Activation of Methanogenesis by Cadmium in the Marine Archaeon *Methanosarcina acetivorans*

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**Abstract**

*Methanosarcina acetivorans* was cultured in the presence of CdCl₂ to determine the metal effect on cell growth and biogas production. With methanol as substrate, cell growth and methane synthesis were not altered by cadmium, whereas with acetate, cadmium slightly increased both, growth and methane rate synthesis. In cultures metabolically active, incubations for short-term (minutes) with 10 μM total cadmium increased the methanogenesis rate by 6 and 9 folds in methanol- and acetate-grown cells, respectively. Cobalt and zinc but not copper or iron also activated the methane production rate. Methanogenic carbonic anhydrase and acetate kinase were directly activated by cadmium. Indeed, cells cultured in 100 μM total cadmium removed 41–69% of the heavy metal from the culture and accumulated 231–539 nmol Cd/mg cell protein. This is the first report showing that (i) Cd²⁺ has an activating effect on methanogenesis, a biotechnological relevant process in the bio-fuels field; and (ii) a methanogenic archaea is able to remove a heavy metal from aquatic environments.

**Introduction**

Methanogenesis is the pathway by which ion (H⁺, Na⁺) gradients across the plasma membrane are generated to drive ATP synthesis, with the concomitant production of methane as an end product. Methanogens are strict anaerobes belonging to the *Archaea* domain, which can be found in a broad variety of environments such as anaerobic digesters of sewage treatment plants, landfills, rice paddies, lakes and in the sea sediments, among others [1]. Indeed, these organisms have an essential role in the global carbon cycle by transforming small carbon molecules such as methanol, methylamines, CO₂ in the presence of H₂, formate, CO and acetate into methane. Because heavy metal pollution may develop in some of these habitats, methanogens may be exposed to this environmental stress with the consequent perturbation of the local ecology. Heavy metal pollution of water resources is now a widespread environmental and public health problem, as a result of their elevated toxicity, which may be exacerbated by their potential bio-magnification effect and accumulation throughout the ecological food webs.

Pollution of coastal zones by heavy metals such as Cd, Pb, Hg and Ni, is a major environmental problem in some regions of the world [2]. Once in the marine environment, these contaminants accumulate in sediments [3]. Cadmium ocean pollution and mobilization has increased exponentially up to 300 thousands per decade, where 40% of the total current pollution derives from anthropogenic activities [4]. In some coastal zones in the Gulf of Mexico, up to 2,530 μg L⁻¹ (22.6 μM) of cadmium has been found, a value far higher than permissible [5]. In other seas and oceans around the world, cadmium concentrations up to 20.5 μg L⁻¹ and 18, 400 μg g⁻¹ in waters and sediments, respectively, have been determined [4]. These cadmium pollution values highlight the importance of determining the toxicity of cadmium in organisms found in sediments, methanogens among them. Depending on the physicochemical environmental characteristics and microbial metabolism, heavy metal contaminants may be released from sediments back into the water. Under such situations, marine sediments may become a secondary source of pollution.

Most studies of microbial communities in the ocean have focused on bacterial diversity in marine sediments and the long-term impact brought about by heavy metals exposure [6]. There are few works about the toxic effect of heavy metals on methanogens in sludge [7] and laboratory strains [8,9]. Remarkably, it has been described that low concentrations of heavy metals are not toxic for methanogens in the sludge, but on the contrary they induce increased methane production [10]. This finding has not been further explored and hence the living components of the consortia affected by the metal have not been identified, and the biochemical mechanisms involved have not been elucidated.

In the present work, the marine archaeon *Methanosarcina acetivorans* was used as a model to assess the effect of cadmium on methanogenesis.
Materials and Methods

Chemicals
Acetate kinase from *M. thermophila*, deoxyribonuclease I from bovine pancreas (DNase I), acetyl-CoA, coenzyme A, acetyl phosphate, ATP and NADH were purchased from Sigma Chem. (St. Louis, Mo, USA). Phosphonoarylpyruvate, pyruvate kinase and NADH-lactate dehydrogenase (both enzymes from rabbit muscle) were from ROCHE (Germany). Absolute methanol, acetate, and CdCl₂ were of analytical grade.

Growth conditions
*Methanosarcina acetivorans* C2A strain, kindly provided by Prof. James G. Ferry (Pennsylvania State University, USA), was cultured under anoxic conditions in the regular high salt medium (HS-medium) described elsewhere [11]. Briefly, Milli Q water was placed into an anaerobic chamber (COY laboratory products, Grass Lake, Michigan, USA) filled up with 80% N₂, 15% CO₂ and 5% H₂. Then, the following salts were added, in g/L: NaCl, 25.4; NaHCO₃, 3.8; KC₁, 1.0; MgCl₂, 11; CaCl₂, 0.2; NH₄Cl, 1.0 and KH₂PO₄, 0.27 and resazurin, 0.001% (w/v) as redox indicator. Vitamin and trace mineral solutions were both added at 1% (v/v) as reported by Sowers et al [12]. Medium was bubbled with the mix of gases describe above for 2–3 h. Next, 120 mM methanol or 100 mM acetate were added as carbon source, followed by 1 g cysteine-HCl (8.2 mM) and 0.25 g Na₂SO₃H₂O (1.04 mM) to ensure complete chemical reduction of growth media. Final pH was 6.8–7.0. 50 mL of medium were poured into 100 mL serum-like bottles (Virmel, Mexico), sealed with a butyl rubber stopper (Virmel, Mexico) and secured with an aluminum crimp collar, 23.7 mM methanol or 100 mM acetate were added as carbon source, followed by 1 g cysteine-HCl (8.2 mM) and 0.25 g Na₂SO₃H₂O (1.04 mM) to ensure complete chemical reduction of growth media. Final pH was 6.8–7.0. 50 mL of medium were poured into 100 mL serum-like bottles (Virmel, Mexico), sealed with a butyl rubber stopper (Virmel, Mexico) and secured with an aluminum crimp collar (Virmel, Mexico). Medium was autoclaved at 121°C for 30 min. After autoclaving a precipitate was formed in the culture media but disappeared after 24 h, approximately. Cultures were started by adding fresh cell inocula and further incubating at 37°C without shaking. Growth was determined by measuring changes in absorbance at 600 nm.

Metabolites content determination
The concentration of the reduced cysteine and sulfide in the fresh medium was determined post column with DTNB (5, 5’-dithiobis-(2-nitrobenzoic acid) by HPLC as described elsewhere [13]. Briefly, 1 mL of fresh medium was taken with a syringe from the anaerobic culture bottles and immediately filtered through a 0.45 µm (pore diameter) filter unit (Millex-HV, Millipore, Ireland) and injected (50 µL) into the HPLC apparatus. The concentration of thiols-groups was calculated by using the DTNB molar extinction coefficient of 13.6 mM⁻¹ cm⁻¹. Sulfide was also determined spectrophotometrically by the methylene blue formation as described by King and Morris [14] with some modifications: in 10 mL anaerobic bottles sealed with a butyl rubber stopper and secured with an aluminum crimp collar, 23.7 mM zinc acetate, 60 mM NaOH, 0.18 mM N,N-dimethyl-β-phenylenediamine (DMPD) dissolved in 5 N HCl and 0.1 mL of culture medium, or different amounts of sulfide, were added by using a syringe and mixed until homogeneity. Then, 2.8 mM FeCl₃ was added and incubated at room temperature for 30 min for color development (methylene blue formation). Final volume was 2.5 mL. Samples were measured at 670 nm under anoxic conditions in an anaerobic chamber. The sulfide content-absorbance relationship was linear up to 350 nmol.

Methane production and methanol were determined by gas chromatography (Shimadzu GC2010 apparatus), equipped with a capillary column HP-PLOT/U of 30 m length, 0.32 mm I.D. and 10 µm film (Agilent, USA) and flame ionization detector. Methane standard curve was carried out as reported by Sowers [12]. For determination of extracellular acetate, aliquots withdrawn from cell cultures with acetate were centrifuged. Samples of the cell-free supernatant were diluted with a buffer containing 50 mM HEPES (4-(2-hydroxyethyl)-1-piperazinethanesulfonic acid), 10 mM MgCl₂ and 1 mM EGTA (ethylene glycol tetraacetic acid) at pH 7.5. The mixture reaction assay also contained 5 mM ATP, 2 mM phosphoenolpyruvate, 0.25 mM NADH, and acetate kinase, pyruvate kinase and lactate dehydrogenase, which were added in excess (>5 U, each) to ensure the complete phosphorylation of acetate coupled to NADH oxidation.

2.4 Cadmium exposure
To determine the effect of cadmium on growth, acetate and methanol cultures were carried out in the presence of different total CdCl₂ concentrations (0, 1, 10, 25, 50 and 100 µM) and the optical density was determined at 600 nm. Such range of cadmium concentrations has been reported to be toxic for a broad range of microorganisms from fresh [15] and marine waters [16].

To determine the effect of cadmium on methane biosynthesis, metabolically active cell cultures in the early stationary growth phase with acetate (at the 10–12 day of culture, with 8 mM of remaining acetate) or methanol (at the 3–4 day of culture, with 5 mM remaining methanol), respectively, were initially subjected to depletion of methane formed by gassing the culture bottles with sterile N₂. Thereafter, 1, 10 or 100 µM total CdCl₂ or other heavy metals (Zn²⁺, Cu²⁺, Hg²⁺, Fe²⁺ and Co²⁺) were added to the cultures and methane production was determined from the head space at different short times up to 12 min. For longer periods of methane synthesis (up to 60 min) in the presence of Cd²⁺, 20 mM acetate was further supplemented to the incubation medium.

2.5 Enzyme activity assays
Cell cultures of 750 mL grown on acetate were harvested under anoxic conditions in the early stationary phase by centrifuging at 3,000 x g for 10 min and washed once with 4 volumes of a solution containing 50 mM Tris-HCl pH 7.5, 20 mM MgCl₂ and 0.02 mM ZnCl₂. Then, the cell pellet was re-suspended in lysis buffer (0.1 M Na-phosphate, pH 8.0 plus some grains of DNase I), stirred strongly for 5 min and centrifuged at 70,000 x g for 30 min. The supernatant (cytosolic fraction; yield 50–100 mg protein) was kept on ice and used immediately for enzyme activities determination. All activities (except for carbonic anhydrase) were determined in the direction of acetate degradation in 50 mM Na₂-Hepes and 10 mM MgCl₂ buffer at pH 7.0 and 27±2°C, in the presence of different CdCl₂ concentrations. In all cases, the reaction assay was started by adding the enzyme (i.e. the cytosol-enriched fraction).

Acetate kinase (AK) activity was determined in cytosolic enriched-fractions of 50–75 µg protein in a reaction medium that also contained 5 mM ATP, 20 mM acetate, 0.2 mM NADH, 2 mM phosphoenol pyruvate and 10 U of both, pyruvate kinase and lactate dehydrogenase. One unit of enzyme (U) is the amount of active enzyme required to transform/produce 1 µmol of substrate/product in 1 min. Phosphotransacetylase (Pa) activity was determined as follows: 3–5 µg of cytosolic protein were incubated in the Hepes-Mg buffer with 5 mM acetyl-phosphate and 160 mM CoA; aliquots were withdrawn at different times (from 5 up to 60 s), mixed with 0.1 M phosphate buffer and 1 mM DTNB and the reaction monitored at 412 nm (representative traces are shown in figure S1).

CODH/acetylCoA synthase activity (CODH/AcCoA synthase activity) was determined anaerobically by mixing 10–25 µg protein with 80 µM
acetyl-CoA and measuring the release of CoA with DTNB at 412 nm. As several different enzymes may release CoA from acetyl-CoA, this activity was also specifically determined by measuring the CO-dependent reduction of methyl viologen as reported elsewhere [12]. Briefly, in an anaerobic sealed bottle the Hepes-Mg buffer +0.5 mM methyl viologen was saturated with CO by bubbling the gas for 30 min (reaction mixture); then, 1.2 mL of reaction mixture was poured into a sealed glass cuvette previously purged with CO. The reaction was started by adding 50 μg of protein and followed at 603 nm. As control of the CODH/AcCoA activity, the cytosolic fraction was gently mixed with air for 10 min, with the remaining activity being lower than 53% (n = 2) of that determined with saturating CO or acetyl-CoA (representative trace is shown in figure S2). Also, 0.5 mM sodium cyanide inhibited the reduction of methyl viologen coupled to CO oxidation by 85±8% (n = 3) as reported previously for the enzyme from M. thermophila [17].

Carbonic anhydrase (CA) activity was determined by incubating 2.5–5 μg of cytosolic protein with 100 mM Na-bicarbonate in a sealed 10 mL bottle with rubber stopper. To detect the CO formation, 5 μL of the head space was taken and injected at different times (0, 30, 60 and 120 s) in a gas chromatograph equipped with a Thermal Conductivity Detector. CO formed in assay reaction buffer without enzyme and with boiled enzyme was subtracted from the CO formed with enzyme (representative traces are shown in figures S3 and S4).

2.6 Cadmium removal and accumulation

Cells were harvested and washed as indicated above with 200 volumes of a solution containing 50 mM Tris-HCl, 2 mM MgCl₂ and 2 mM EGTA (TME buffer) at pH 7.5; the pellet was re-suspended in fresh buffer to give 5–10 mg protein/mL and frozen at −70°C until use. Aliquots of the cell suspension were digested with H₂SO₄+HNO₃ (1:3) for 2 h at 100°C and the intracellular cadmium content determined by atomic absorption spectrophotometry (Varian Spectra AA 640).

2.7 Ultrastructure analysis

Methanol-grown cells with or without 100 μM CdCl₂ were fixed by immersion in glutaraldehyde (3%, v/v, in phosphate buffer, pH 7.4), after removal from the culture medium, and dehydrated in graded ethanol. Samples of 1 mm² containing the cells were cut out in cross section with a diamond knife and embedded in 1:1 epoxy resin. To determine cadmium and sulfur localization inside the cells, atomic-resolved high angle annular dark-field scanning-transmission electron microscopy (HAADF-STEM) was used as reported previously [18].

The protein content was determined after cells were washed once with TME buffer by the Biuret method with bovine serum albumin as standard as described previously [13]. For the statistical analysis of the data, the Student’s t-test or a two way ANOVA and Bonferroni post analyses were performed using the Graph Pad PRISM version 5.01 software.

Results and Discussion

Cadmium solubility and effect on cell growth

Because cysteine and sulfide present in the culture medium bind the cadmium added with high affinity, the soluble free Cd²⁺ concentrations were estimated (see Table I) by using the program Chelator [19] and the following physico-chemical conditions. The concentration of the reduced cysteine and sulfide in the medium determined experimentally were for cysteine 1.7±0.03 mM and for sulfide 1.21±0.4 and 0.95±0.03 mM as determined by HPLC.

| Total Cd(Cl₂) (μM) | Methane produced mmol/240 h (acetate) | Methane produced mmol/240 h (methanol) | Cd removed and accumulated nmol/total 240 h (acetate) | Cd removed and accumulated nmol/total 240 h (methanol) |
|-------------------|--------------------------------------|--------------------------------------|--------------------------------------|--------------------------------------|
| 0                 | 0                                    | 0                                    | 0                                    | 0                                    |
| 1                 | 21                                   | 58                                   | 0.2                                  | 0.1                                  |
| 10                | 54                                   | 56                                   | 0.1                                  | 0.01                                 |
| 25                | 109                                  | 59                                   | 0.1                                  | 0.02                                 |
| 50                | 226                                  | 64                                   | 0.1                                  | 0.02                                 |
| 100               | 206                                  | 45                                   | 0.1                                  | 0.02                                 |

Data shown were obtained from cell cultures at the end of the growth curve. Values are the mean ± SD of at least 4 cultures from different batches.

Table 1. Methane production and cadmium accumulation in M. acetivorans cultured on acetate or methanol.

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(DTNB) and spectrophotometrically (methylene blue), respectively (mean ± SE, n = 4); the ionic strength = 0.77, pH = 7.0 and temperature = 36°C. The log values of the equilibrium constants ($K_{eq}$) for the association of the complexes were 13.4 and 20.13 for Cys-cadmium and Cys-Cd-Cys, and 6.1 for sulfide-cadmium [20].

Cells cultured on methanol showed similar growth either in the absence or presence of up to 100 μM total CdCl$_2$ (Fig. 1A and inset). With acetate, growth was slightly faster in cultures with 100 μM cadmium during the exponential phase (Fig. 1B and inset). To undoubtedly establish that the turbidity increase induced by cadmium was indeed reporting cell growth in acetate cultures, the growth rate (GR) and the duplication time (DT) were determined by using the curve of methane production vs time and assuming that methane production is proportional to the number of living cells present in the culture. The duplication times were similar to those reported previously for *M. acetivorans* [11]. No significant differences in GR values (0.064±0.003 versus 0.0625±0.003 h$^{-1}$) and DT values (10±2.3 h versus 11±2.7 h) were found for cells cultured in methanol without or with 100 μM total Cd$^{2+}$. In contrast, in acetate cultures the GR value was significantly higher in cultures with 100 μM Cd$^{2+}$ (0.028±0.004 versus 0.030±0.006 h$^{-1}$; n = 5, P<0.05). DT did not significantly changed (26±3 versus 24±2 h; n = 5) for cells cultured without or with 100 μM Cd, respectively (Fig. 1). Furthermore, two way ANOVA analyses on the global data showed that cadmium exerted a positive effect on the growth curve with acetate but not with methanol as carbon source (Figs. 1C and 1D).

The rate of methane production in acetate cultures with 100 μM total CdCl$_2$ was slightly but significantly higher than in its absence in the time-period from 110 up to 230 h of culture (Fig. 1D). It is worth noting that the methane yield, at the end of the growth curves (96 and 244 h for methanol and acetate, respectively), was the same under all conditions, because the total amount of carbon source added was identical (Table 1): 4.1±0.13 (control) and 4.1±0.02 mmol methane (+100 μM CdCl$_2$) for methanol and 4.5±0.3 (control) and 4.7±0.3 mmol methane (+100 μM CdCl$_2$) for acetate. On the other hand, when 500 μM total CdCl$_2$ was added to cultures with acetate or methanol as carbon source, no growth or methane synthesis were detected (data not shown) indicating that these high cadmium levels were indeed extremely toxic to the cells.

**Effect of cadmium on methane synthesis**

For short-term experiments, cultures of cells in the early stationary growth phase (10 days for acetate- and 4 days for methanol-grown cells) were incubated with CdCl$_2$ at 25–27°C. The concentrations of acetate and methanol remaining in the cultures were 8±3 mM (400±150 μmol acetate; n = 5) and 5±1 mM (250±50 μmol methanol; n = 5), respectively. Under these conditions, cadmium exerted a remarkably stimulating effect on the synthesis of methane in control cells not previously exposed to Cd$^{2+}$; the most potent activation was reached at 10 μM total CdCl$_2$ (Fig. 2A). Moreover, the rate of the methane production increased 9 and 6.5 fold for acetate- and methanol-grown cells, respectively, in 2 min (Fig. 2B).

After 10 min of incubation the methane produced, in the absence or presence of 10 μM total CdCl$_2$, by stationary acetate-grown cells was 18±3 and 26±4 μmol methane, and after 60 min

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**Figure 1.** Growth curves and methane synthesis of *M. acetivorans* cultured on methanol (A, C) or acetate (B, D), respectively, and in the absence (filled squares) or presence of 100 μM CdCl$_2$ (open squares). Values represent the mean ± SD of at least 4 different cell batches. *a:* P<0.05 vs control curve without cadmium using two way ANOVA. Inset; curves with 1 (filled circles), 10 (filled triangles), 25 (open squares) and 50 (open circles) μM CdCl$_2$.

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These experiments were started after thoroughly purging the flasks and adding the indicated CdCl₂ concentrations (time-point equal to zero). (C) Activation of methane synthesis by other heavy metals. Acetate-grown cells cultures were incubated for 5 min in the absence or presence of 100 μM of the metals indicated. At t = 0 (before metal addition), the methane remaining in the bottle cultures was 8.8 ± 1.2 μmol methane per culture. P < 0.05 using the Student's t-test for non-paired samples for vs control (without cadmium or other metal ion); vs cells exposed to 1 μM cadmium; vs methanol cultures exposed to cadmium.

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was 24 ± 2 and 43 ± 8 μmol methane, respectively (Fig. S5). Hence, the methane produced was the same regardless the carbon source concentration, sub-saturating or growth-limiting (8 mM acetate, Fig. 2B) for the 10 min experiments and saturating (20 mM acetate, Fig. S5) for the 60 min experiments.

Activation of methanogenesis was not exclusive for cadmium, since also 100 μM of the essential trace metals Co²⁺ or Zn²⁺ had a similar effect, whereas Cu²⁺ and Fe²⁺, also essential trace metals, or Hg²⁺ were poor activators of the methane production (Fig. 2C).

These data suggested that the activation of methane production by cadmium was not due to the precipitation of sulfur that may be toxic for the cell, as copper, iron and mercury can also form complexes with sulfur; in fact, insoluble complexes were apparent with iron. A copper inhibitory effect was previously described for methanogenesis derived from anaerobic sludge digestion [10,21]. On this regard, a positive effect of P₃M concentrations cadmium on growth presumably due to activation of CA was reported for Thalassiosira weissflogii, a diatom that also comes from the marine habitat [22]. Interestingly, acetate-grown M. acetivorans cells have significant higher AK, Pta, CODH/AcCoAs and CA protein content than methanol-grown cells [23,24]. Hence, the influence of cadmium on enzymes activities from the upper part of the aceticlastic pathway, which have not been previously determined in M. acetivorans, was here examined (Table 2).

AK activity was 10 fold lower (see legend to Table 2 for values) than that reported for the enzyme from M. thermophila [25]; the AK activity slightly increased (25–30%) by 10 μM total cadmium. This cadmium activating effect is intriguing because no metal has been reported to be required for AK activity in M. thermophila [26]. Pta activity under our conditions was 15 times lower than that reported for the enzyme from M. thermophila [17]. The last two enzymes were not activated by 0.01-10 μM total CdCl₂, but they were rather partially inhibited (Table 2). With a novel strategy to determine CA activity which was based on measuring by gas chromatography the CO₂ produced, the M. acetivorans CA showed a higher activity than that reported by semiquantitative electrometric method at alkaline pH for the M. thermophila enzyme [28] and marked activation by 1–10 μM total cadmium (Table 2).

Methanococcus CA is promiscuous respect to the metal bound into its active centre, because the presence of zinc, cobalt and even iron has been reported for this enzyme in M. thermophila and M. acetivorans [29,30]. Indeed, the recombinant purified CA showed activity even with Cd²⁺ [31]; hence, cadmium might also be able to bind and activate CA in vivo. Thus, activation of CA and AK by cadmium may be involved in the higher methane production in acetate-grown cells. Another possible explanation for the stimulation of the methane production was that cadmium uncoupled the methanogenic pathway by collapsing the ion gradient across the plasma membrane. However, the total protein determined at the end of culture in cells grown with cadmium suggested that ATP content was not compromised. On the other hand, cadmium activation of methanogenesis suggested metal internalized into cells; hence, the cadmium removal from cultures by cells was determined.

**Cadmium removal**

Under our culture conditions, in which the cysteine and sulfide concentrations were high, the added micromolar CdCl₂ concentrations yielded free Cd²⁺ concentrations in the pM range (Table 1). It is known that organic and inorganic sulfur may attenuate the toxicity of Cr (VI) in yeasts isolated from industrial wastes [32]. Hence, the low toxicity of cadmium in M. acetivorans may be due to the low free Cd²⁺ available in the medium. Nevertheless, cells surprisingly removed up to 70% and 40% of total added cadmium from the medium in the cultures with acetate or methanol, respectively (Table 1). In this regard, with 100 μM added CdCl₂, an accumulation of 0.54 and 0.23 μmol cadmium/mg cell protein (Table 1) was determined for acetate and methanol-grown cells, respectively, which were harvested after 10 or 4 days culture and washed once with an EGTA (e.g., potent metal ion chelating agent)-containing buffer. The cell-free culture medium contained 1.4 ± 0.1 μM total cadmium. In turn, 0.04 ± 0.01 and 0.1 ± 0.03 μmol total cadmium/mg cell protein were found in the supernatant after the EGTA-washing treatment in acetate- and methanol-grown cells, respectively (i.e., adsorbed Cd²⁺ to the cell outer layers), revealing that most of the cadmium associated with the cells was indeed intra-cellularly trapped.

Due to the extremely low free Cd²⁺ concentration, it seems likely that the complexes formed between cadmium and sulfur compounds, and not the free Cd²⁺, were the species that preferentially entered into cells (Table 1). To further demonstrate that cadmium was indeed inside the cells, methanol-grown cells cultured in 100 μM total CdCl₂ were prepared as previously
Table 2. Effect of cadmium on enzyme activities of the acetoclastic pathway upper part from *Methanosarcina acetivorans*.

| [CdCl₂] μM | Acetate kinase times | Phosphotranacetylase times | CODH/AcCoA synthase times | Carbonic anhydrase times |
|------------|----------------------|---------------------------|--------------------------|--------------------------|
|            |                      |                           | with Acetyl-CoA           | with CO                   |
| 0          | 1                    | 1                         | 1                        | 1                        |
| 0.01       | 1.38±0.27            | Not determined            | 1.0±0.1                  | 1.03±0.11                | 2.9±0.8 **           |
| 0.1        | 1.35±0.18**          | Not determined            | 1.02±0.3                 | 0.66±0.21                | 4.9±3.2 *            |
| 1          | 1.1±0.16             | 0.73±0.15                 | 0.98±0.18                | 0.60±0.16                | 4.2±2.7 *            |
| 10         | Not determined       | 0.77±0.17                 | 0.74±0.2                 | 0.5±0.16                 | 1.7±0.8              |
| 100        | ND                   | 0.47±0.08                 | ND                       | ND                       | ND                   |

All activities were determined by using freshly prepared cytosolic fraction as described in the Methods section. Values are the mean ± SD of at least three independent preparations. Control activities were for AK: 0.75±0.21 U/mg protein (n = 4); for Pta: 1.48±0.8 U/mg protein (n = 4); for CODH/AcCoA synthase with acetyl-CoA: 0.37 ±0.12 U/mg protein (n = 5); and with CO: 0.68±0.11 U/mg protein (n = 3); CA: 26±12 U/mg protein.

*P < 0.05 vs control for independent samples;
**P < 0.05 vs control for paired samples. ND: Not determined.

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Figure 3. Identification of cadmium clusters in *M. acetivorans*. Spectral analysis by HAADF-STEM from methanol-grown cells with 100 μM CdCl₂ (A) or without cadmium (B). C: carbon; O: oxygen, Cu: cupper; S: sulfur; Cd: cadmium.

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reported [18] for HAADF-STEM. Although the images were diffused (Fig. S6), high intracellular contents of electro-dense grains of cadmium and sulfur were definitively identified (Fig. 3). A similar accumulation profile has been reported for cobalt, where the complexes cobalt-chloride and cobalt-citrate are the forms that preferentially are retained in the granular sludge and affect methane production [33]. Such high capacity for cadmium removal in M. acetivorans suggests that this Archaea may have developed strategies to contend against heavy metals different to those reported for the eubacteria domain, which are mainly based on the extrusion of the metal by means of pumps that use ATP as motive force [34].

It has been documented that an excess of zinc, copper, or cobalt, all essential heavy metals, inhibit methanogenesis [9,31,35,36]. In contrast, cadmium seems to be less toxic for methanogenesis in the granular sludge [10], although the free metal concentration or the organic complexes formed were not described. In the present study the free Cd\(^{2+}\) concentration in the culture medium was estimated to be in the pM range. This suggested that M. acetivorans has very high affinity transporters for Cd\(^{2+}\) or that the metal ion may permeate the plasma membrane as a complex with the external cysteine and sulfur.

Concluding remarks

Despite the very low concentration calculated of free Cd\(^{2+}\), this non-essential heavy metal was able to activate a biological process, i.e., methanogenesis in M. acetivorans, due in part to a direct activation of acetylthioenzyme enzymes. M. acetivorans removed and accumulated cadmium; hence, M. acetivorans may become a suitable model for studying the effect of heavy metals on marine methanogens and its mechanisms of heavy metal resistance in the Archaea domain. Moreover, further optimization of the enhanced methane production by cadmium, and other heavy metals, may place this process in the biotechnological leading frontier for generation of biogas.

Supporting Information

Figure S1 Activity of phosphotransacetylase from M. acetivorans. An aliquot of the cytosolic fraction was incubated with the substrates acetyl-Pi and CoA in the absence (■) or presence of 0.1 (▲), 1 (▲) or 10 (▼) \(\mu\)M total CdCl\(_2\). In the absence of protein the CoA concentration remained constant (□). (TIF)

Figure S2 CODH/AcCoAs complex activity from M. acetivorans; representative traces of the activity by adding cytosolic fraction (containing the enzyme complex, ferredoxin and THIPT) and 80 \(\mu\)M AcCoA. Representative trace with: 125 (□), 62 (▲) and 25 (■) \(\mu\)g of cytosolic fraction without cadmium in the absence or presence of 0.01 (▲), 0.1 (▲), 1 (▼) and 10 \(\mu\)M total CdCl\(_2\) (◆). (TIF)

Figure S3 Formation of CO\(_2\) by CA from M. acetivorans in the absence (chromatogram A), or presence of 1 \(\mu\)M CdCl\(_2\). In chromatogram C, cytosolic fraction was previously heated. The different peaks shown represent consecutive sample injections at different times. It should be noted that y-scale is higher for chromatogram B. Under these conditions, CA activity was linear for at least 1 min (Fig. S4). Reverse reaction in turn, produced less reliable values by this method (not shown). (TIF)

Figure S4 Activity of CA in the cytosolicfraction of M. acetivorans in the absence (■) or presence of 0.01 (▲), 0.1 (▲), 1 (▼) or 10 \(\mu\)M total CdCl\(_2\) (◆). A representative data with heated cytosolic fraction in presence of 0.1 \(\mu\)M CdCl\(_2\) is also shown (□). (TIF)

Figure S5 Activation of methanogenesis by cadmium. Cultures on acetate were purged by passing N\(_2\) for 5 min. Then, samples of the head space were withdrawn from the cultures at 0 and 5, 10, 20, 30 and 60 min of incubation with 0 (filled symbols) or 10 \(\mu\)M CdCl\(_2\) (open symbols) for GC analysis. These experiments were started with the addition of 20 mM acetate. Values are the mean ± SD of 3 independent cell preparations. P<0.05 using the Student’s t-test for * vs control (without cadmium). (TIF)

Figure S6 Intracellular cadmium clusters in M. acetivorans. HAADF-STEM projection images of methanol-grown cells cultured in methanol in the absence (A) or in the presence of 100 \(\mu\)M CdCl\(_2\) for 5 days (B). The image in B revealed cadmium grains along the cell (white spots). (TIF)

Text S1 Methods and Results.

Acknowledgments

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Author Contributions

Conceived and designed the experiments: RJC RMS. Performed the experiments: ELS MGSM VHJ RGC. Analyzed the data: RJC. Contributed reagents/materials/analysis tools: RJC. Wrote the paper: RJC RMS.

References

1. Ferry JG, Kastead KA (2007) Methanogenesis, In: Archaea, Molecular and cellular biology Ed. Ricardo Cavicchioli (ASM press), Washington, DC. USA. pp 100–123.
2. Cesar A, Marìn A, Marin-Guirao L, Vita R, Lloret J, et al (2009) Integrative genome analysis of M. acetivorans, an alternative energy producer for large-scale bio-gas production and 80 \(\mu\)M CoA. Representative trace with: 125 (□), 62 (▲) and 25 (■) \(\mu\)g of cytosolic fraction without cadmium in the absence or presence of 0.01 (▲), 0.1 (▲), 1 (▼) and 10 \(\mu\)M total CdCl\(_2\) (◆). (TIF)

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References

1. Ferry JG, Kasteck KA (2007) Methanogenesis, In: Archaea, Molecular and cellular biology Ed. Ricardo Cavicchioli (ASM press), Washington, DC. USA. pp 100–123.
2. Cesar A, Marìn A, Marin-Guirao L, Vita R, Lloret J, et al (2009) Integrative genomics of sediment in Portman Bay [northeast Spain]. Ectocontul Environ Saf 72: 1832–1841.
3. Gillan DC, Danis B, Pernet P, Joly G, Dubois P (2005) Structure of sediment-associated microbial communities along a heavy-metal contamination gradient in the marine environment. Appl Environ Microbiol 71: 679–690.
4. Simpson WR (1981) A critical review of cadmium in the marine environment. Progr Oceanogr 10: 1–70.
5. Villanueva S, Alliono V. (1992) Heavy metals in the coastal zone from Gulf of Mexico and Mexican Caribbean: a review (Metales pesados en la zona costera del Golfo de México y Caribe Mexicano: una revisión). Red Rev Cient Amér Latin Carib Esp y Port 8: 47–61 (Web page: http://redalyc.uaemex.mx/src/ inicio/HomRevRed.jsp?iCveEntRev=370. Accessed 2012 Oct 5).
6. Toes AC, Finke N, Kuenen JG, Muyzer G (2008) Effects of deposition of heavy-metal-polluted harbor mud on microbial diversity and metal resistance in sandy marine sediments. Arch Environ Contam Toxicol 55: 372–385.
7. Florencio L, Field JA, Lettinga G (1994) Importance of cobalt for individual trophic groups in an anaerobic methanogen-degrading consortium. Appl Environ Microbiol 60: 227–234.
8. Jarrell KF, Saulnier M, Ley A (1987) Inhibition of methanogenesis in pure cultures by ammonia, fatty acids, and heavy metals, and protection against heavy metal toxicity by sewage sludge. Can J Microbiol 33: 551–554.
9. Mori K, Hattu M, Kimura R, Takamizawa K (2000) Effect of heavy metals on the growth of a methanogen in pure culture and co culture with a sulfate-reducing bacterium. J Bacteriol 90: 260–265.
10. Altas L (2009) Inhibitory effect of heavy metals on methane-producing anaerobic granular sludge. J Hazard Mater 162: 1551–1556.
11. Sowers KR, Baron SF, Ferry JG (1984) Methanosarcina acetivorans. Appl Environ Microbiol 47: 971–978.
12. Sowers KR (1995) Methanogens: growth and identification. In: Archaea. Halton GJ, Schlesner HJ (eds.) Cold Spring Harbor Press, Cold Spring Harbor, NY. pp 200–245
13. Liu-Silva E, Ramírez-Lima IS, Oliva-Sandoval V, García-García JD, Garcia-Contreras, et al (2011) Removal, accumulation and resistance to chromium in heterotrophic Euglena gracilis. J Hazard Mater 193: 216–224.
14. King TE, Morris RO (1967) Determination of Acid-Labile Sulfide and Sulphhydryl Groups. Methods Enzymol 10: 634–641.
15. Maldon P, Romoee MG (2006) Acute toxicity of heavy metals towards freshwater ciliated protists. Environ Pollut 141: 1–7.
16. Wang M, Wang WX (2011) Cadmium sensitivity, uptake, subcellular distribution and thiol induction in a marine diatom: Recovery from cadmium exposure. Aquat Toxicol 101: 387–393.
17. Abbanat DR, Ferry JG (1990) Synthesis of acetyl coenzyme A by carbon monoxide dehydrogenase complex from acetate-grown Methanosarcina thermophila. J Bacteriol 172: 7143–7150.
18. Mendoza-Co´zatl DG, Rodrı´guez-Zavala JS, Rodrı´guez-Enrı´quez S, Mendoza-Hernández G, Briones-Gallardo R, et al (2006) Phytochelatin-cadmium-sulfide high-molecular-mass complexes of Euglena gracilis. FEBS J 273: 5703–5713.
19. Schoenmakers ThJM, Visser G, Flik G, Theuvenet APR (1992) Chelator: an improved method for computing metal ion concentrations in physiological solutions. Biotech 12: 870–879.
20. Sillen LG, Martell AE (1964) Stability constants of metal-ion complexes. The Chemical Society, Burlington House, London.
21. Hickey RF, Vanderswielen J, Swietenbaum MS (1989) The effect of heavy metals on methane production and hydrogen and carbon monoxide levels during batch anaerobic sludge digestion. Wat Res 23: 207–218.
22. Lane TW, Morel MM (2000) A biological function for cadmium in marine diatoms. Proc Natl Acad Sci USA 97: 4627–4631.
23. Li J, Li Q, Rohlín I, Kim U, Salmon K, et al (2007) Quantitative proteomic and microarray analysis of the archaeon Methanosarcina acetivorans grown with acetate versus methanol. J Proteom Res 6: 759–771.
24. Rohlín I, Guanahini RP (2010) Carbon-dependent control of electron transfer and central carbon pathway genes for methane biosynthesis in the Archaea, Methanosarcina acetivorans strain C2A. BMC Microbiol 10: 62.
25. Aceti DJ, Ferry JG (1988) Purification and characterization of acetyl kinase from acetate-grown Methanosarcina thermophila. Evidence for regulation of synthesis. J Bacteriol 263: 15444–15448.
26. Ingram-Smith C, Gorrell A, Lawerence SH, Iyer P, Smith K, et al (2005) Characterization of the acetyl binding pocket in the Methanosarcina thermophila acetyl kinase. J Bacteriol 187: 2386–2394.
27. Lundie Jr LL, Ferry JG (1989) Activation of acetate by Methanosarcina thermophila. Purification and characterization of phosphotransacetylase. J Biol Chem 264: 18392–18396.
28. Alber BE, Ferry JG (1994) A carbonic anhydrase from Methanosarcina thermophila. Proc Natl Acad Sci USA 91: 6099–6013.
29. Innocenti A, Zimmerman S, Ferry JG, Scozzafava A, Supuran CT (2004) Carbonic anhydrase inhibitors. Induction of the zinc and cobalt gamma-class enzyme from the archaon Methanosarcina thermophila with anions. Bioorg Med Chem Lett 21: 3327–3331.
30. Macauley SR, Zimmerman S, Apolamario EE, Evilia C, Hou YM, et al (2009) The archetype gamma-class carbonic anhydrase (Cah) contains iron when synthesized in vitro. Biochem 48: 817–819.
31. Tripp BC, Bell CR 3rd, Cruz F, Krebs C, Ferry JG (2004) A role for iron in an ancient carbonic anhydrase. J Biol Chem 279: 6683–6687.
32. Pent P, Baldi F (1992) Modulation of chromation (VI) toxicity by organic and inorganic sulfur species in yeasts from industrial wastes. Biomet 5: 179–185.
33. Fermoso FG, Bartacek J, Manzano R, van Leeuwen HP, Lens PNL (2010) Productive anaerobic granular sludge bioreactors with cobalt: Impact of cobalt retention on methanogenic activity. Biore Technol 101: 9429–9437.
34. Silver S, Misra TK, Laddaga RA (1989) DNA sequence analysis of bacterial high-molecular-mass complexes of Euglena gracilis. FEBS J 273: 5703–5713.
35. Karri S, Sierra-Alvarez R, Field JA (2006) Toxicity of copper to acetoclastic and hydrogenotrophic activities of methanogens and sulfate reducers in anaerobic sludge. J Bacteriol 172: 7145–7150.
36. Kim BK, Conway de Macario E, Nolling J, Daniels L (1996) Isolation and characterisation of a copper-resistant methanogen from a copper-mining soil sample. Appl Environ Microbiol 62: 2629–2635.