Molecular Underpinnings of Nitrite Effect on CymA-Dependent Respiration in *Shewanella oneidensis*

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**Keywords:** Shewanella, nitrite, regulation, inhibition, respiration

Shewanella exhibit a remarkable versatility of respiration, with a diverse array of electron acceptors (EAs). In environments where these bacteria thrive, multiple EAs are usually present. However, we know little about strategies by which these EAs and their interaction affect ecophysiology of *Shewanella*. In this study, we demonstrate in the model strain, *Shewanella oneidensis* MR-1, that nitrite, not through nitric oxide to which it may convert, inhibits respiration of fumarate, and probably many other EAs whose reduction depends on quinol dehydrogenase CymA. This is achieved via the repression of cyclic adenosine monophosphate (cAMP) production, a second messenger required for activation of cAMP-receptor protein (Crp) which plays a primary role in regulation of respiration. If nitrite is not promptly removed, intracellular cAMP levels drop, and this impairs Crp activity. As a result, the production of nitrite reductase NrfA, CymA, and fumarate reductase FccA is substantially reduced. In contrast, nitrite can be simultaneously respired with trimethylamine N-oxide, resulting in enhanced biomass.

**INTRODUCTION**

*Shewanella oneidensis* MR-1 is a Gram-negative facultative anaerobe with remarkable respiration abilities that permit the use of a diverse array of terminal electron acceptors (EAs), including fumarate, nitrate, nitrite, thiosulfate, trimethylamine N-oxide (TMAO), dimethylsulfoxide (DMSO), Fe(III), Mn(III) and (IV), Cr(VI), and U(VI), among others (Fredrickson et al., 2008). As anticipated there are a large number of respiratory pathways, some of which have been elucidated over the past two decades (Gon et al., 2002; Maier et al., 2003; Gralnick et al., 2006; Cruz-Garcia et al., 2007; Gao et al., 2009; Shirodkar et al., 2011; Shi et al., 2012). For nitrate and nitrite respiration, most bacteria use the quinol dehydrogenases NapC and NrfBCD (or NrfH) to transfer electrons to the terminal reductases NapA and NrfA, respectively. In contrast, *S. oneidensis* lacks these dehydrogenases (Jepson et al., 2006; Gao et al., 2009; Simpson et al., 2010; Fu et al., 2014). Instead, CymA, a cytoplasmic-membrane-bound cytochrome c, is recruited to function for the missing proteins (Marriott et al., 2012). The role of CymA could be partially, but significantly less effectively, fulfilled by the \( b_{c_1} \) complex and SirCD, which primarily function as quinol dehydrogenases to transport electrons to the cytochrome \( cbb_3 \) oxidase and to sulfite reductase, respectively (Cordova et al., 2011; Zhou et al., 2013; Fu et al., 2014). Interestingly, only two components are needed for the formation of functional NAP and NRF complexes, CymA-NapA and CymA-NrfA (Gao et al., 2009). One consequence of sharing CymA is that reduction of...
nitrite to ammonium by NrfA does not commence until nitrate is thoroughly exhausted, resulting in a characteristic two-step reduction of nitrate (Gao et al., 2009).

In addition to being EAs for respiration, nitrate (as a precursor to nitrite) and nitrite have been used for centuries as preservatives in meat products to inhibit growth of bacterial pathogens. The antimicrobial action of nitrite is generally attributed to the formation of nitric oxide (NO), which interferes with protein cofactors, such as Fe–S clusters, heme, and lipoamide, or promotes the formation of reactive nitrogen species (Reddy et al., 1983; Hyduke et al., 2007; Mason et al., 2009; Richardson et al., 2011). In S. oneidensis, the cytochrome cbb3 oxidase, the enzyme complex predominantly responsible for oxygen respiration, is the primary target of nitrite stress (Fu et al., 2013; Zhou et al., 2013; Yin et al., 2015). This corresponds to the fact that the Escherichia coli cytochrome bo terminal oxidase is highly susceptible to NO (Mason et al., 2009).

When nitrite is added to S. oneidensis cultures grown under aerobic conditions, cells are constantly under nitrite threat until entry into the stationary phase, when reduction of nitrite to non-harmful ammonium ion occurs (Dong et al., 2012; Zhang et al., 2013). In contrast, when oxygen is absent, nitrite can be immediately consumed as an EA, leading to a rapid decrease in its concentrations (Gao et al., 2009). Despite this, levels at which nitrite completely prevents growth are ~25 and ~5 mM for aerobic and anaerobic growth (nitrite as sole EA) respectively (Dong et al., 2012; Zhang et al., 2013), implying that nitrite is more toxic anaerobically than aerobically. One handy explanation is that nitrite is converted to NO faster under anoxic conditions. In parallel, nitrite may be attacking a cell process that is not, or less, necessary for aerobic growth.

In this study, we examined the role of nitrite during anaerobiosis and found that inhibition of growth on fumarate, probably many other CymA-dependent EAs, by nitrite is associated with cyclic adenosine 3′,5′-monophosphate (cAMP). In bacteria, the physiological function of cAMP signaling, which has been extensively studied for decades, is diverse (Green et al., 2014). For regulation of metabolism, cAMP as an effector forms a regulatory system with cAMP receptor protein (Crp) to coordinate the allocation of proteomic resources with different metabolic demands in different nutrient environments (You et al., 2013). In S. oneidensis, the CAMP-Crp system is the primary regulator mediating respiration of various EAs, as well as many other biological processes (Saffarini et al., 2003; Dong et al., 2012; Fu et al., 2013; Zhou et al., 2013; Gao et al., 2015). Proper CAMP levels must be maintained, by adenylate cyclases (ACs) and phosphodiesterase for synthesis and degradation respectively (Charania et al., 2009; Yin et al., 2016), because either too much or too little negatively impacts aerobic growth (Zhou et al., 2013; Yin et al., 2016). We further showed that nitrite via a yet unknown mechanism triggers repression of cAMP production, by which nitrite eventually compromises Crp activation, leading to substantial decreases in the production of NrfA, CymA, and the fumarate reductase, FccA.

### MATERIALS AND METHODS

#### Bacterial Strains, Plasmids, and Culture Conditions

The bacterial strains and plasmids used in this study are listed in Table 1. In-frame deletion strains derived from S. oneidensis MR-1 used in this study were constructed and verified in previous reports. Sequences of the primers used in this study are available upon request. E. coli and S. oneidensis were grown aerobically in Lysogeny broth (LB, Difco, Detroit, MI, USA) at 37 and 30°C for genetic manipulation. When appropriate, the growth medium was supplemented with the following: 2, 6-diaminopimelic acid (DAP), 0.3 mM; ampicillin, 50 μg/ml; kanamycin, 50 μg/ml; gentamycin, 15 μg/ml; and streptomycin, 100 μg/ml. All chemicals were obtained from Sigma-Aldrich (St. Louis, MO, USA) unless otherwise noted.

Growth of S. oneidensis strains under aerobic or anaerobic conditions was measured at 600 nm (OD600). MS defined medium containing 30 mM lactate as electron donor was used as previously described (Shi et al., 2015). For aerobic growth, mid-log phase cultures (~0.2 of OD600) were inoculated into fresh medium to ~0.02 and shaken at 200 rpm at 30°C. For anaerobic growth, mid-log phase aerobic cultures were pelleted by centrifugation, purged with nitrogen, suspended in fresh medium prepared anaerobically to an OD600 of ~0.02. EAs used in this study include nitrite (2 mM), fumarate (20 mM), and TMAO (20 mM). To assay the effect of NO on growth, DETA NONOate (t1/2, 20 h at 37°C and 56 h at 25°C) was used because it releases NO slowly and can maintain a relatively steady NO concentration (Zhang et al., 2013). For NO scavenging, carboxy-PTIO (Invitrogen, Carlson, CA, USA) was added to a final concentration of 0.1 mM. For chemical complementation, cAMP of various levels was used.

#### Controlled Expression of cymA, fccA, nrfA, and scyA Genes

To assess effects of the four genes expressed at varying levels on nitrite-associated physiology, we placed each of them under the control of isopropyl-β-D-thiogalactopyranoside (IPTG)-inducible promoter P_{lac} within pHGE-P_{lac} (Luo et al., 2013). After verification by sequencing, the vectors were transferred into the relevant strains via conjugation. Cells carrying vectors of interest were grown in media indicated in the text and/or figure legends in the presence of IPTG of varying levels.

#### Chemical Assays

Concentrations of nitrite in culture supernatants were measured by a modified Griess assay and quantitated spectrophotometrically at 540 nm (Miranda et al., 2001). Intracellular cAMP concentrations were measured using a commercially available kit (cAMP direct immunoassay kit; BioVision) essentially the same as described before (Yin et al., 2016). Standard curves were made with commercial agents each time.
Promoter Activity Assay

The activity of various promoters was assessed using a single-copy integrative lacZ reporter system as described previously (Fu et al., 2014). A fragment containing the sequence upstream of each operon from −300 to +1 (relative to the translation start codon) was amplified and cloned into the reporter vector pHGEI01 and verified by sequencing. These plasmids were then transferred by conjugation into relevant S. oneidensis strains. Plasmid pHGEI01 containing promoters of interest integrates into the chromosome and the antibiotic marker is then removed by an established approach (Fu et al., 2013, 2014). Cells grown to the mid-log phase were collected and β-galactosidase activity assays were performed with an assay kit as described previously (Jin et al., 2013).

Expression of GFP Fusions and Quantitation of Fluorescence

To validate protein overproduction driven by the IPTG-inducible Ptac, constructs expressing GFP protein were prepared. After verification by sequencing, the vectors were moved into relevant S. oneidensis strains by conjugation. Expression of GFP fusions was visualized using a confocal microscope as described previously (Luo et al., 2013). For quantitation, mid-log phase cultures were collected, washed with phosphate-buffered saline containing 0.05% Tween 20, and resuspended in the wash buffer to an OD600 of 0.1. 100 µl of the cell suspensions were transferred into black 96-well plates at various time intervals and fluorescence was measured using a fluorescence microplate reader (M200 Pro Tecan) with excitation at 485 nm and detection of emission at 515 nm.

SDS-PAGE, Heme-Staining, and Immunoblotting Assays

Unless otherwise noted, mid-log phase cells were harvested, washed with phosphate buffered saline (PBS), resuspended in the same buffer, and sonicated. Protein concentrations of the cell lysates were determined by the bicinchoninic acid assay (Pierce Chemical). The cell lysates were resolved by SDS-PAGE using 12% polyacrylamide gels and stained with 3,3′,5,5′-tetramethylbenzidine (TMBZ) as described elsewhere (Thomas et al., 1976). Immunoblotting analysis was performed essentially as previously described (Dong et al., 2012). Proteins separated by SDS-PAGE were electrophoretically transferred to polyvinylidene difluoride (PVDF) membranes according to the manufacturer’s instructions (Bio-Rad). The gels were blotted for 2 h at 60 V using a Criterion blotter (Bio-Rad). The blotting membrane was probed with a rabbit polyclonal antibody against S. oneidensis Crp (Dong et al., 2012). Goat anti-rabbit IgG-HRP (horseradish peroxidase; Roche Diagnostics) was used as the secondary antibody (1:5,000) and the signal was detected using an enhanced chemiluminescence kit (Thermo Scientific). 

Table 1: Strains and plasmids used in this study.

| Strain or plasmid               | Description                                                                 | Reference or source            |
|--------------------------------|-----------------------------------------------------------------------------|--------------------------------|
| **Strain**                     |                                                                             |                                |
| Escherichia coli               |                                                                             |                                |
| DH5α                           | Host for cloning                                                            | Lab stock                      |
| WM3064                         | Donor strain for conjugation, ΔdapA                                         | W. Metcalf, UIUC               |
| Shewanella oneidensis          |                                                                             |                                |
| MR-1                           | Wild type                                                                   | Lab stock                      |
| HG0970                         | ΔfocA derived from MR-1                                                     | Gao et al., 2010a              |
| HG3286-4                       | ΔcysA derived from MR-1                                                     | Chen et al., 2015              |
| HG3980                         | ΔnfrA derived from MR-1                                                     | Gao et al., 2009               |
| HG3982                         | ΔnarP derived from MR-1                                                     | Dong et al., 2012              |
| HG4961                         | ΔcymA derived from MR-1                                                     | Gao et al., 2009               |
| **Plasmid**                    |                                                                             |                                |
| pHGC01                         | Integrative vector for complementation                                        | Fu et al., 2013                |
| pHGEI01                        | Integrative E. coli lacZ reporter vector                                     | Fu et al., 2014                |
| pHBE-Cre                       | Helper vector for antibiotic marker removal                                  | Fu et al., 2013                |
| pHGE-Ptac                      | Km′, IPTG-inducible Ptac expression vector                                   | Luo et al., 2013               |
| pHGE-Ptac-cymA                 | Inducible expression of cymA                                                | This study                     |
| pHGE-Ptac-fccA                 | Inducible expression of fccA                                                | This study                     |
| pHGE-Ptac-nfrA                 | Inducible expression of nfrA                                                | This study                     |
| pHGE-Ptac-gfp                  | Inducible expression of gfp                                                 | This study                     |
| pHGE-Ptac-scyA                 | Inducible expression of scyA                                                | This study                     |
| pHGEI-Popr-lacZ                | E. coli lacZ under control of crp promoter                                  | This study                     |
| pHGEI-PoyA-lacZ                | E. coli lacZ under control of cyaA promoter                                 | This study                     |
| pHGEI-PoyC-lacZ                | E. coli lacZ under control of cyaC promoter                                 | This study                     |
| pHGEI-PcpdA-lacZ               | E. coli lacZ under control of cpdA promoter                                 | This study                     |
detected using a chemiluminescence Western blotting kit (Roche Diagnostics) in accordance with the manufacturer’s instructions. Images were visualized with a UVP imaging system.

**Nitrite Sensitivity Assay**

*Shewanella oneidensis* strains grown to an OD<sub>600</sub> of ~0.4 were adjusted to approximately 10<sup>7</sup> CFUs/ml, followed by 10-fold serial dilutions. Ten microliters of each dilution was spotted onto LB plates containing 3 mM nitrite. The plates were incubated at 30°C before being read. The assays were repeated at least three times with similar results.

**Other Analyses**

LC/MS/MS analyses of excised proteins were carried out essentially as described previously (Sun et al., 2013). Student’s t-test was performed for pairwise comparisons. Values are presented as mean ± SD in the relevant figures.

**RESULTS**

**Inhibitory Effect of Nitrite on Anaerobic Growth of *S. oneidensis***

*Shewanella oneidensis* grows extremely poorly on nitrite as sole EA (Zhang et al., 2013). Two mM nitrite facilitates growth most effectively, resulting in an OD<sub>600</sub> up to 0.06, which is barely visible (Cruz-Garcia et al., 2007; Gao et al., 2009). To determine whether this poor growth is a result of nitrite toxicity, we measured growth of *S. oneidensis* on fumarate, DMSO, TMAO, or sulfite in the presence of 2 mM nitrite. While the former two rely on CymA for electrons, the remaining are CymA-independent (Schwalb et al., 2003). As shown in Figure 1A, growth with fumarate was substantially retarded in the presence of nitrite, manifesting that nitrite inhibits fumarate respiration. Inhibition by nitrite was also observed with DMSO, albeit less significantly (Figure 1B). In contrast to fumarate and DMSO, nitrite had a negative impact on the early phase of growth with TMAO or sulfite but higher cell densities were eventually obtained (Figures 1C,D). We reasoned that this difference may be due to the reduction of nitrite. Thus we measured the remaining nitrite in supernatants of cultures grown on nitrite with one of these EAs. Consistent with the growth phenotype, nitrite reduction in cells grown on fumarate or DMSO hardly occurred, whereas nitrite was consumed in the presence of TMAO or sulfite (Figure 1E). In fact, TMAO dramatically promoted reduction of nitrite as the latter was depleted in 4 h. It was worth noting that compared to fumarate and TMAO, DMSO and sulfite were poor EAs to support growth, making it difficult to assess influence of nitrite; hence, we used fumarate and TMAO as EAs for subsequent experiments through this study. As impacts of nitrite on respiration of different EAs vary, the data suggest that nitrite toxicity may not be the only reason explaining inhibition of growth on fumarate or DMSO.

**Nitrite Inhibition of Fumarate Respiration is Not via NO**

Previously we had shown that an *S. oneidensis* strain lacking the cytochrome bd oxidase is highly susceptible to nitrite but not to NO under aerobic conditions, implicating a difference in toxicities of nitrite and NO to cells grown aerobically (Fu et al., 2013; Zhang et al., 2013). Despite this, whether the inhibitory effect of nitrite is due to the production of NO requires further validation given that *S. oneidensis* produces NO endogenously in the presence of either nitrate or nitrite under anaerobic conditions only (Price et al., 2007). To this end, we first examined the growth of *S. oneidensis* on fumarate in the presence of 0.1–0.8 mM DETA NONOate (NO generating agent; Figure 2A). Up to 0.5 mM DETA NONOate did not
Rather it showed a similar negative impact on growth of the overproduction of CymA did not alleviate nitrite inhibition. As mentioned earlier, in addition to terminal reductases the pathways for fumarate and TMAO reduction differ in that they employ distinct quinol dehydrogenases, CymA and TorC, respectively (Schwalb et al., 2003). To evaluate whether CymA may be a target of nitrite inhibition, we placed the cymA gene under the control of the IPTG-inducible promoter \( P_{\text{lac}} \) and determined whether its overexpression suppresses the nitrite inhibition of fumarate-supported growth (Figure 3A). As expected, the \( \Delta \text{cym}A \) strain hardly grew on fumarate (Schwalb et al., 2003). While complete complementation was obtained with 0.01 mM IPTG, but without IPTG addition (low expression as \( P_{\text{lac}} \) is slightly leaky (Luo et al., 2013; Shi et al., 2014; Sun et al., 2014; Fu et al., 2015) or with higher concentrations there was only partial rescue of the growth defect, indicating that CymA in excess has a negative effect on growth (Figure 3A). Importantly, overproduction of CymA did not alleviate nitrite inhibition. Rather it showed a similar negative impact on growth of the \( \Delta \text{cym}A \) strain in the presence of nitrite. These data rule out the possibility that CymA is one of the primary targets of nitrite inhibition (Figure 3B). We then tested whether or not fumarate reductase FccA is inhibited by nitrite using the same protocol. Similar results were obtained, including a dose-dependent complementation and a negative effect of excess FccA with or without nitrite (Figures 3C,D). To test the possibility that the growth defect resulting from CymA or FccA overproduction is a protein-burden effect, we repeated the experiment with the \( \text{gfp} \) gene under the same conditions. Overproduction of GFP did not impair growth (Supplementary Figure S1). Furthermore, Given that both CymA and FccA are \( c \)-type cytochromes we overproduced periplasmic \( c \)-type cytochrome ScyA to determine whether the phenotype is specific to any \( c \)-type cytochrome when in excess. Excess ScyA induced by 0.2 mM IPTG increased nitrite resistance (Supplementary Figure S1), which is in agreement with a previous study (Yin et al., 2015). However, it did not inhibit growth on fumarate in the absence of nitrite (Supplementary Figure S1), indicating that growth inhibition by excess CymA or FccA may not be applicable to other \( c \)-type cytochromes. Overall, these data indicate that neither CymA nor FccA is directly inhibited by nitrite.

Neither CymA nor the Fumarate Reductase Is Inhibited by Nitrite

Experiments were performed at least three times with error bars representing the standard deviation.

Nitrite Alters the Content of \( c \)-Type Cytochromes

The respiratory diversity of \( S. \text{oneidensis} \) is, at least in large part, due to its abundant \( c \)-type cytochromes, which confer to colonies or cell pellet an orange color (Meyer et al., 2004; Gao et al., 2010b). We noticed that the color of cell pellets grown on fumarate in the presence of nitrite was significantly lighter (Figure 4). Interestingly, this phenomenon was neither observed with either as sole EA nor with the TMAO/nitrite pair, implying that nitrite may reduce cytochrome \( c \) production in cells grown with fumarate but not TMAO. To test this possibility, we assayed heme \( c \) levels (Figure 4). The amount of total \( c \)-type cytochromes decreased significantly (to \( \sim 54\% \) relative to the non-treated control) by including nitrite with fumarate but was not significantly altered with TMAO.

We then compared by SDS-PAGE the cytochrome \( c \) profiles of extracts of cells grown on fumarate or TMAO with or without nitrite and found that the overall amounts of \( c \)-type cytochromes were generally consistent with the pellet colors.
and the heme c levels (Figure 4). Three bands predicted to be NrfA, TorC, and CymA based on previous studies (Dos Santos et al., 1998; Myers and Myers, 2000; Dong et al., 2012) were excised, digested, and confirmed using mass spectrometry (MS). While TorC was visible as long as TMAO was present, NrfA showed a significantly different pattern. NrfA was evident in extracts of cells grown on nitrate, nitrite, or TMAO with nitrite but significantly weak in cells grown on TMAO only (Dong et al., 2012; Zhang et al., 2013). More importantly, this band was barely visible from extracts of cells grown on fumarate even in the presence of nitrite, suggesting that NrfA production is somehow blocked when both fumarate and nitrite are present.

**Reduced Production of c-Type Cytochromes in the Presence of Nitrite Is Partially Due to Reduced Crp-cAMP Activity**

Previously, we had found that cells lacking global regulator Crp produced c-type cytochromes at a level similar to that for cells grown fumarate plus nitrite (Gao et al., 2010b), implying a link between nitrite treatment and Crp regulation. Therefore, we monitored expression of the crp gene with a lacZ-reporter. The crp promoter activity did not alter significantly in cells grown on fumarate with nitrite compared to those grown on just fumarate or for cells with TMAO in the absence or presence of nitrite (Figure 5A, left panel). Western blotting analysis of Crp revealed a similar result with a crp null mutant as control (embedded in Figure 5A). Given that the activity of Crp depends on formation of a Crp-cAMP complex (Charania et al., 2009; Zhou et al., 2013), we then assayed cAMP levels in these cells (Figure 5A, right panel). Compared to cells grown on fumarate in the absence of nitrite (which was set to 100%), those grown on TMAO with or without nitrite had significantly greater cAMP levels. In contrast, there was a pronounced decrease in cAMP in cells grown on fumarate with nitrite. These results suggest that the inhibitory effect of nitrite on growth in the presence of fumarate is, at least in part, due to a compromised Crp activity primarily by reducing intracellular cAMP levels.
and phosphodiesterase, respectively (Botsford and Harman, 1992; Imamura et al., 1996). In *S. oneidensis*, there are three functional ACs, CyaA, CyaB, and CyaC, of which CyaC is the major AC for cAMP production and CyaA contributes slightly whereas CpdA is the only cAMP phosphodiesterase (Charania et al., 2009; Yin et al., 2016). To reveal direct factors for lowered cAMP levels in the presence of nitrite, we monitored expression of cya (cyaA and cyaC) and cpdA genes in the wild-type cultures grown on fumarate or TMAO w/o nitrite. As shown in Figure 5D, expression of cyaC, the major AC gene, but neither cyaA nor cpdA, was substantially repressed in the cells grown on both fumarate and nitrite, suggesting that the reduction in the cAMP amount is likely due to decreased production.

**Differential Expression of nap and nrf Operons in the Presence of Nitrite**

As shown above (Figure 4), nitrite is able to induce expression of *nrfA* in cells grown on TMAO but not on fumarate. To determine how reduced activity of the cAMP-Crp complex affects expression of relevant genes, we assayed their promoter activities by using an integrative *lacZ* reporter system (Fu et al., 2014). In the either absence or presence of nitrite, *nrfA* expression was extremely low in cells grown on fumarate but was induced more than threefold by nitrite in cells grown on TMAO (Figure 6A). In the case of *cymA* and *fccA* genes, both were downregulated by nitrite (Figure 6A). While the *fccA* gene had an approximately 70% decrease in expression, the expression reduction of the *cymA* gene in the presence of nitrite was less extensive (Figure 6A). We could not assess directly influence of the Crp loss on expression of these genes because the *Δcrp* strain could not grow on fumarate. As an alternative, we examined effect of cAMP addition on expression of these genes in cells grown on fumarate or TMAO w/o nitrite. Exogenous cAMP enhanced expression of the *nrfA*, *fccA*, and *cymA* genes in fumarate-grown cells when nitrite was present, which is in agreement with growth restoration by cAMP as shown in Figures 5 and 6A. Furthermore, these observations were supported in general by heme-staining data presented in Figure 6B, which included a verified *nrfA* null mutant (*ΔnrfA*) as negative control (Gao et al., 2009).

Given that forced expression of the *fccA* gene could not correct the growth defect in media containing fumarate and nitrite (Figure 3D), we reasoned that reduced production of NrfA underlies the growth defect. To test this, we first examined growth of the *ΔnrfA* strain on both fumarate and nitrite (Gao et al., 2009). Expectedly, it had further worsened growth than the wild-type (Figure 6C). We then introduced the *nrfA* gene placed under the control of *P_{lac}* promoter into the wild-type and assayed growth. The effect was obvious. Without IPTG, growth was much improved because of expression from the leaky promoter (Figure 6C). In the presence of IPTG of 0.01 mM, with fumarate and nitrite cells carrying the construct displayed a growth pattern similar to that observed from cells grown on TMAO and nitrite (Figure 6C). These data thus confirm that the growth defect was due to low production of NrfA.

![Figure 4](https://example.com/figure4.png) **Figure 4** Nitrite reduces levels of c-type cytochromes in cells grown on fumarate. Mid-log phase cultures (approximately 0.2 of OD_{600} grown on either fumarate or TMAO were pelleted and photographed (upper panel). Cell samples were lysed and the heme c levels were determined (middle panel). The average amount of heme c from the fumarate-grown samples without nitrite was set to 100%. Proteins (10 μg per lane) extracted from the indicated samples (with or without 2 mM nitrite) were resolved by SDS-PAGE and analyzed by heme staining (lower panel, a composite image in which a marker is placed next to the gel). NrfA of 2 μM was used. NitrA (2 μM) was included for comparison. The positions of NrfA, TorC, and CymA were marked. All experiments were performed at least three times with standard deviations presented as error bars or similar results were obtained.

To further confirm the involvement of the Crp-cAMP complex in the process, we examined effect of exogenous cAMP on cytochrome c levels and growth on fumarate or TMAO with nitrite. As shown in Figure 5B, the amounts of total c-type cytochromes in cells grown on TMAO and nitrite in the presence of cAMP up to 20 μM were similar but increased significantly (to ~138% relative to the nitrite-free control) when cAMP was supplemented to 50 μM. In contrast, the cAMP addition proportionally enhanced c-type cytochrome production in cells grown on fumarate plus nitrite, with restoration to the wild-type level being achieved by 20 μM. We then assessed whether exogenous cAMP also helps growth in the presence of nitrite (Figure 5C). Indeed, cAMP at 20 μM greatly improved growth of the wild-type strain on both fumarate and nitrite (Figure 6A). As an alternative, we examined effect of cAMP addition on expression of these genes in cells grown on fumarate or TMAO w/o nitrite. Exogenous cAMP enhanced expression of the *nrfA*, *fccA*, and *cymA* genes in fumarate-grown cells when nitrite was present, which is in agreement with growth restoration by cAMP as shown in Figures 5 and 6A. Furthermore, these observations were supported in general by heme-staining data presented in Figure 6B, which included a verified *nrfA* null mutant (*ΔnrfA*) as negative control (Gao et al., 2009).

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![Figure 5](https://example.com/figure5.png) **Figure 5** Nitrite reduces levels of c-type cytochromes in cells grown on fumarate. Mid-log phase cultures (approximately 0.2 of OD_{600}) grown on either fumarate or TMAO were pelleted and photographed (upper panel). Cell samples were lysed and the heme c levels were determined (middle panel). The average amount of heme c from the fumarate-grown samples without nitrite was set to 100%. Proteins (10 μg per lane) extracted from the indicated samples (with or without 2 mM nitrite) were resolved by SDS-PAGE and analyzed by heme staining (lower panel, a composite image in which a marker is placed next to the gel). NrfA of 2 μM was used. NitrA (2 μM) was included for comparison. The positions of NrfA, TorC, and CymA were marked. All experiments were performed at least three times with standard deviations presented as error bars or similar results were obtained.
FIGURE 5 | Effect of exogenous cAMP on levels of c-type cytochromes and growth. (A) Effect of nitrite on the expression of the crp gene (left panel) and cAMP levels (right panel) in cells grown on fumarate, fumarate and nitrite, TMAO, as well as TMAO and nitrite. Expression was assayed by using an integrative lacZ reporter as described in Section “Materials and Methods.” Results of western blotting of Crp were embedded with Δcrp as negative control (left panel). The average levels of expression (left panel) and cAMP (right panel) from the fumarate samples without nitrite were set to 100%. Cells were prepared as described in Figure 4. (B) Mid-log phase wild-type cultures under indicated conditions were collected and processed for determination of levels of c-type cytochromes as in Figure 4. (C) Growth of the wild-type under indicated conditions. cAMP concentration: 20 µM. Nitrite-free (−) cultures were included for comparison. (D) Expression of the genes involved in cAMP production (cyAC and cyA) and degradation (cpdA) in cells grown on fumarate, fumarate and nitrite, TMAO, as well as TMAO and nitrite. All experiments were performed in triplicate and error bars indicate the standard error.

Effects of Nitrite on cAMP Production Is Independent of NarP-NarQ

In S. oneidensis, the NarP-NarQ two-component system (TCS) senses both nitrate and nitrite, and then regulates transcription of both the nap and nrf operons accordingly (Dong et al., 2012). We then tested whether the nitrite inhibition of cAMP production in cells grown on fumarate depends on NarP-NarQ. A narP null mutant (ΔnarP), which is impaired in nitrite reduction (Dong et al., 2012), was indistinguishable from the wild-type when grown on fumarate and nitrite (Figure 7). In contrast, with TMAO and nitrite, this strain grew substantially slower than the wild-type, a phenotype in agreement with reduced NrfA production. Importantly, addition of exogenous cAMP had no significant effect when the TCS was absent (Figure 7). We then examined expression of the nap and nrf operons in a narP null mutant (Dong et al., 2012). In the ΔnarP strain grown on either fumarate or TMAO, neither the nap nor the nrf operon was induced by nitrite with or without cAMP (data not shown). These results manifest that the TCS is the immediate regulator for the nap and nrf operons in response to nitrite while the cAMP-Crp complex functions at higher levels in the regulatory hierarchy.

DISCUSSION

Nitrite can enter the cytoplasm in the form of undissociated nitrous acid and through specific transporters (Samouilov et al., 2007; Lü et al., 2013). Despite this, it is likely that the majority of its targets reside in the periplasm, where it is at the highest concentrations. This may be particularly important for S. oneidensis as more than 40 c-type cytochromes are presented, either membrane-bound or soluble in the periplasm (Myers and Myers, 2000; Gao et al., 2010a; Maia and Moura, 2014). Nitrite inhibits the cytochrome cbb3 oxidase, the predominant system for oxygen respiration (Fu et al., 2013; Zhou et al., 2013). In addition, it has been reported that nitrite impedes respiration of soluble iron species in S. oneidensis (DiChristina, 1992). Furthermore, nitrite is more inhibitory anaerobically than aerobically and its targets under anoxic conditions remain undefined.
In this study, we aimed to unravel the mechanism of nitrite inhibition during anaerobiosis in *S. oneidensis*. By performing a series of experiments, we generated three important contributions to the current understanding of nitrite-associated physiology. First, we show that NO plays a negligible role in the phenotypes caused by nitrite. This is surprising because the antimicrobial action of nitrite has long been attributed to the formation of NO (Reddy et al., 1983; Hyduke et al., 2007; Mason et al., 2009; Richardson et al., 2011). Second, we present data to suggest that terminal reductases may not be direct targets of nitrite, contrasting the *cbb*$_3$ oxidase for oxygen respiration (Fu et al., 2013). Third, we demonstrate that nitrite inhibition of growth on fumarate, presumably most of CymA-dependent EAs, is a result of reduced cAMP levels, leading to repression of both fumarate reductase and nitrite reductase.

Unlike the *E. coli* Crp which plays a central role in carbon catabolite repression (Görke and Stülke, 2008), the *S. oneidensis* counterpart is the predominant regulator controlling respiration (Saffarini et al., 2003; Gao et al., 2010b). Not surprisingly, a large number of genes involved in respiration including *nrfA, cymA*, and *fccA* are likely to be under direct control of Crp as Crp-binding motifs are predicted to be located in their promoter regions (Saffarini et al., 2003; Schwalb et al., 2003; Gao et al., 2010b; Novichkov et al., 2013). The present results show that nitrite compromises Crp activity by inhibiting production of the major AC CyaC and thereby reducing cAMP levels, resulting in the reduced expression of many *c*-type cytochrome genes.
This notion is reasonable because lose of either two cyclases that are responsible for cAMP production results in a substantial reduction in the activity of Crp (Charania et al., 2009). Moreover, overproduction of Crp could not fully complement a crp mutant (Fu et al., 2013) and cAMP in excess also affects Crp’s activity in S. oneidensis (Fu et al., 2013; Yin et al., 2016). Therefore, a strict stoichiometric ratio of Crp to cAMP molecules appears important for activity of Crp.

With low levels of cAMP, production of both fumarate and nitrite reductases lessens substantially in the presence of both fumarate and nitrite (Figure 8). Although the shortage of the former slows fumarate respiration and limits biomass yield, it is the reduced amount of the latter that is the determining factor for the growth defect, revealed by that the forced production of NrfA rather than FccA is able to correct the defect. In contrast, nitrite can be simultaneously utilized by cells respiring on TMAO, supporting significantly enhanced biomass. Unlike the reduction pathways for fumarate and soluble iron species, the reduction pathways for nitrite seem to be more likely a result of the signal-sensing role as it lacks redox-sensing domains. Unlike the reduction pathways for fumarate and soluble iron species as well as many others, the TMAO pathway does not use CymA as a quinol dehydrogenase (Marritt et al., 2012). Instead TMAO reduction depends on TorC, which along with the reductase, TorA, is not subject to Crp regulation (Gon et al., 2010). Apparently, with TMAO supporting growth, intracellular cAMP levels can be maintained sufficiently high for NrfA production, allowing rapid removal of toxic nitrite.

The cAMP-Crp complex functions as an activator for both the nap and nrf operons, and so does NarP-NarQ TCS. The data presented here is in agreement with the previous proposal that the TCS is the immediate regulator for the nap and nrf operons while the cAMP-Crp complex functions at higher levels in the regulatory hierarchy (Dong et al., 2012). Intriguingly, although NarP-NarQ senses and responds to nitrite, it is not required for the role of nitrite in cAMP production. Exactly how nitrite triggers repression of cAMP production is not known.

How nitrite triggers reduction of cAMP biosynthesis remains unknown. As a matter of fact, despite intensive studies for decades we still know little about bacterial ability to maintain cAMP homeostasis in response to environmental and metabolic signals (Green et al., 2014). Crp has been a global protein under extensive investigation in the field of global transcription machinery engineering (gTME; Geng and Jiang, 2015). Numerous E. coli crp mutants showing improved performance under various stressful conditions, but the predominant mechanism for the elevated resistance is attributed to altered transcriptional profile, and stresses per se do not serve as a signal to mediate activity of cAMP-Crp (Geng and Jiang, 2015). Clearly, nitrite is different. We have previously suggested that regulation by cAMP-Crp is particularly critical in adaptation of S. oneidensis to redox-stratified environments. Although the major AC CyaC resides in the inner-membrane and is accessible to nitrite (Charania et al., 2009), nitrite may not directly inhibit the enzyme because the lowered cAMP concentration caused by nitrite seems to be more likely a result of reduced CyaC production (Figure 8). Nevertheless, this possibility will be tested in future. Crp is unlikely to be able to assume the signal-sensing role as it lacks redox-sensing domains (Fu et al., 2013; Zhou et al., 2013). Therefore, there must be some factors that sense redox signals (different EAs) and subsequently mediate cAMP levels (Figure 8). Identification of such factors is undoubtedly a significant challenge. Nevertheless, our data indicate that nitrite could serve as a good signal for unraveling the signal transduction cascade, which is under way.

Intriguingly, both CymA and FccA display a dose-dependent impact on cells grown on fumarate. FccA is a unique fumarate reductase located in the periplasm, and CymA is responsible for electron transfer to FccA from the quinol pool (Leys et al., 1999). Although, further investigation is needed, a recent finding offers a feasible explanation (McMillan et al., 2013). It is known that the quinol dehydrogenase activity of CymA is triggered by formation of a CymA-FccA complex. Respiratory electron flux relies to some extent on the stability of the complexes. Without FccA, CymA could only reduce the quinol species, menaquinol-7. Hence, the formation of...
the complex regulates the activity of CymA by changing the direction of electron flow. Accordingly, we speculate that optimum respiration depends on a certain stoichiometric ratio of CymA to FccA molecules and overproducing either would undermine the interaction, leading to the impaired respiration of fumarate.

AUTHOR CONTRIBUTIONS

Jie Yuan and HG conceived the idea and designed the project. MJ, HF, and Jianhua Yin carried out the experiments. MJ and HG analyzed data. MJ and HG wrote the paper.

ACKNOWLEDGMENTS

This research was supported by National Natural Science Foundation of China (31270097, 41476105), and the Fundamental Research Funds for the central Universities (2015FZA6001, 2016FZA6003).

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: http://journal.frontiersin.org/article/10.3389/fmicb.2016.01154

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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