

Kinetics and Mechanism of Mineral Respiration: How Iron Hemes Synchronize Electron Transfer Rates

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Abstract: Anaerobic microorganisms of the *Geobacter* genus are effective electron sources for the synthesis of nanoparticles, for bioremediation of polluted water, and for the production of electricity in fuel cells. In multistep reactions, electrons are transferred via iron/heme cofactors of *c*-type cytochromes from the inner cell membrane to extracellular metal ions, which are bound to outer membrane cytochromes. We measured electron production and electron flux rates to $5 \times 10^5 \text{ e}^- \text{ s}^{-1}$ per *G. sulfurreducens*. Remarkably, these rates are independent of the oxidants, and follow zero order kinetics. It turned out that the microorganisms regulate electron flux rates by increasing their $\text{Fe}^{2+}/\text{Fe}^{3+}$ ratios in the multiheme cytochromes whenever the activity of the extracellular metal oxidants is diminished. By this mechanism the respiration remains constant even when oxidizing conditions are changing. This homeostasis is a vital condition for living systems, and makes *G. sulfurreducens* a versatile electron source.

Some anaerobic microorganisms are able to use extracellular metal salts as terminal oxidants in their respiratory pathways (mineral respiration).^[1] Important examples are bacteria of the *Geobacter* genus,^[2] which turned out to be effective electron sources for the synthesis of nanoparticles,^[3] for bioremediation of toxic and radioactive metals,^[4] and for the current production in microbial fuel cells.^[5] The respiratory pathways are based on electron transfer (ET) from the inside of the cells to the exterior. ET reactions can be mediated by proteins of bacterial filaments or by *c*-type cytochromes.^[1,6,7] We studied ET processes, where *c*-type

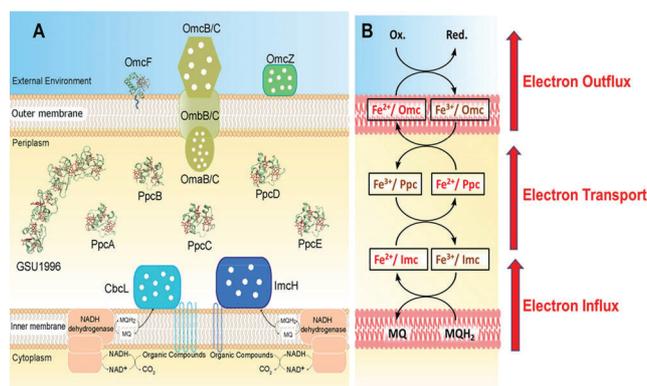


Figure 1. Outline of the inner and outer cell membrane of *Geobacter sulfurreducens*. A) Multiheme *c*-type cytochromes that are involved in anaerobic (mineral) respiration. Electrons originating from the oxidation of organic compounds are transferred to a menaquinone pool via an NADH hydrogenase located in the inner membrane. Depending on the redox potential of the final electron acceptor, the electrons are transferred to one of two pathways: the CbcL-dependent pathway, which operates when the final oxidants are at or below redox potentials of -100 mV vs. NHE, or the ImcH pathway, which is used when the final oxidants are above these redox potential values. B) During mineral respiration, several ET steps between the iron/hemes of the multiheme cytochromes transport electrons through the periplasm and the outer membrane to the extracellular oxidant.

cytochromes are the electron carriers (Figure 1). The respiratory chain starts at the inner cell membrane (electron influx) by enzymatic reduction of the $\text{Fe}^{3+}/\text{heme}$ cofactors of inner membrane cytochromes (Imc). During electron transport through the periplasm, several ET steps occur within and between the periplasmic multiheme cytochromes (Ppc). Finally, $\text{Fe}^{3+}/\text{hemes}$ of the outer membrane cytochromes (Omc) accept these electrons, and reduce the extracellular metal ions (electron outflux).

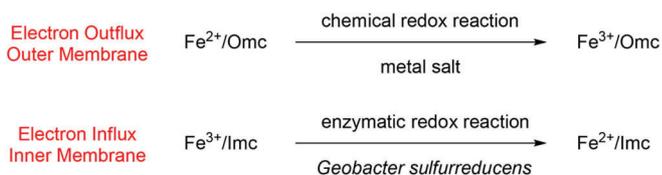
We have measured the electron production and flux through the cell by *in vivo* experiments with *Geobacter sulfurreducens* using AgNO_3 , Na_2CrO_4 , and AgCl as oxidants (Scheme 1). It turned out that each microorganism produces 5×10^5 electrons per second, and that the electron flux rate is nearly independent of oxidation potentials, aggregation states, charges and concentrations of the extracellular metal salts. This surprising behavior is accomplished by the bacteria, which are able to vary the $\text{Fe}^{2+}/\text{Fe}^{3+}$ ratios in the hemes of the cytochromes, allowing to synchronize the electron outflux rates with the constant electron influx rate.

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Scheme 1. Chemical oxidation of Fe^{2+} /hemes by extracellular metal ions (electron outflux), and enzymatic reduction of the Fe^{3+} /hemes at the inner membrane (electron influx). By measuring the changes in the concentrations of the Fe^{2+} /hemes and the oxidizing metal ions, the ET rates were determined.

First experiments were performed with AgNO_3 , which was reduced by *G. sulfurreducens* to Ag nanoparticles (AgNPs). Redox reactions between the oxidant and the Fe^{2+} /hemes rely on complexation of Ag^+ ions by the amino acids of Omc proteins. High complexation constants prevent the migration of Ag^+ ions into the cell, and thus protect the bacteria against the toxicity of Ag^+ ions. To verify Ag^+ complexation, NMR titration experiments were carried out with OmcF, the only Omc of *G. sulfurreducens* whose structure was determined in NMR studies.^[8] In the 11 kDa OmcF a methionine, Met86, and a histidine, His34, are well oriented for the coordination of a metal ion in a linear fashion (Figure 2 A). Both amino acids are known to bind Ag^+ ions in peptides.^[9] ^1H NMR titration of OmcF with silver nitrate in D_2O confirmed silver complexation by the imidazole group of His34 as well as by the sulfur atom of Met86 (Figure S1).

This titration furthermore indicated the formation of a 3:1 Ag^+/OmcF complex (Figure 2B), implying other coordinating residues whose proton signals could not be followed, likely because of the neighboring paramagnetic iron/heme entity. In order to determine the binding constant, we added imidazole- d_4 as competing ligand, whose affinity constant to Ag^+ ions is known.^[10] This created two sets of protein signals (Figures 2C and Figure S2), which were due to a partial exchange of the axial His39-heme ligand by the added imidazole- d_4 . This type of exchange had already been observed in other iron/heme systems,^[11] and we confirmed this ligand exchange for OmcF by variation of the imidazole- d_4 concentration (Figure S3). From the signals of Met86 and His34, two of the three Ag^+/OmcF binding constants were determined to be $\log K_{1,1} = 6.1(\pm 0.3)$ and $\log K_{2,1} = 6.2(\pm 0.5)$ (Figure S4). The third complexation could not be fitted with a reasonable error. The binding constants correspond to typical values reported for other histidine/silver/methionine complexes.^[9,12] Binding between Ag^+ and two (protein-free) histidine molecules has also a binding constant on the order of 10^6 .^[10] The NMR structures of the other outer membrane cytochromes of *G. sulfurreducens* such as OmcB and OmcC are not known. They are large, multiheme cytochromes with a high number of free histidine residues, which should bind Ag^+ ions also very efficiently.^[13]

Experiments to determine the ET rates were carried out with 0.6–0.9 μM *G. sulfurreducens* and 50 μM AgNO_3 solutions with low Cl^- and Fe^{2+} concentrations (Figure S5).^[14] UV/Vis spectra exhibited Soret and Q-bands of the iron hemes, as well as scattering contributions of the microorganisms (Figure 3A). As the AgNP concentration was low, the plasmon

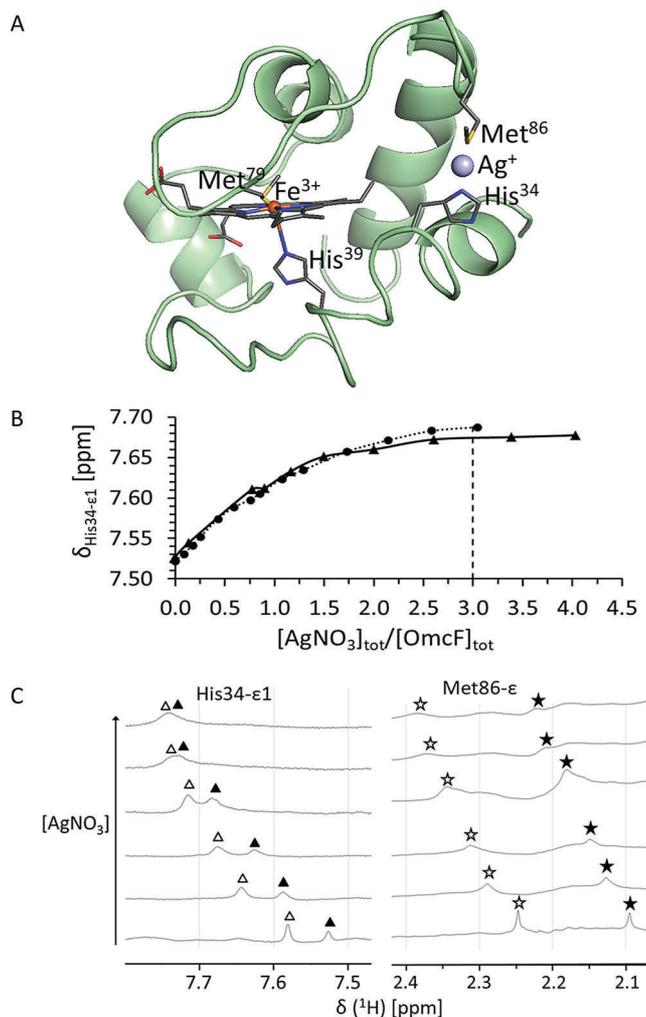


Figure 2. A) Complexation site of one Ag^+ ion on the outer membrane cytochrome OmcF, containing a Fe^{3+} /heme cofactor. The complexation sites of the two other Ag^+ ions bound by OmcF could not be determined. B) Change in the NMR chemical shift of one aromatic C–H group of His34 during the addition of AgNO_3 on OmcF (1.1 mM: circles, and 0.2 mM: triangles) in HEPES buffer (pD 7.8). A 3:1 Ag^+/OmcF stoichiometry yielded a good fit of the experimental data. C) NMR chemical shifts of one aromatic C–H group of His34 and of the CH_3 group of Met86 (OmcF: 1 mM) during addition of AgNO_3 (0–7 mM). The NMR experiments were carried out in competition with imidazole- d_4 , which duplicated the signals (blank and filled) due to displacement of His39.

resonance of the AgNPs was weak and overlapped with the iron/heme Soret bands (Figure S6). Therefore, the concentration changes of the Fe^{2+} /hemes were analyzed by their Q-bands (Figure S7). AgNP formation was determined by the increasing scattering of the microorganisms at 610 nm (Figure 3C,D)^[15] and EDS experiments (Figure S8).

Addition of AgNO_3 to *G. sulfurreducens* oxidized in less than two seconds more than 98% of the Fe^{2+} /hemes to Fe^{3+} /hemes (Figure S9), and the $\text{Fe}^{2+}/\text{Fe}^{3+}$ ratio remained nearly constant for the next 4–5 minutes (Figure 3B). During this time, the light scattering increased in a nearly linear way (Figure 3C), which indicated a constant AgNP formation that was independent of the Ag^+ concentration.^[15] Thus, its rate

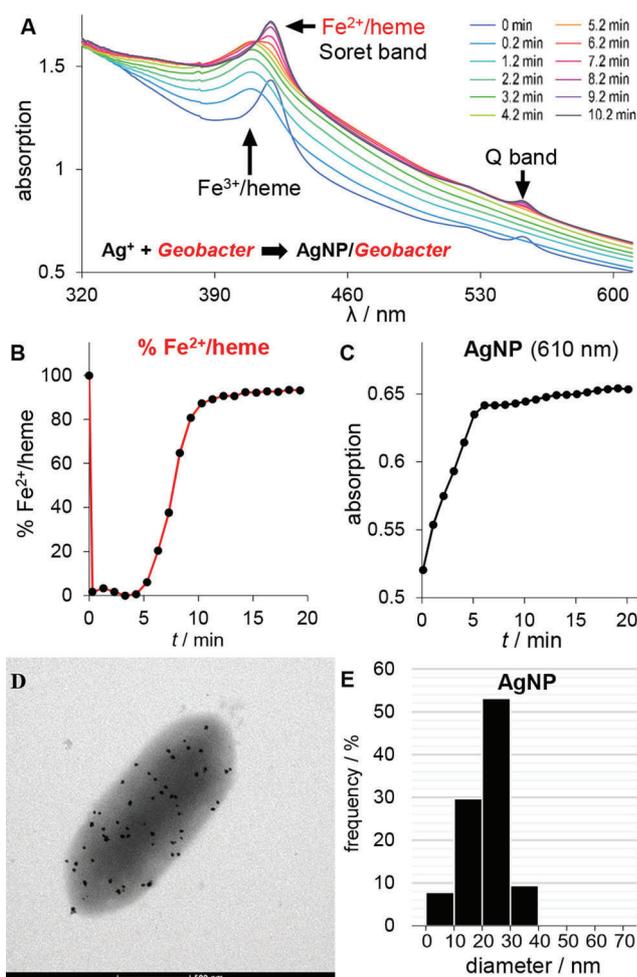
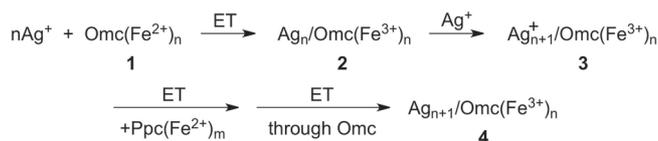


Figure 3. Reduction of Ag^+ ions (0.05 mM) by *G. sulfurreducens* (0.7 μM). A) UV/Vis spectra of the bacterial suspension during Ag^+ reduction (indicated times correspond to the measurement start at 610 nm; scan time = 1 min). The spectrum at 0 min was registered before AgNO_3 addition. B) Time dependence of the Fe^{2+} /heme concentration, determined from its Q-band (Figure S7). C) Time dependence of the scattering (610 nm) demonstrating AgNP formation. D) TEM image recorded at the end of the redox reaction showing AgNPs attached to the microorganism. E) Size distribution of AgNPs at the microorganism.

followed zero order kinetics. After consumption of the Ag^+ ions, Fe^{3+} /hemes were reduced back to Fe^{2+} /hemes (Figure 3B). The observed zero order kinetics during AgNP formation can be explained by a constant number of reaction sites at which AgNPs grew on the surface of the bacteria (Figure 3D), as well as an excess of Ag^+ ions (0.7 μM *G. sulfurreducens*; 0.05 mM AgNO_3) at the reaction start.^[16] As long as the Ag^+ ions were in large excess, their addition to the AgNPs $2 \rightarrow 3$ was obviously faster than ET between and through the cytochromes.^[17] Thus, the ET steps $3 \rightarrow 4$ and not the Ag^+ addition to 2 were rate determining (Scheme 2). We repeated the measurements more than 10 times with different generations of *G. sulfurreducens* and calculated an average ET rate of $3.0 \times 10^5 \text{ es}^{-1}$ per bacterium with a variation of $\pm 40\%$ at 30°C (Table 1). When the reaction temperature was



Scheme 2. Formation of AgNPs starts with complexation of Ag^+ ions to Omc 1 and subsequent oxidation of Fe^{2+} /hemes to Fe^{3+} /hemes in 2 . The AgNP growth occurs by Ag^+ addition ($2 \rightarrow 3$), as well as ET steps between and through the *c*-type cytochromes leading to 4 , which continues the AgNP growth.

Table 1: Electron flux rates with different oxidants and different generations of *G. sulfurreducens*.

Oxidant	Redox potential [V] ^[a] (pH 7)	ET rate [es^{-1}] ^[b]	Temperature [°C]
AgNO_3	0.80	3.0×10^5	30
		1.5×10^5	20
		0.8×10^5	15
Na_2CrO_4	0.62	5.0×10^5	30
AgCl	0.21	7.5×10^5	30

[a] Standard reduction potentials against NHE were used. They were not corrected for the influence of the reaction mixture and the reaction temperature. [b] The variation was $\pm 40\%$.

decreased to 20 and 15°C, the electron flux slowed down by a factor of 2.0 and 3.8, respectively.

Interestingly, the initial oxidation of the Fe^{2+} /hemes of *G. sulfurreducens* in the resting state by Ag^+ ions ($1 \rightarrow 2$) was much faster than the electron flux during respiration. It took less than 2 seconds to oxidize nearly all Fe^{2+} /hemes to Fe^{3+} /hemes (Figure S9). During this initial phase, about 10% of the AgNPs were generated. As Ag^+ ions were in 7×10^7 fold excess to *G. sulfurreducens*, the reduction of 10% of the Ag^+ ions was caused by 0.7×10^7 Fe^{2+} /hemes. This number of iron/hemes per bacterium is in good agreement with direct iron/heme measurements by Esteve-Núñez et al.^[18]

A characteristic of the silver ion system is that not the decrease of the oxidants, the Ag^+ ions, but the increase of the reaction products, the AgNPs, was determined. In order to analyze directly the change of the oxidants by UV/Vis spectroscopy, we used Na_2CrO_4 as a reagent (Figure 4A,C). Another difference in these two oxidants is that CrO_4^{2-} anions instead of Ag^+ cations were bound by the amino acids of the Omc proteins, and Cr^{3+} cations instead of Ag^0 nanoparticles were formed as reaction products. Several experiments with 0.7–0.9 μM *G. sulfurreducens* and 0.05 or 0.1 mM Na_2CrO_4 solutions at 30°C were carried out.^[19] Figure 4C shows that the concentration of the oxidant decreased in a constant way. Thus, also with Na_2CrO_4 , a concentration change did not influence the redox rate. The reaction was of zero order in chromate, so that its rate could be calculated directly from the decrease in the chromate concentration. An average ET rate of $5.0 \times 10^5 \text{ es}^{-1}$ per bacterium at 30°C was determined (Table 1). Interestingly, the Fe^{2+} /heme concentration increased slightly during the experiment (Figure 4B). This rise could compensate for a slowdown of the redox reaction caused by a decrease of the chromate concentration, or by a change of the redox

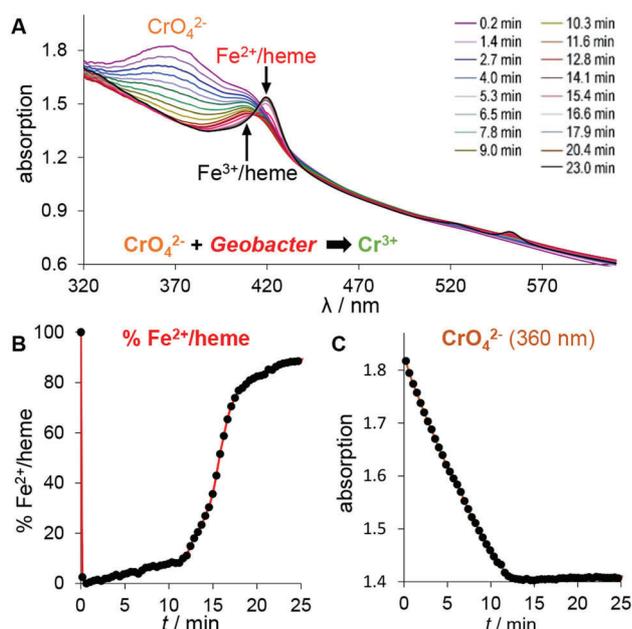


Figure 4. Reduction of CrO_4^{2-} ions (0.1 mM) by *G. sulfurreducens* (0.8 μM). A) UV/Vis monitoring of the bacterial suspension upon CrO_4^{2-} reduction (indicated times correspond to the measurement start at 610 nm; scan time = 25.3 s). B) Time dependence of the Fe^{2+} /heme concentration, determined from its Q-band (see also Figure S7). C) Time dependence of the chromate concentration, measured at 360 nm.

potentials induced by Cr^{3+} ions attached to amino acids of Omc.

A much more pronounced increase of the Fe^{2+} /heme concentration was observed when solid AgCl nanocrystals served as the oxidant.^[3a] AgCl nanocrystals were generated by addition of AgNO_3 to a Cl^- -containing growth medium (Figure S10),^[20] to which the same volume of a bacteria suspension was injected, leading to a 0.1 mM AgCl and 0.45 μM *G. sulfurreducens* solution. Within 5 minutes (Figure 5A,C) the microorganisms reduced the insoluble AgCl nanocrystals (Figure 5D) into insoluble AgNPs (Figure 5E).^[21] The AgNP formation rate remained constant, demonstrating that the product formation again followed a zero order kinetics. However, in sharp contrast to experiments with soluble Ag^+ ions (Figure 3B), the reduction of solid AgCl was accompanied by an increase of the Fe^{2+} /heme concentration from 15 to 95% (Figure 5B). This implies different reaction mechanisms. With soluble Ag^+ ions as the oxidant, complexation by Omc proteins led to a certain number of AgNPs at the microorganisms. These are the reaction sites to which the large excess of Ag^+ ions diffused and became reduced.

The situation with insoluble AgCl nanocrystals as oxidants was completely different as nearly all Ag^+ ions were stored in the solid AgCl nanocrystals.^[22] The reduction of each AgCl particle to an AgNP therefore decreased the concentration of the redox sites, thus the electron flux rate should decrease continuously. However, Figure 5C shows that the rate of the AgNP growth remained constant. This was actually accomplished by the concentration increase of the reaction

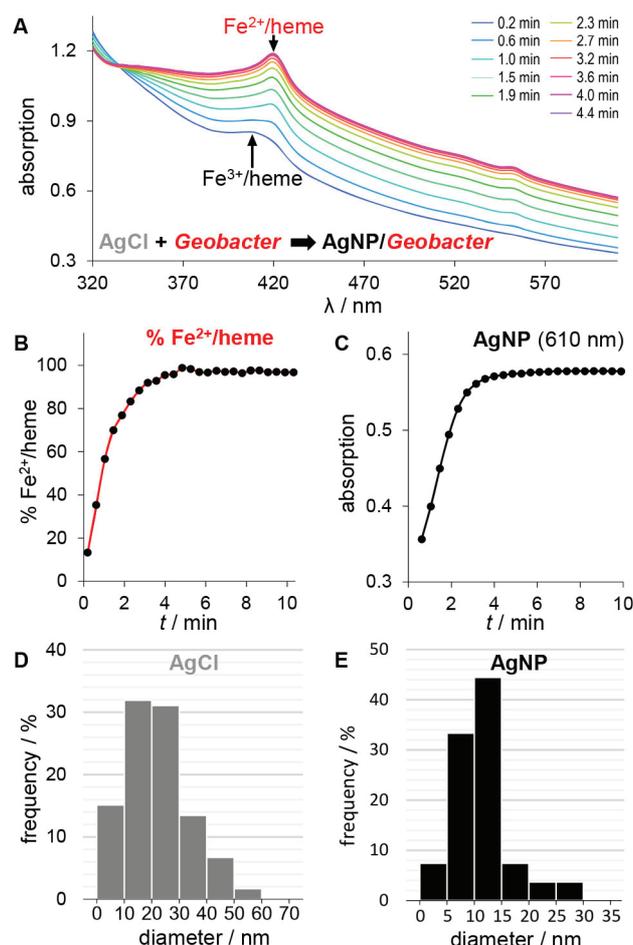


Figure 5. Mineral respiration of *G. sulfurreducens* (0.45 μM) using solid silver chloride nanocrystals (0.1 mM) as an extracellular oxidant. A) UV/Vis monitoring of the bacterial suspension upon AgCl reduction (indicated times correspond to the measurement start at 610 nm; scan time = 25.3 s). B) Increase of the Fe^{2+} /heme concentration during the reaction, determined from the Q-bands. C) Linear increase of the scattering at 610 nm demonstrating zero order kinetics for AgNP formation. D) Size distribution of the oxidizing AgCl nanocrystals. E) Size distribution of the generated AgNPs.

partners, the Fe^{2+} /hemes (Figure 5B). Obviously, the increasing Fe^{2+} /heme: Fe^{3+} /heme ratio in a multiheme Omc raised its reactivity to such an extent that the AgNP formation rates remained constant. We measured an average ET rate at 30 °C of $7.5 \times 10^5 \text{ e s}^{-1}$ per bacterium with a variation of $\pm 40\%$ (Table 1). The chemical rates in Table 1 show that respiration of *G. sulfurreducens* using different oxidants occurs with EET rates of $5 \times 10^3 (\pm 2 \times 10^5) \text{ e s}^{-1}$ per bacterium at 30 °C. This corresponds to a current production of $80 (\pm 30) \text{ fA}$, which is similar to electrochemical data. Jian et al. have measured a current of $90 (\pm 30) \text{ fA}$ in experiments where a single *G. sulfurreducens* cell was in contact with a gold electrode,^[23] and El-Naggar et al. determined currents of $60 (\pm 40) \text{ fA}$ in electrochemical measurements with *Shewanella oneidensis*.^[24]

Our experimental results show that in mineral respiration of *G. sulfurreducens*, the electron flux remains constant during the decrease of the oxidant concentration, demonstrating zero order reaction kinetics. In addition, the electron

flux rates are nearly independent of the type of the oxidant. This could be a general effect during mineral respiration of anaerobic bacteria; it might occur also with other members of the *Geobacter* genus or with *S. oneidensis* microorganisms. The independence from the environmental conditions demonstrates the flexibility of these bacteria, which adapt to changing external redox potentials by changing $\text{Fe}^{2+}/\text{heme}:\text{Fe}^{3+}/\text{heme}$ ratios in the *c*-type cytochromes. Decreasing the reactivity or the number of the reaction sites of the oxidizing metal ions is compensated by an increase of the $\text{Fe}^{2+}/\text{heme}$ population in the multiheme outer membrane cytochromes. A constant ET is needed to maintain constant ATP synthesis during respiration.^[25] Such ATP homeostasis is a vital condition for living systems.^[26] Obviously, the $\text{Fe}^{3+}/\text{heme}$ concentrations in the inner membrane cytochromes remain high enough to stabilize the enzymatic respiration reaction. This might be one of the reasons for the high concentration of iron/hemes in these microorganisms (10^7 per *G. sulfurreducens*). Thus, ATP homeostasis is the result of an interplay between chemical redox reactions at the outer cell membrane and enzymatic redox reactions at the inner cell membrane (Figure 1B and Scheme 1). The iron/hemes of *c*-type cytochromes, which are the carriers of the electrons, act as buffers that synchronize the ET rates of the electron influx and the electron outflux. This makes *G. sulfurreducens* a predictable reaction tool for redox reactions.

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Conflict of interest

The authors declare no conflict of interest.

Keywords: electron transfer · *Geobacter sulfurreducens* · iron heme · silver nanoparticles · silver proteins

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- [1] For a recent review see: L. Shi, H. Dong, G. Reguera, H. Beyenal, A. Liu, H.-Q. Yu, J. K. Fredrickson, *Nat. Rev. Microbiol.* **2016**, *14*, 651–662.
 [2] a) D. R. Lovley, J. F. Stolz, G. L. Nord, E. J. P. Phillips, *Nature* **1987**, *330*, 252–254; b) F. Caccavo, D. J. Lonergan, D. R. Lovley,

- M. Davis, J. F. Stolz, M. J. McNemy, *Appl. Environ. Microbiol.* **1994**, *60*, 3752–3759.
 [3] a) N. Law, S. Ansari, F. R. Livens, J. R. Renshaw, J. R. Lloyd, *Appl. Environ. Microbiol.* **2008**, *74*, 7090–7093; b) J. R. Lloyd, J. M. Byrne, V. S. Coker, *Curr. Opin. Biotechnol.* **2011**, *22*, 509–515.
 [4] “*Geobacter*: The Microbe Electrics Physiology, Ecology, and Practical Applications”: D. R. Lovley, T. Ueki, T. Zhang, N. S. Malvanker, P. M. Shrestha, K. A. Flanagan, M. Aklujkar, J. E. Butler, L. Giloteaux, A.-E. Rotaru, D. E. Holmes, A. E. Franks, R. Orellana, C. Risso, K. P. Nevin, in *Advances in Microbial Physiology* (Ed.: R. K. Poole), Academic Press, San Diego, **2011**, pp. 1–100.
 [5] For a recent review see: A. J. Slate, K. A. Whitehead, D. A. C. Brownson, C. E. Banks, *Renewable Sustainable Energy Rev.* **2019**, *101*, 60–81.
 [6] a) G. Reguera, K. D. McCarthy, T. Mehta, J. S. Nicoll, M. T. Tuominen, D. R. Lovley, *Nature* **2005**, *435*, 1098–1101; b) S. E. Childers, S. Ciuffo, *Nature* **2002**, *416*, 767–769.
 [7] The composition of bacterial filaments (pili) is in some cases still a matter of discussion: a) F. Wang, Y. Gu, J. P. O’Brien, S. M. Yi, S. E. Yalcin, V. Srikanth, S. Shen, D. Vu, N. L. Ing, A. L. Hochbaum, E. H. Engelmann, N. S. Malvanker, *Cell* **2019**, *177*, 361–369; b) D. L. Lovley, D. Walker, *Front. Microbiol.* **2019**, *10*, 2078.
 [8] a) J. M. Dantas, M. Silva e Sousa, C. A. Salgueiro, M. Bruix, *Biomol. NMR Assignments* **2015**, *9*, 365–368; b) J. M. Dantas, M. A. Silva, D. Pantoja-Uceda, D. L. Turner, M. Bruix, C. A. Salgueiro, *Biochim. Biophys. Acta Bioenerg.* **2017**, *1858*, 733–741.
 [9] V. Chabert, M. Hologne, O. Sénéque, O. Walker, K. M. Fromm, *Chem. Commun.* **2018**, *54*, 10419–10422.
 [10] R. Czoik, A. Heintz, E. John, W. Marczak, *Acta Phys. Pol. A* **2008**, *114*, A51–A56.
 [11] a) J. Hirst, S. K. Wilcox, J. Ai, P. Moënné-Loccoz, T. M. Loehr, D. B. Goodin, *Biochemistry* **2001**, *40*, 1274–1283; b) D. E. McRee, G. M. Jensen, M. M. Fitzgerald, H. A. Siegel, D. B. Goodin, *Proc. Natl. Acad. Sci. USA* **1994**, *91*, 12847–12855.
 [12] V. Chabert, M. Hologne, O. Sénéque, A. Crochet, O. Walker, K. M. Fromm, *Chem. Commun.* **2017**, *53*, 6105–6108.
 [13] a) UniProtKB-Q749K5(CYCB_GEOSL); b) UniProtKB-Q749L1 (CYTC_GEOSL).
 [14] In a study using different AgNO_3 amounts these concentrations turned out to be appropriate for detailed experiments. The preparation of the *G. sulfurreducens* solution followed the procedure described in S. I. Vasylevskiy, S. Kracht, P. Corcosa, K. M. Fromm, B. Giese, M. Fütég, *Angew. Chem. Int. Ed.* **2017**, *56*, 5926–5930; *Angew. Chem.* **2017**, *129*, 6020–6024. The growth medium is given in Figure S5. The experiments were carried out under strictly anaerobic conditions. The OD_{600} values for different *G. sulfurreducens* generations varied between 0.45 and 0.75. Dilution experiments showed that OD_{600} changed in a linear way. The kinetic experiments were carried out in 3.0 mL of thermostatted *G. sulfurreducens* suspensions to which 0.1 mL AgNO_3 solutions were added with a syringe, so that the concentration in the reaction mixture was 0.05 mM. The concentrations of the Cl^- and Fe^{2+} ions in the solutions were 2 and about 10 μM , respectively. The $\text{Fe}^{2+}/\text{heme}$ concentrations were determined by integration of the Q-band areas.
 [15] We detected that the intensity at 610 nm increased linearly in experiments with increasing Ag^+ concentration. We checked the Ag^+ concentrations in the mother liquor by ICP spectroscopy: when the intensity increase had stopped, Ag^+ ions in the mother liquor could no longer be detected, and the analysis during the redox reaction showed a decrease in the Ag^+ concentration. Thus, we concluded that this intensity change might be an acceptable way to measure the formation of AgNPs. We decided

to use this observable, because ICP experiments were not very precise, as the bacteria had to be removed by filtration or centrifugation. Both procedures took time during which the microbiological reaction continued. In addition, during filtration some Ag^+ ions might be removed by the bacteria in the filter pores. A further complication was that the redox process were fast, requiring only a few minutes (Figure 3), which caused large errors in the ICP experiments. Therefore, we used the intensity at 610 nm to measure the formation and growth of AgNPs. In future work we will further study this phenomenon.

- [16] Complexation of Ag^+ ions by Omc at the beginning of the experiment generated the AgNP growth sites. Analyses of several TEM pictures showed that the number of AgNPs number is always less than 10^2 per microorganism (see, for example, Figure 3D). The excess of the Ag^+ ions per microorganism at the beginning of the experiment was 0.7×10^8 .
- [17] From the rate coefficient of $0.5 \times 10^{10} \text{ L mol}^{-1} \text{ s}^{-1}$ for the addition of Ag^+ ions to neutral Ag^0 one can assume that also the reaction with AgNPs might be very fast, see: A. Henglein, *Chem. Rev.* **1989**, *89*, 1861–1873; A. Henglein, *Ber. Bunsen-Ges.* **1977**, *81*, 556–561. ET reactions through the multiheme cytochrome MtrF of *Shewanella oneidensis* had been calculated to be about 10^4 s^{-1} : M. Breuer, K. M. Rosso, J. Blumberger, *Proc. Natl. Acad. Sci. USA* **2014**, *111*, 611–616. ET between the iron/hemes of different cytochromes should be even slower.
- [18] The number of iron/hemes had been determined to be about 10^7 per *Geobacter sulfurreducens*: A. Esteve-Núñez, J. Sosnik, P. Visconti, D. R. Lovley, *Environ. Microbiol.* **2008**, *10*, 497–505.
- [19] The kinetic experiments were carried out as described for AgNO_3 reactions. Under strictly anaerobic conditions, 0.1 mL Na_2CrO_4 solutions were added to the bacterial suspension in such concentrations that the chromate concentrations were 0.05 or 0.1 mM. Variation of the Cl^- concentration had no effect on the EET rate.
- [20] Under strictly anaerobic conditions, 0.05 mL of a 0.3M KCl solution was combined with 1.35 mL of the growth medium. Injection of 0.1 mL of a 3 mM AgNO_3 solution generated AgCl nanocrystals, as powder X-ray measurements proved (Figure 5D and Figure S9). After a few seconds, the mixture became clear and was combined with 1.5 mL of a suspension of *G. sulfurreducens* in the resting state.
- [21] Using the dimensions of the AgCl unit structure, the number n of Ag^+ ions in roundish AgCl nanocrystals with a diameter d is $n = 12 \cdot d^3$. The number of Ag^0 in AgNPs is $n = 30 \cdot d^3$, see: A. Henglein, M. Giersig, *J. Phys. Chem. B* **1999**, *103*, 9533–9539.
- [22] A medium-sized (20 nm) AgCl nanocrystal contains about 10^5 Ag^+ ions.
- [23] X. Jiang, J. Hu, E. R. Petersen, L.-A. Fitzgerald, C. S. Jackan, A. M. Lieber, B. R. Ringeisen, C. M. Lieber, J. C. Biffinger, *Nat. Commun.* **2013**, *4*, 2751.
- [24] B. Gross, M. Y. El-Naggar, *Rev. Sci. Instrum.* **2015**, *86*, 064301.
- [25] M. Saraste, *Science* **1999**, *283*, 1488–1493.
- [26] H. Y. Kueh, P. Niethammer, T. J. Mitchison, *Biophys. J.* **2013**, *104*, 1338–1348.

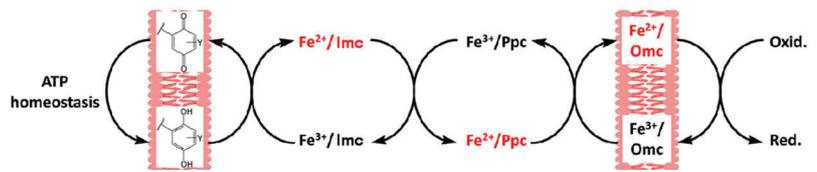
Communications



Electron Transfer

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Kinetics and Mechanism of Mineral
Respiration: How Iron Hemes
Synchronize Electron Transfer Rates



The bacterium *Geobacter sulfurreducens* requires extracellular metal salts for respiration. Each microorganism produces 5×10^5 electrons per second, which reduce the metal ions bound to the outer

membrane cytochromes. With iron heme cofactors of *c*-type cytochromes, the electron flow is independent of the concentration and type of oxidant. This results in ATP homeostasis.