



Silver and Copper Acute Effects on Membrane Proteins and Impact on Photosynthetic and Respiratory Complexes in Bacteria

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ABSTRACT Silver (Ag+) and copper (Cu+) ions have been used for centuries in industry, as well as antimicrobial agents in agriculture and health care. Nowadays, Ag+ is also widely used in the field of nanotechnology. Yet, the underlying mechanisms driving toxicity of Ag+ ions in vivo are poorly characterized. It is well known that exposure to excess metal impairs the photosynthetic apparatus of plants and algae. Here, we show that the light-harvesting complex II (LH2) is the primary target of Ag⁺ and Cu⁺ exposure in the purple bacterium Rubrivivax gelatinosus. Ag⁺ and Cu⁺ specifically inactivate the 800-nm absorbing bacteriochlorophyll a (B800), while Ni²⁺ or Cd²⁺ treatment had no effect. This was further supported by analyses of CuSO₄or AgNO₃-treated membrane proteins. Indeed, this treatment induced changes in the LH2 absorption spectrum related to the disruption of the interaction of B800 molecules with the LH2 protein. This caused the release of B800 molecules and subsequently impacted the spectral properties of the carotenoids within the 850-nm absorbing LH2. Moreover, previous studies have suggested that Ag+ can affect the respiratory chain in mitochondria and bacteria. Our data demonstrated that exposure to Ag⁺, both in vivo and in vitro, caused a decrease of cytochrome c oxidase and succinate dehydrogenase activities. Ag+ inhibition of these respiratory complexes was also observed in Escherichia coli, but not in Bacillus subtilis.

IMPORTANCE The use of metal ions represents a serious threat to the environment and to all living organisms because of the acute toxicity of these ions. Nowadays, silver nanoparticles are one of the most widely used nanoparticles in various industrial and health applications. The antimicrobial effect of nanoparticles is in part related to the released Ag⁺ ions and their ability to interact with bacterial membranes. Here, we identify, both *in vitro* and *in vivo*, specific targets of Ag⁺ ions within the membrane of bacteria. This include complexes involved in photosynthesis, but also complexes involved in respiration.

KEYWORDS chlorophyll, copper, membrane complexes, metal homeostasis, photosynthesis, respiration, silver, toxicity

etal accumulation in the environment results in toxicity and defects in metabolism, leading to impaired growth of microorganisms, as well as to a variety of metabolic disorders in higher organisms. In most bacteria, metals such as Cu^+ , Cd^{2^+} , or Ag^+ would diffuse through nonspecific importers within the membrane. This induces the expression of the detoxification systems that allow the cell to tolerate the presence of metals in its environment (1–6). Among these systems, metal efflux systems are very efficient to detoxify excess metal. The P_1B -type ATPases are the most fre-

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quently present heavy metal transporters in bacteria (7). They extrude excess or toxic metal ions such as Cu⁺, Zn²⁺, Cd²⁺, Co²⁺, Pb²⁺, or Ag⁺ from the cytoplasm to the periplasm, where metal is handled by other detoxifying proteins. In Escherichia coli, the Cu+ detoxifying system includes the Cu+ efflux ATPase CopA, the CusFCBA efflux system, and the CueO oxidase (8, 9). These systems are also involved in Ag+ detoxification in E. coli and other species (9-11). In mutants defective in the efflux system, metal accumulation in the cytoplasm can disrupt different metabolic pathways. Indeed, Cu⁺, Ag⁺, or Cd²⁺ can disrupt the solvent-exposed 4Fe-4S clusters of dehydratases (12, 13). In the purple photosynthetic bacterium Rubrivivax gelatinosus, Cu⁺ induces the expression of the CopA-ATPase and the periplasmic blue copper protein CopI (14, 15). Recent in vivo studies showed that Cu+ accumulation in R. gelatinosus and the human pathogen Neisseria gonorrhoeae ΔcopA mutants affects cell growth by altering heme biosynthesis in the cytoplasm (14, 16) or cytochrome c assembly in the periplasm for the $\triangle copl$ mutant in R. gelatinosus (15). Interestingly, similar effect of tellurite on cytochrome c-type assembly was recently reported in Rhodobacter capsulatus (17). Cu+ can also compete with iron for the metal binding site of IscA and inhibit the 4Fe-4S cluster assembly pathway in E. coli (18). In plants and algae, metals exert their toxic action mostly by damaging chloroplasts, which leads to decreased efficiency of photosynthesis. Plants subject to excess metals usually exhibit a decrease in the photosystem amount and chlorophyll content (19-22). However, the toxicity mechanisms are not well known. Assessing the effect of metals on the growth of photosynthetic bacteria can provide new insights into the toxicity mechanisms and identify metal targets in phototrophs. Purple photosynthetic nonsulfur bacteria can grow by aerobic and anaerobic respiration or photosynthetically in the light under anaerobic or microaerobic conditions, using a cyclic electron transport chain. Aerobic respiration usually involves a branched energy-transducing electron transfer chain (23). The cytochrome c-dependent branch usually involves the NADH dehydrogenase, succinate dehydrogenase, the bc_1 complex, and the terminal cytochrome c oxidase (aa_3 or cbb_3). Under light-exposed condition, photosynthesis takes place within the membranous photosynthetic apparatus. The photosystem is usually composed of three pigmentprotein complexes, namely, the two light-harvesting antennae (light-harvesting complex I [LH1] and light-harvesting complex II [LH2]) and the reaction center (RC), associated with carotenoids and bacteriochlorophylls (24). During the process, the light-harvesting complexes (LH) capture light energy and direct it to the RC, where conversion of the excitation energy/charge separation takes place. The LH antenna system consists of two large pigment-protein complexes, the core light-harvesting complex, LH1, that surrounds the RC, and the peripheral light-harvesting complex, LH2, induced under low-light conditions to increase light trapping efficiency in some species. Both LH antennae are composed of two integral membrane polypeptides (α and β) that associate with bacteriochlorophyll (BChI) and carotenoid molecules (25-27). The LH2 antennae contain two spectrally distinct bacteriochlorophylls, a (B800) and B850, which absorb in the near-infrared range, at 800 and 850 nm, respectively. The crystal structure of the LH2 from Rhodopseudomonas acidophila was previously resolved (25). The B850 molecules are sandwiched between the α and β subunits and are perpendicular to the membrane surface. In contrast, the B800 molecules are localized between the β subunits and aligned parallel to the membrane surface. The structures of the RC-LH1 core complexes of Rhodopseudomonas palustris and Thermochromatium tepidum are available (26, 27). In this study, we analyzed the effect of extended exposure to metals on photosynthesis and respiration in the photosynthetic purple bacterium R. gelatinosus. The data indicated that the B800 of LH2 was specifically removed upon exposure to AgNO₃ and CuSO₄. We then assessed the impact on the respiratory chain and showed that metal ions also damaged the succinate dehydrogenase and the terminal cytochrome c oxidase, thereby affecting respiration.



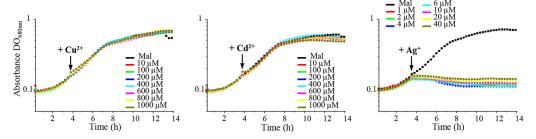


FIG 1 AgNO₃ growth inhibition of *R. gelatinosus*. Wild-type cells were grown in 96-well microplates in the Tecan Infinite M200 luminometer. Indicated concentrations of CuSO₄, CdCl₂, or AgNO₃ were added to the growth medium after 3.5 h, when cells reached an OD_{680} of 0.16 (arrow). Black line represents growth curves of untreated cells (malate) and red, green, blue, cyan, magenta, yellow, and brown lines represent growth curves of cells treated with increasing concentrations of Cu2+, Cd2+, or Ag+, as indicated.

RESULTS

Silver is highly toxic for R. gelatinosus, and the Cu+-ATPase CopA is not involved in Ag+ response. To assess the toxicity of Ag+ in comparison with those of other toxic metals, wild-type cells were treated with increasing concentration of AgNO₂, CuSO₄, or CdCl₂ during the exponential growth phase, and overnight growth was monitored. Growth was not affected by the addition of CuSO₄ or CdCl₂, even at 1 mM. In contrast, addition of 1 μ M AgNO₃ was enough to fully inhibit growth (Fig. 1). Similar results were reported in E. coli cells, highlighting the acute toxicity of Ag+ compared to that of other metal cations (13). We should note that toxicity was reduced when AgNO₃ was added to a higher density of cells (Fig. S1), as previously reported for E. coli. It was suggested that metal ions could interact and be sequestered on the cell surface; the high cell density will therefore affect the dose response (13). To cope with excess toxic metal, bacteria usually induce the genes encoding the metal-efflux ATPases. Ag+ tolerance in E. coli involves the metal efflux P₁B-type ATPase CopA, which translocates Cu⁺ and Ag⁺ from the cytoplasm to the periplasm (4, 9). Therefore, the R. gelatinosus efflux-defective copA mutant was used to check the involvement of CopA in Ag+ efflux. Unlike CuSO₄, which inhibits copA mutant growth (Fig. 2A), no difference in growth inhibition was observed between copA mutant and wild-type cells subjected to excess AgNO₃. These data suggested that in contrast to Ag⁺ tolerance in E. coli, CopA is not involved in Ag+ tolerance in R. gelatinosus. Although the ZntA/CadA ATPase is known to translocate divalent cations, we also checked whether the ΔcadA mutant was sensitive to Ag+ (A. S. Steunou, A. Durand, M.L. Bourbon, M. Babot, S. Liotenberg, and S. Ouchane, submitted for publication). As for the copA mutant strain, no difference in growth was observed between wild-type and $\Delta cadA$ strains in the presence of AgNO₃ (not shown). Cells were also spotted on solid medium supplemented with the same metals. Both copA and ΔcadA mutants showed growth inhibition on 500 μM CuSO₄ and 500 μM CdCl₂, respectively. However, none of the mutants exhibited an altered growth phenotype on 5 μ M AgNO₃ (Fig. 2B). We therefore concluded that the CopA and CadA ATPases were not involved in the AgNO₃ response. To further support this conclusion, we analyzed by Western blot the expression of CopA and CopI in response to metal shock in a strain expressing a His-tagged version of CopA (CopA-H₆) (Fig. 2C). Cells were grown under photosynthetic condition and shocked with CuSO₄ and CdCl₂, known to induce the expression of the Cu⁺ efflux system (Steunou et al., submitted for publication), or with AgNO₃. Untreated cells showed a basal level of CopA and CopI expression because of the presence of 1.6 µM CuSO₄ in the growth medium. As expected, addition of CuSO₄ or CdCl₂ to the growing cells led to significant increases in the amounts of CopA and CopI (Fig. 2C). In contrast, AgNO₃ did not induce the expression of both proteins. Collectively, these results showed that the R. gelatinosus CopA efflux ATPase is not involved in AgNO₃ stress response and detoxification, in contrast to that in E. coli.

Silver and copper excess specifically affected the LH2 complexes in the membrane. To assess the effects of Ag⁺ ions on photosynthesis, cells (optical density at

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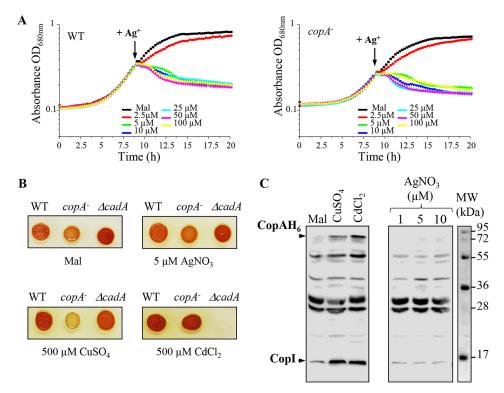


FIG 2 CopA and CadA are not involved in AqNO₃ response. (A) Wild-type (WT) and copA mutant (copA⁻) cells were grown in microplates under microaerobic conditions. Indicated concentrations of AgNO₃ were added to the growth medium after 8.5 h, when cells reached an OD₆₈₀ of 0.3 (arrow). (B) Growth phenotype of the WT, copA and $\triangle cadA$ mutants in the presence of indicated concentrations of AgNO₃, CuSO₄, or CdCl₂ on solid malate media. Cells were grown aerobically for 24 h at 30°C prior to photography. (C) Induction of CopA-H₆ and CopI expression in cells shocked for 1 h with 0.5 mM CuSO₄, 0.5 mM CdCl₂, or 1 to 10 µM AqNO₃. Cells were grown in photosynthetic condition and metals were added to the growth medium when cells reached an OD₆₈₀ of 0.8. Total protein extracts from the same amount of cells (OD $_{680}$ of 0.1) were separated on 14% SDS-PAGE. The proteins were revealed on a Western blot using an HisProbe-HRP.

 $680 \text{ nm } [OD_{680}] = 2)$ grown overnight under photosynthetic condition were treated with 1 mM AgNO₃ and grown further for 2, 10, or 20 h. The bacteriochlorophyll a absorbance in the photosynthetic complexes was measured to monitor changes in response to excess AgNO₃ in the cell in the reaction center and in light-harvesting antenna LH1 complexes and LH2 complexes. The effect of Ag+ on the photosynthetic (PS) complex spectra are presented in Fig. 3A. The B860 (RC-LH1-LH2) and B800 (LH2) wavelength band intensity variations, depending on the length of AgNO₃ exposure, are represented. Ag⁺ induced no apparent effect on the 860-nm band. However, a time exposure-dependent decrease of the B800 band intensity was observed (Fig. 3A). This suggested that the LH2 antennae were affected by AgNO₃ exposure. We also question whether this effect was specific to AgNO₃. For that purpose, cells were also subjected to metal excess stress as described above, but with different metal cations (Fig. 3B). Interestingly, only CuSO₄ caused the same effect as AgNO₃ on the LH2 complexes. Exposure to $\mathrm{CdCl_2}$ or $\mathrm{NiSO_4}$ did not affect the photosynthetic complexes. These data demonstrated that AgNO₃ and CuSO₄ extended exposure affected the LH2 in the photosynthetic membranes.

Silver and copper specifically induced the loss of the 800-nm absorbing bacteriochlorophyll a in LH2. The LH2 antenna (B800 and B850) complexes are spectrally characterized by the 800- and 850-nm absorption bands that arise from the nearinfrared (Qy) transitions of the bacteriochlorophyll a. The loss of the B800 band suggested the loss of the LH2 in the membrane. However, the spectra presented in Fig. 3 also suggested that the LH2 B850 bacteriochlorophyll was not affected, since no shift was observed in the 860-nm band that encompasses the RC (870-nm), the LH1



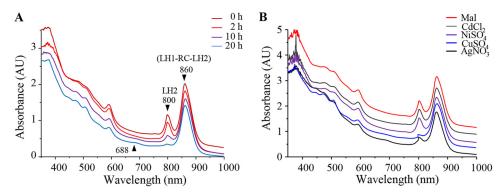


FIG 3 AgNO₃ exposure impact on photosystem in vivo. (A) Spectral analyses of wild type (WT) cells grown overnight by photosynthesis and exposed or not to 1 mM AgNO₃ after they reach an OD_{680} of 2. Spectra (350 to 1,000 nm) were recorded after 2- (light red line), 10- (purple line), or 20-h (blue line) exposure on a double-beam Cary 500 spectrophotometer. (B) Spectral analyses of the WT cells exposed or not to 1 mM CdCl₂ (gray line), NiSO₄ (purple line), CuSO₄ (blue line), or AgNO₃ (black line) after they reached an OD₆₈₀ of 2. Spectra were recorded after 20 h of exposure. Mal (dark red line), untreated cells grown in malate medium.

(875-nm), and the LH2 (850-nm) bands. To confirm this assumption, we compared the spectra of the untreated or AgNO₃-shocked wild-type cells to the spectrum of the pucBA LH2-deficient mutant (28) that only assembles the RC-LH1 core (Fig. 4A). Deletion of the LH2 genes resulted in a significant decrease of the 800-nm band and a substantial red shift of 15 nm (from 860 to 875 nm) of the 860-nm band (28). The resulting peak at 875 nm corresponds to the RC-LH1 core absorption bands. In sharp contrast with the LH2-deficient mutant, the AgNO₃-shocked wild-type cell spectrum showed the decrease of the 800-nm band and no changes in the 860-nm-absorbing bacteriochlorophyll molecules arising from the RC-LH1 core and a modified (B800-free) LH2 (Fig. 4A). Similar impact on LH2 complexes was observed when cells were subjected to CuSO₄ treatment (Fig. S2). These spectra showed that the B800 molecules can selectively be extracted or released from the LH2 complexes in the presence of metals without disrupting the interaction of the LH2 polypeptides with the B850 molecules.

The effect of metals on the RC-LH1 core was also assessed using the pucBA LH2-deficient mutant. Spectra of the exposed cells showed that AgNO₃ and CuSO₄ slightly affected the amount of the RC-LH1 (Fig. 4B). Moreover, total protein lysates from all untreated or treated samples were also loaded onto SDS-PAGE. The Coomassie blue staining showed that all wild-type samples have comparable amounts of LH2 subunits, indicating that the release of B800 molecules did not affect the LH2 protein

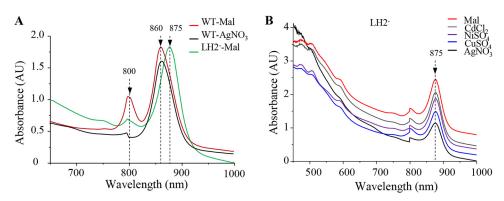


FIG 4 Effect of AgNO₃ on LH2 complexes in vivo. (A) Absorption spectra of wild-type (WT) cells grown overnight by photosynthesis untreated (Mal, red line) or exposed to 1 mM AqNO₃ (black line) in comparison with LH2deficient mutant (LH2-) cells grown overnight by photosynthesis in malate medium (green line). (B) Spectral analyses of the LH2-deficient mutant cells untreated (Mal, red line) or exposed to 1 mM CdCl₂ (gray line), NiSO₃ (purple line), CuSO₄ (blue line), or AgNO₃ (black line) after they reach an OD₆₈₀ of 2. Spectra were recorded after 20 h of exposure.

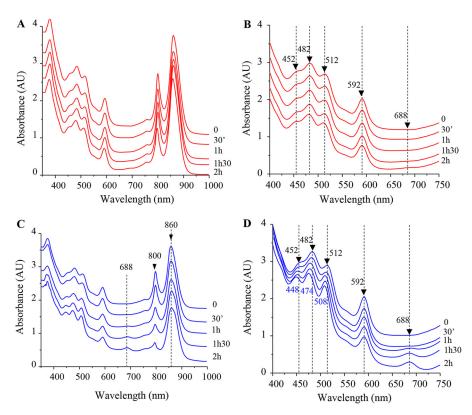


FIG 5 Effect of CuSO₄ on LH2 complexes of isolated membranes. (A) Spectra (350 to 1,000 nm) of untreated membranes were recorded every 30 min. (B) Enlargement of the 400- to 750-nm spectrum absorbance region of untreated membranes. (C) Enriched membrane fractions were mixed with CuSO₄ at 2 mM final concentration; 350- to 1,000-nm spectra were then recorded every 30 min. (D) Enlargement of the 400- to 750-nm spectrum absorbance region, highlighting the shift in the carotenoids bands and the increase in the 688-nm band in the CuSO₄-treated membranes.

stability (Fig. S2). Altogether, these data indicated that AgNO₃- and CuSO₄-induced alterations in the structure of the LH2 complexes, targeting the B800 molecules of the complex. However, this specific and rather limited effect on B800 and the LH2 could not explain the drastic growth inhibition by Ag+, suggesting that Ag+ affects other crucial components or complexes of the cell.

Metal-specific impact on the 800-nm band attested by the release of bacteriochlorophyll and shift in carotenoid absorbance. In the LH2 structure of Rhodopseudomonas acidophila, the 800-nm absorbing bacteriochlorophyll a molecules lie between the β -apoprotein helices, where phytyl moieties interact with the carotenoids. The structural data showed that at least one of the carotenoid molecules makes close Van der Waals contacts with the B800 pigment (25, 29). We therefore assumed that the release of B800 molecules following metal stress should also impact the B800carotenoid interaction. To verify this assumption, enriched membranes from wild-type cells were incubated in phosphate buffer supplemented or not with 2 mM CuSO₄ or AgNO₃. Spectra were then recorded every 30 min to monitor the effect of metals on photosynthetic complexes on isolated membranes (Fig. 5). For untreated membranes, no changes in the amount or in the spectral properties of the RC, LH1, and LH2 were observed (Fig. 5A and B). However, in the membranes subjected to CuSO₄ treatment, a significant decrease in the intensity of the 800-nm band was observed in association with an absorption increase at 688 nm (Fig. 5C and D). This later absorption peak very likely arose from oxidized Bchl in solution, as previously reported (30). Furthermore, with extended exposure to CuSO₄, a shift was also observed in the carotenoid absorption region. Indeed, untreated proteins exhibited three peaks at 452, 482, and 512 nm (Fig. 5B), while CuSO₄ treatment resulted in a shift of the carotenoid absorbance to 448,



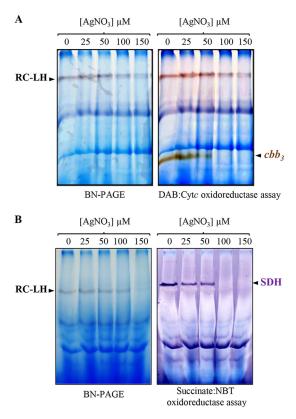


FIG 6 AgNO₃ effect on respiratory complexes in vivo. The wild-type (WT) cells were grown under microaerobic respiratory condition and shocked for 1 h with increasing concentration of AgNO₃. DDMsolubilized membrane proteins were separated on a 5% to 12% BN-PAGE. (A) cbb₃ cytochrome c oxidase DAB:Cytc in-gel activity assay. (B) Succinate dehydrogenase (SDH) in-gel activity assay.

474, and 508 nm (Fig. 5D). Similar effects were obtained in the presence of AgNO₃ treatment (Fig. S3). Thus, we concluded that in the presence of CuSO₄ or AgNO₃, changes in the LH2 absorption spectrum are related to the disruption of the interaction of B800 molecules with LH2, which causes the release of the B800 molecules and subsequently impacts the spectral properties of the carotenoids within the B850 LH2.

Silver damages the cytochrome c oxidase and the succinate dehydrogenase in the respiratory chain. Previous studies in 1974 and 2005 established that Ag⁺ ions inhibit the respiratory chain of E. coli (31, 32). In eukaryotes, Ag+ ions can induce mitochondrial dysfunction, partly by inhibiting respiration (33, 34). However, the complexes targeted by Ag⁺ ions were not yet identified. To check the effect of AgNO₃ on two respiratory complexes (succinate dehydrogenase and cbb₃ cytochrome c oxidase) from R. gelatinosus, exponentially growing wild-type cells under respiratory conditions were subjected to increasing concentration of AgNO₃ (25 to 150 μ M) and grown for another hour. To examine the effect of AgNO₃ on the cbb₃ oxidase, membrane proteins were solubilized and cytochrome c oxidase activity was assayed on blue native PAGE (BN-PAGE) (35). As shown in Fig. 6A, comparable diaminobenzidine (DAB)-positive bands corresponding to the cbb₃ oxidase were revealed in the solubilized membrane proteins from untreated and 25 μ M AgNO₃-stressed cells. Decreased activity was detected in the 50 μ M AgNO₃-stressed cells. However, no active cbb_3 oxidase was detected on membrane fractions isolated from 100 and 150 μ M AgNO₃-stressed cells (Fig. 6A). We should note that there was a slight effect on the amount of RC-LH with 100 and 150 μ M AqNO₃. This could be the consequence of membrane protein solubilization; indeed, the loss of B800 may destabilize LH2 in the presence of detergent. Nonetheless, the other blue-stained complexes on BN-PAGE did not seem to be affected (Fig. 6A and B), suggesting that AgNO₃ targets only some complexes, including

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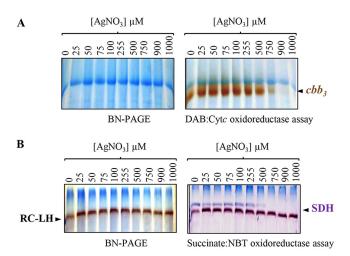


FIG 7 Effect of $AgNO_3$ treatment on respiratory complexes in membrane-enriched fractions. These fractions were mixed with increasing concentration of $AgNO_3$ for 1 h. DDM-solubilized membrane proteins were then separated on a 5% to 12% gradient BN-PAGE. (A) cbb_3 cytochrome c oxidase in-gel activity assay. (B) Succinate dehydrogenase (SDH) in-gel activity assay.

the cbb_3 cytochrome c oxidase and the RC-LH. Similarly, succinate:nitroblue tetrazolium (NBT) in gel assay revealed an active band, likely corresponding to succinate dehydrogenase (SDH). Treatment with increasing concentration of $AgNO_3$ resulted in a partial or full inhibition of this activity (Fig. 6B) suggesting that $AgNO_3$ also affected succinate dehydrogenase in exposed cells.

As for photosynthetic complexes, we analyzed the *in vitro* effect of AgNO $_3$ on respiratory complexes in isolated membrane protein fractions (Fig. 7). For that purpose, membranes from wild-type cells were incubated in buffer supplemented or not with increasing concentration of AgNO $_3$ (from 25 to 1,000 μ M). Similarly to the *in vivo* data, incubation of membrane proteins with increasing concentration of AgNO $_3$ led to a decrease in cytochrome c oxidase cbb_3 (Fig. 7A) and succinate dehydrogenase activities (Fig. 7B). Altogether, these data suggested that AgNO $_3$ could inhibit respiration by directly damaging the respiratory complexes, including cytochrome c oxidase and succinate dehydrogenase.

Silver damages respiratory complexes in *Escherichia coli* but not in *Bacillus subtilis*. The findings above prompted us to test the activity of respiratory complexes in other bacterial species after AgNO $_3$ treatment. To this aim, *E. coli* and *Bacillus subtilis* cells grown to exponential phase were subjected to increasing concentration of AgNO $_3$ (25 to 150 μ M) and grown for another hour. Membranes proteins were isolated, and activity assays for respiratory complexes were performed by BN-PAGE. As *E. coli* cells do not express any cytochrome c oxidase, we only assayed the activity of succinate dehydrogenase. We detected changes in the activity of this complex when cells were subjected to AgNO $_3$ stress, as the SDH activity decreased with increasing concentration of AgNO $_3$ (Fig. 8A). To ascertain that the detected band corresponds to the SDH, the succinate-NBT in-gel assay was also performed with membrane proteins isolated from the *sdhA* deletion mutant (36) (Fig. 8A). These results confirmed that AgNO $_3$ can affect respiration in *E. coli* and provide evidences that the SDH complex is a target of AgNO $_3$. In *B. subtilis*, however, AgNO $_3$ treatment did not affect the activity of the cytochrome c oxidase caa_3 , nor the activity of the SDH (Fig. 8B).

DISCUSSION

The use and spread of metal ions or nanoparticles represent a serious threat to the environment and to all living organisms because of the acute toxicity of these ions. Silver and copper ions have been used for their antimicrobial activities for several years. Nowadays, Ag⁺ nanoparticles are one of the most widely used nanoparticles in many industrial and health applications (37). The antimicrobial effect of Ag⁺ nanoparticles is



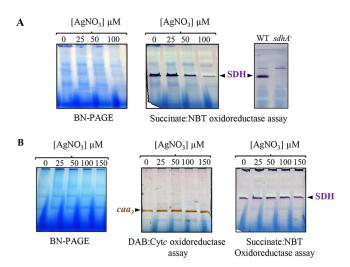
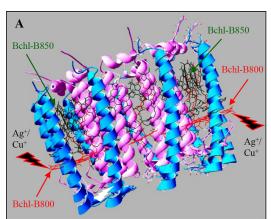


FIG 8 AgNO₃ effect on respiratory complexes in E. coli and B. subtilis cells. Strains were grown under aerobic respiratory condition and shocked for 1h with increasing concentration of AgNO₃. DDMsolubilized membrane proteins were separated on a 5% to 12% gradient BN-PAGE. (A) Succinate dehydrogenase (SDH) in-gel activity assay of E. coli membrane fractions. Membrane fraction from the SDH-deficient mutant ($sdhA^-$) was used as a control. (B) caa_3 cytochrome c oxidase and succinate dehydrogenase (SDH) in-gel activity assays of B. subtilis membranes.

in part related to the released Ag+ ions and their ability to interact with bacterial membranes (37–39). It is therefore important to characterize the toxicity of Ag⁺ ions and to identify cellular targets of this metal. Previous studies reported that the acute toxicity of Ag+ lies in its ability to interact with membranes but also in its ability to affect iron sulfur proteins (13). To identify targets of Ag+, we compared the impact of different metal ions on the stability and activity of membrane complexes in the purple photosynthetic bacterium R. gelatinosus. AgNO₃ was found to be more toxic than the other ions used, including CuSO₄ and CdCl₂. This may be related to the absence of an efficient efflux system to detoxify Ag+ and/or to its bioactivity and ability to damage molecules. Indeed, the Cop system involved in detoxification of Ag+ in other bacteria (9-11) is not induced by AgNO₃ in R. gelatinosus, which may increase the susceptibility of the bacterium to AgNO₃. In E. coli, although the Cop efflux is effective in expelling Ag+ ions outside the cells, AgNO₃ remains very toxic and targets different cellular components. In Chlamydomonas reinhardtii (39, 40) and Arabidopsis thaliana (41), Ag+ and Cu⁺ exposures were both found to significantly inhibit growth and to induce decreases in photosynthesis and chlorophyll content. Here, we found that both metals target the bacterial LH2. Both Ag⁺ and Cu²⁺ specifically target the B800 molecules but not the B850 ones (Fig. 9). Similar results were reported when LH2 complexes from R. sphaeroides and R. acidophila were subjected to high atmospheric pressure (30). This could be related to the structure of this complex and the position of the chlorophyll molecules in the complex. In fact, the B850 bacteriochlorophylls with the carotenoid molecules are buried between the concentric rings formed by the α and β subunit outer rings (25) and are therefore well shielded from the external buffer. In contrast, the B800 molecules are located between the outer rings formed by helices of the etasubunits and are parallel to the lipid surface near the cytosolic side (Fig. 9). This positions the B800 molecules in contact with the solvent, where they would be more exposed than the B850 molecules, in agreement with water molecules being found close to the B800 molecules in the R. acidophila LH2 crystal structure (25, 29). The effect on B800 did not modify the complex stability, since the B850 molecules were not affected. Nevertheless, the resulting complex would be inefficient for light energy capture and photon transfer to the photochemical reaction center. Indeed, the LH2 light energy is transferred from B800 to B850. Energy transfer then occurs between B850 and B875 molecules in the light-harvesting complex LH1 to the RC (42). Thus, we



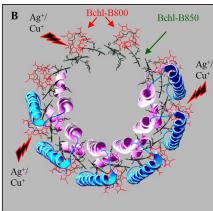


FIG 9 Ag⁺ and Cu⁺ specifically target the B800 within the LH2. Structure of the LH2 and arrangement of pigments within the complex from Rhodopseudomonas acidophila (1KZU.pdb) (25). The figure was generated using the Protein Data Bank (Swiss-PdbViewer). View parallel to the plane of the membrane (A) and from the top of the complex (B) showing the exposed B800 molecules (in red) located between the β subunit helices, which form the outer ring (cyan). The B850 molecules (green) are buried between the concentric rings formed by the helices of the α subunits (pink) and the outer β subunit helix ring. For better viewing, carotenoids and helices of α and β subunits were hidden.

assume that Ag⁺ or Cu⁺ exposure will result in reduced excitation transfer to the B870 in the RC and decreased photosynthesis yield. Only a slight effect of Ag+ and Cu+ was observed on the LH1-RC in the LH2-deficient strain. In the LH1-RC structures from R. palustris and T. tepidum (26, 27), the bacteriochlorophyll molecules in LH1 complexes are found sandwiched between the concentric ring formed by the α subunits and the external ring formed by the β helices, like the B850 molecules in the LH2 structure. These Bchl molecules and the RC-Bchls could therefore be shielded from the external buffer and therefore from damages that may be caused by the presence of metal ions. The mechanism by which Ag⁺ or Cu⁺ release B800 from the LH2 remains to be studied. Nonetheless, previous studies have shown that Mg²⁺ in chlorophylls could be substituted, both in vitro and in vivo, by heavy metal ions (20, 43, 44). It was shown that in vivo substitution of the Mg²⁺ atom of chlorophyll by heavy metals, including Cu⁺, Cd²⁺, or Pb²⁺, is a major damage mechanism in stressed plants. Indeed, substitution of Mg²⁺ affects the LHCII and the photosystem PSII, thereby causing a decrease in photosynthesis (20). Likewise, it was shown that Ni²⁺, Cu²⁺, and Zn²⁺ induced a destabilization of heme binding to b-type hemoproteins and led to the release of heme from myoglobin, ferricytochrome b_s , indoleamine-dioxygenase, hemopexin, and cytochrome P450 (45, 46). Formation of a bioconjugate of human hemoglobin with ${\rm Ag^+}$ ions was also reported (47, 48). Finally, both Ag+ and Cu+ can displace metal- or damage-exposed 4Fe-4S clusters in proteins (13). By theses means, such metal ions can inhibit the activity of hemoproteins and metaloproteins in the membrane and the cytosol. AgNO₃ impacts on the activity of cytochrome c oxidase and succinate dehydrogenase in R. gelatinosus, as well as on the succinate dehydrogenase in E. coli, were demonstrated in this study. This could arise from the disruption of the interaction between the cofactors (heme or 4Fe-4S) and the proteins. Nevertheless, we should note that the effects of Ag+ on complexes reported in this study were obtained with high concentration of AgNO₂. No effect was shown in the Gram-positive bacterium B. subtilis. This may be related to the difference in the cell wall structure between Gram-positive and Gram-negative bacteria. The much thicker peptidoglycan layers in Gram-positive bacteria are crucial in protecting the cell from environmental stress, including that of external metal ions (49). In agreement with this, Staphylococcus aureus is less sensitive to AgNO₃ than E. coli, as AgNO₃ treatment was shown to strongly affect the membrane integrity of E. coli but not that of S. aureus (50–52). In a recent study, AgNO₃ was shown to affect the activity of the cytochrome c oxidase in B. subtilis (53). However, AgNO₃ interfered with the biogenesis process of the oxidase by displacing Cu²⁺ from the Sco



assembly protein (53). In our study, we checked the impact of AgNO₃ on already-assembled complexes. Nevertheless, prolonged exposure to metal should be further studied to better characterize the response to Ag⁺ stress in *B. subtilis*. Beside the direct effect of Cu²⁺ and Ag⁺ on these membrane complexes, heme or 4Fe-4S cluster degradation is expected to release iron, which may magnify the toxicity of metals, as excess free iron gives rise to hydroxyl radicals and induces oxidative stress. Most toxicity studies of Ag nanoparticles in bacteria and eukaryotes shed light on oxidative stress response. For a full characterization of nanoparticle toxicity mechanisms, future experiments should address the issue of interaction between membrane complexes involved in cellular bioenergetics and Ag nanoparticles.

MATERIALS AND METHODS

Bacterial strains and growth. *E. coli* and *B. subtilis* cells were grown aerobically (500-ml flasks containing 50 ml medium) at 37° C in LB medium. *R. gelatinosus* cells were grown at 30° C, in the dark microaerobically (low oxygenation in 50-ml flasks containing 50 ml medium) or in light by photosynthesis (filled tubes with residual oxygen in the medium) in malate growth medium (54). The antibiotics kanamycin (Kan) and trimethoprim (Tmp) were used at a final concentration of 50 μ g/ml.

Growth inhibition curves were monitored at $OD_{680'}$ with measurements taken every 15 min for 24 h, using an Infinite M200 luminometer (Tecan, Mannerdorf, Switzerland) for aerobic condition. For photosynthesis conditions, strains were grown as described above, and OD was measured after 24 h using the Tecan luminometer.

Membrane protein preparation. Cells were disrupted by sonication in 0.1 M sodium phosphate buffer (pH 7.4) containing 1 mM phenylmethylsulfonyl fluoride. Unbroken cells were removed by a low-speed centrifugation step (25,000 \times g, 30 min, 4°C), and supernatants were subjected to ultracentrifugation (200,000 \times g, 90 min, 4°C) to collect the membrane fraction. Membrane fractions were then resuspended in the same buffer. Membrane protein concentration was estimated using the bicinchoninic acid assay (Sigma), with bovine serum albumin as the standard. For membrane protein metal treatment, required concentrations of metal solution were mixed with 50 mg/ml membrane proteins at room temperature. Spectra were recorded every 30 min.

Spectrophotometric measurements. Absorption spectroscopy was performed with a Cary 500 spectrophotometer. For spectra on whole cells, cells were resuspended in a 60% (wt/vol) sucrose solution. Membrane fractions were in 0.1 M sodium phosphate buffer (pH 7.4).

Blue native gel electrophoresis and in-gel assays. To assay *cbb*₃ and succinate dehydrogenase activities, *R. gelatinosus* wild-type cells were grown microaerobically. For *E. coli* and *B. subtilis*, cells were grown aerobically. Membranes were prepared as previously described. Blue native polyacrylamide gel electrophoresis (BN-PAGE) and in-gel Cox activity assays (DAB:Cytc staining) were performed as described in (35), and succinate dehydrogenase activity was assayed using succinate and NBT (nitroblue tetrazolium), as described in reference (55).

Western blot analysis and HisProbe-HRP detection. Equal amounts of cells (${\rm OD_{680}}=1$) were disrupted in SDS loading buffer, and proteins were then separated on a 15% SDS-PAGE and further transferred to a Hybond ECL polyvinylidene difluoride (PVDF) membrane (GE Healthcare). Membranes were then probed with the HisProbe-horseradish peroxidase (HRP) (Pierce), according to the manufacturer's instructions, and positive bands were detected using a chemiluminescent HRP substrate, according to the method of Haan and Behrmann (56). Image capture was performed with a ChemiDoc camera system (Bio-Rad).

SUPPLEMENTAL MATERIAL

Supplemental material for this article may be found at https://doi.org/10.1128/mBio.01535-18.

FIG S1, PDF file, 0.1 MB.

FIG S2, PDF file, 0.1 MB.

FIG S3, PDF file, 0.1 MB.

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We declare that we have no conflicts of interest with the contents of this article.

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