 µM was added and current was again recorded after a 6-h equilibration period (gray bars). Representative data from at least two experiments are shown. (C) Maximum current production versus protein attached to electrode surface for different strains. Current production data were recorded for each strain at a flavin concentration of 0.5 μ M. Experiments were performed using MR-1 (\blacksquare), $\Delta omcA$ (**A**), $\Delta mtrC$ (**\diamond**), $\Delta mtrA$ (\Box), $\Delta mtrB$ (\triangle), and $\Delta omcA$ $\Delta mtrC$ (\diamond) . Error bars indicate standard errors from triplicate experiments.

rent at electrodes due to lower per-cell rates of electron transfer or simply less attachment by cells to electrode surfaces, both of which are required for current production (30). To differentiate between these two possibilities, total protein attached to electrode surfaces was quantified after measuring the maximum current for each of the mutant strains and the wild type under identical flavin concentrations. The most electrodeattached protein was observed with wild-type biofilms (50.1 \pm 6.1 µg total protein per electrode), while strains lacking omcA $(34.9 \pm 4.3 \ \mu g$ total protein per electrode) and *mtrC* (24.7 \pm 10.1 \ \mu g 1.4 µg total protein per electrode) attached to electrodes 70 and 50% as effectively as the wild type, respectively (Fig. 4C). Protein quantification revealed that $\Delta mtrA$, $\Delta mtrB$, and $\Delta omcA \ \Delta mtrC$ did not significantly colonize electrode surfaces (Fig. 4C). Calculating electron transfer rates as specific activities revealed the current production per unit of attached protein to be 0.81 µA/µg protein for MR-1, 0.60 µA/µg protein for $\Delta omcA$, and 0.090 $\mu A/\mu g$ protein for $\Delta mtrC$ (Fig. 4C). The complementation of $\Delta omcA$ returned iron reduction and current production to wild-type levels (see Fig. S2 in the supplemental material). These results showed that while $\Delta omcA$ produced less current per electrode, this difference was due more to a colonization defect than an electron transfer defect per se.

Confocal microscopy of wild-type, $\Delta omcA$, and $\Delta mtrC$ strains on electrodes harvested after measuring the maximum current and visualized using live/dead fluorescent staining agreed with total protein differences observed between these strains (Fig. 5). Total volume calculations of biofilms estimated using the Image Structure Analyzer software from the Center for Biofilm Engineering at Montana State University (57) were consistent with attached biomass measurements. Surface plots to aid the visualization of the very thin (1 to $2 \mu m$) biofilms formed by the mutant strains are provided in the supplemental material (see Fig. S2). Interestingly, the small amount of $\Delta mtrC$ biomass on the electrode surface stained red, suggesting that these cells possessed compromised membranes, and this explained the very low current output per unit protein observed (Fig. 4C). These results were consistent with the hypothesis that OmcA is required for attachment and/or biofilm formation on electrodes, while MtrC is the primary terminal reductase for electron transfer under electrode-reducing conditions.

DISCUSSION

Although it was known that secreted flavins increased the rates of electrode (30) and insoluble iron (51) reduction by *S. oneidensis*, the exact mechanism of this enhancement remained elusive prior to this study. In every species of *Shewanella* we have tested to date, flavins are found to accumulate in culture supernatants. Intriguingly, one of the few characterized *Shewanella* species unable to reduce insoluble iron oxide minerals, *S. denitrificans*, accumulated flavins to similar levels. The genome analysis of *S. denitrificans* strain OS217 indicates that the genes encoding the Mtr respiratory pathway (*omcA, mtrC, mtrB*, and *mtrA*) are absent (14). One possibility is that *S. denitrificans* recently lost the Mtr pathway, and the secretion of flavins is a metabolic relic. On the other hand, extracellular flavins may be critical for a non-respiratory role, such as iron acquisition (7, 52, 55), the sensing of redox gradients, or pro-



FIG. 5. Confocal microscopy of *Shewanella* biofilms attached to electrode surfaces. For each panel, the left side is a top-down maximum projection of the electrode surface and the right side is a single representative cross-section (z) view. Bars indicate a length of 40 μ m. (A) *S. oneidensis* strain MR-1 wild-type cells; (B) *omcA* mutant cells; (C) *mtrC* mutant cells.

moting associations with other organisms. Such multiple roles for extracellularly secreted molecules may help explain the metabolic investment of producing these compounds.

Several mechanisms have been proposed for the transfer of electrons from the cytoplasm to insoluble extracellular electron acceptors in Shewanella. One explanation involves direct contact by OM proteins, where OmcA and/or MtrC directly reduces substrates unable to diffuse into the periplasm. Indeed, recent research has shown that purified OM cytochromes can reduce electrodes in vitro (12, 54). Electrically conductive appendages also have been observed in MR-1 cultures under certain conditions and have been proposed to be involved in electron transfer (17). When proteins cannot directly contact metals, siderophore production is a common strategy to increase available soluble iron (38), and MR-1 could use a similar mechanism to make soluble iron available to the cell for respiration (18, 49). Small redox-active molecules also can accept electrons from cells and pass them to insoluble substrates, a process termed electron shuttling (18). The observation that MR-1 can reduce insoluble iron at a distance (25) suggests that chelation and/or electron shuttling are utilized, as these mechanisms do not require contact between MR-1 and the iron surface. The work presented here supports the model of electron shuttling using excreted flavins by MR-1. Chelators could increase rates of insoluble iron reduction at high concentrations, but the effect of chelation on iron oxide reduction near natural flavin concentrations appears to be unimportant (Fig. 1). Consistently with this observation, the addition of 12 µM citrate did not stimulate reduction rates of resting cells containing 12 µM AQDS (data not shown).

To verify that flavins can act as electron shuttles, we demonstrated that MR-1 reduces extracellular flavins, and this activity is dependent on the electrons gained from carbon source oxidation. We also demonstrated that the reduction of flavins by MR-1 requires components of the Mtr respiratory pathway. The finding that OmcA does not contribute to the flavin reduction rate in incubations with wild-type cells but can partially compensate for the loss of MtrC suggests a level of functional redundancy or modularity between these OM cytochromes. Importantly, flavins stimulated insoluble iron reduction more effectively than AQDS, suggesting that MR-1 has evolved to use flavins for electron shuttling and that acceleration seen from other exogenous shuttles is a consequence of pathway promiscuity (18).

Though strains lacking OmcA had no defect in flavin reduc-

tion in washed-cell incubations, by standardizing for levels of attached protein, controlling for external flavin concentrations, and maintaining a constant electrode potential, we identified a new phenotype for this mutant. Lower current production (per unit of surface area) by strains could be ascribed to both a lower current production rate per cell and a lower degree of attachment by $\Delta omcA$ strains. These experiments highlighted the importance of accounting for all variables when using electrodes as electron acceptors. OmcA previously was thought to be only a terminal reductase of metals, which was supported by in vitro biochemical evidence that it can reduce various forms of iron (46, 47, 56). OmcA previously was shown to bind insoluble forms of iron in vitro with a high affinity (56), a greater affinity than that of MtrC (28). We suggest that while OmcA plays a small role in electron transfer in vivo, it also is involved in cellular attachment to solid surfaces or other cells. Our observation is consistent with atomic force microscope experiments demonstrating the unfolding of proteins in the size range of OmcA and MtrC on the outer surface of S. oneidensis bound to iron oxide-coated cantilevers (29). Indeed, mutants lacking OmcA have a marked deficiency in the reduction of insoluble substrates such as electrodes (Fig. 4A) and manganese oxide (36) but not for the reduction of soluble substrates like flavins (Fig. 3), cobalt(III)-EDTA (21), and ferric citrate (8, 36).

Another observation made in this study is that strains lacking MtrC attach to electrodes but produce only 10% of the wild-type current when expressed per unit of attached protein (Fig. 4C). This conflicts with the observation that cell suspensions of $\Delta mtrC$ are able to reduce flavins at nearly half the rate of MR-1 (Fig. 3). Several factors may account for this discrepancy; most important is the fact that flavin reduction assays were performed for a short period of time using resting cells, while all electrode experiments require continuous growth and colonization by the strains being tested. Another difference between these assays was the concentration of flavins used: 250 nM in the bioreactor and as much as 120 µM in the resting-cell experiments (which were used to obtain a robust signal for rate calculations). OmcA may replace MtrC more effectively under conditions of high flavin concentrations. It also is clear that a majority of $\Delta mtrC$ cells simply were dead (or at least unable to maintain a sufficient membrane potential), although they remained attached to the electrode surface (as shown via confocal microscopy) (Fig. 5), diluting the per-cell respiration rates of $\Delta mtrC$. Regardless, the respiration rates observed for wildtype resting cells (15 relative fluorescence units/s = \sim 4 nmol electrons/mg protein/s) were similar to those observed for colonized electrodes (0.8 μ A/ μ g = \sim 8 nmol electrons/mg protein/s). Thus, our resting-cell assay accurately reflected the maximal rates of electron transfer in both cases.

The availability of OmcA to promote electrode interaction also may be impaired in cells lacking MtrA (D. Richardson, personal communication) and MtrB (32), as strains lacking MtrB mislocalize the outer membrane proteins MtrC and OmcA. MtrC mutants are able to reduce soluble substrates such as ferric citrate and cobalt(III)-EDTA much more efficiently than mutants lacking MtrA, MtrB, or both OmcA and MtrC (5, 6, 8, 21). If the same trend follows for electrode reduction, cells lacking MtrC may be able to respire enough to begin the colonization of the electrode, whereas $\Delta mtrA$, $\Delta mtrB$, and $\Delta omcA \ \Delta mtrC$ cannot. The defect in the electrode reduction of strains lacking MtrA and MtrB could be linked to the fact that these proteins form a complex with MtrC in MR-1 (43). Thus, if any of the components are missing, the entire complex could be inactivated or mislocalized. Overall, this model could explain the observation that while mutant strains lacking OmcA are deficient only in solid substrate reduction, MtrC mutants are defective in the reduction of both soluble and insoluble substrates (8, 9, 35, 36).

We recently demonstrated that OM cytochromes can reduce electrode surfaces in the absence of exogenous flavins, although these rates are dramatically lower than flavin-assisted rates (4), a finding that corroborates experiments that showed that flavins were needed to accelerate the turnover of purified membrane cytochromes to explain observed reduction rates of whole cells (42). While purified cytochromes (MtrC and OmcA) are able to reduce electrodes in the absence of electron shuttles (12, 20), under cell growth conditions, the majority of the current produced in three electrode bioreactors requires the presence of flavins and is critically dependent on the Mtr pathway (Fig. 4). Both direct contact and electron-shuttling strategies absolutely depend on close cell surface contact, because electron shuttles are predicted to be effective only across very short distances ($\sim 1 \mu$ m) (40).

We have shown in this study that the Mtr pathway is essential for growth on electrodes and accounts for at least 90% of flavin reduction activity. The specific contribution of flavin shuttling in the reduction of iron oxide, manganese oxide, and many other insoluble substrates by *Shewanella* still has not been quantified and will require mutants or isolates unable to actively secrete flavins. While previous work suggests that the Mtr pathway accounts for most of the insoluble substrate reduction in *S. oneidensis*, it remains unknown if these studies were monitoring deficiencies in direct iron/manganese/electrode reduction (5, 6, 9, 11, 37, 44) or simply reflecting deficiencies in flavin reduction by these mutant strains.

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