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Methods for Detecting Microbial Methane Production and Consumption by Gas Chromatography

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Abstract

Methane is an energy-dense fuel but is also a greenhouse gas 25 times more detrimental to the environment than $CO₂$. Methane can be produced abiotically by serpentinization, chemically by Sabatier or Fisher-Tropsh chemistry, or biotically by microbes (Berndt *et al.*, 1996; Horita and Berndt, 1999; Dry, 2002; Wolfe, 1982; Thauer, 1998; Metcalf et al., 2002). Methanogens are anaerobic archaea that grow by producing methane gas as a metabolic byproduct (Wolfe, 1982; Thauer, 1998). Our lab has developed and optimized three different gas chromatograph-utilizing assays to characterize methanogen metabolism (Catlett *et al.*, 2015). Here we describe the end point and kinetic assays that can be used to measure methane production by methanogens or methane consumption by methanotrophic microbes. The protocols can be used for measuring methane production or consumption by microbial pure cultures or by enrichment cultures.

Materials and Reagents

- (Figure 1A)
- **4.** 20 mm Aluminum Seal Crimps (Wheaton, catalog number: 224178-01)
- **5.** 18 G and 22 G BD PrecisionGlide Needle (Becton Dickinson, catalog number: 305195)
- **6.** 22 G BD PrecisionGlide Needle (Becton Dickinson, catalog number: 305155)
- **7.** Hamilton Gas Tight Syringe [1705 Sl 50 μl Syr (22s, 2″, 2) L] (Hamilton Company, catalog number: 80956) (Figure 1)
- **8.** Autosampler vials (National Scientific, catalog number: G4012-1W)** (Figure 1A)

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Equipment

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accommodate other atmospheric and temperature conditions.

2. A standard curve needs to be generated each time the FID detector is turned on. Once the standard has been created, export the following data to Microsoft Excel: Date and Time, Sample_Name, Vial #, Retention Time (min), Height, Area, and Injection_DataFileDirectory. Copy and paste the exported data into the attached spreadsheet's "Standard Data" tab, making sure numbers do not transfer as text and everything is in the correct column. If using Agilent OpenLAB CDS ChemStation software, a report layout can be created that only includes these fields in the order of the spreadsheet.

3. Once the "Standard Data" tab is filled out, fill out the "Constants" tab with injection volume, vial volume, and the volumes of methane gas added to each standard vial, corresponding to what was entered on the "Standards" tab. The spreadsheet will complete the calculations. The standard curve created by default is in Peak Area vs Methane (nmoles). It is important to note that this standard curve is the amount of methane injected into the GC (2 μl injected from each autosampler vial). This can be used in a direct comparison to 2 μl sampled in the same way elsewhere. The number of nmoles has not been multiplied by the dilution factor that would represent the number of moles in the entire autosampler vial. (For a 1.99 ml autosampler vial, the dilution factor is 995.)

4. When using the standard curve created by the attached spreadsheet, data obtained directly from the GC can be compared. When completing the End Point Assay, the peak areas obtained from each sample can be directly plugged into the standard curve equation. In a Kinetic Assay, the volume needs to be adjusted to account for the vial volume displaced by the cell suspension (intact resuspended cells). Once the amount of methane detected by the GC is calculated, dilution factors must be used to calculate the total amount of methane in the headspace of the given culture.

C. End point assay

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- **1.** Grow pure or enrichment cultures in Balch tubes to the desired optical density. For example, for pure cultures of *Methanosarcina acetivorans* C2A grown at 35 °C in HS medium with 125 mM methanol as carbon and energy source with a 1:100 inoculum early stationary phase ($OD_{600} = \sim$ 0.9) is reached between 50 and 75 h (Catlett et al., 2015).
- **2.** Prepare autosampler vials by flushing with air. Stopper, crimp, and label vials.
- **3.** In the fume hood, take culture tubes and insert an 18 G needle three quarters of the way through the blue butyl stopper (Figure 5A). Do not push the needle through the stopper completely.
- **4.** Using a gas tight Hamilton syringe, insert the syringe needle into the 18 G needle (Figure 5B). Be sure that the Hamilton syringe stopcock is closed at this time (Figure 2C). The 18 G needle is used as a guide to help

puncture the butyl stopper with the Hamilton syringe needle while keeping the syringe needle straight.

- **5.** Push the Hamilton syringe needle through the butyl blue stopper until the end of the needle becomes visible in the headspace of the culture tube (Figure 5B). Once again, do not push the 18 G needle completely through the stopper.
- **6.** Open the stopcock of the Hamilton syringe. Quickly and steadily withdraw the Hamilton syringe plunger to the desired volume of headspace. (When using the standard curve described above, it is best to withdraw 50 μl of headspace from the culture.) Once at the desired volume of gas, quickly turn the stopcock of the Hamilton syringe, sealing the gas inside the syringe. Do not let the gas escape to equilibrate with atmospheric pressure (as in section A). Depending on the pressure in the growing culture, the plunger on the Hamilton syringe may continue to move as the gas expands in the syringe.
- **7.** Withdraw the Hamilton syringe and 18 G needle from the culture's butyl stopper.
- **8.** With the Hamilton stopcock still closed, insert the Hamilton syringe needle into a stoppered and crimped autosampler vial.
- **9.** Open the stopcock and inject all the gas from the syringe into the autosampler vial.
- **10.** Close the stopcock and withdraw the Hamilton syringe from the autosampler vial.
- **11.** Once the headspace from all the cultures has been collected, place the vials in the autosampler.
- **12.** Run the "Methane" method on the GC (Table 1).

D. Kinetic assay

- **1.** Before beginning, have the following materials and reagents completely anaerobic and ready in the chamber: Mupirocin stock (3.5 mg/ml) (or appropriate protein synthesis inhibitor depending on the susceptibility profile of the organisms you are assaying), sterile 16 mm test tubes, autosampler vials, autosampler stoppers and crimps, IEC Medilite Microcentrifuge. Prepare a stock of plain medium (no carbon source), and a second stock of growth medium with twice the concentration of carbon source (2x C medium). It is recommended to store these stocks in the anaerobic chamber.
- **2.** Grow cultures in Balch tubes to desired optical density (exponential phase $OD_{600} = -0.3-0.5$. Record OD of the culture.
- **3.** Bring into the anaerobic chamber: Balch tube cultures in a 4 °C Nalgene Labtop Cooler, sterile labeled microcentrifuge tubes, autosampler crimper,

liquid waste container, solid waste container, autosampler vial rack, and forceps.

- **4.** Measure the amount of plain medium and 2x C medium required for the experiment (Table 2). Add protein synthesis inhibitor to desired concentration. For methanogens, add mupirocin to 70 μM.
- **5.** Once items are brought into the chamber, gently resuspend any settled cells in the 10 ml culture. Withdraw 5 ml using a syringe and place in a sterile test tube. Withdraw the remaining 5 ml and place in a separate test tube.
- **6.** Spin down cell cultures in the centrifuge for 5 min at 1,228 x g.
- **7.** Decant supernatant and resuspend each pellet in 5 ml/test tube of plain medium. Use the syringe to disrupt the pellet and spin gently to avoid lysing cells.
- **8.** Spin the resuspended cells for 5 min in the centrifuge at 1,228 x g.
- **9.** As the centrifuge is running, place autosampler vials on an autosampler vial rack. Seven vials are required to assay one strain: five vials for sample replicates, a medium-only control (medium without cells) and a no substrate cells-only control (cells in plain medium only, no carbon source).
- **10.** For 500 μl cell suspensions, Add 250 μl 2x C medium to the five sample replicates and to the medium-only control. Add 250 μl of plain medium to both the medium-only and to the no-cells controls (Table 2).
- **11.** When the centrifuge has halted, immediately decant supernatant. Tap the test tube onto a piece of paper towel to remove all residual media.
- **12.** Use 2 ml plain medium to resuspend and combine pellets into one tube. This should be 2 ml total resuspension for each 10 ml Balch tube culture. Keep cold in a 4 °C Nalgene Labtop Cooler.
- **13.** Add 250 μl of cell suspension to the sample replicates and the cells-only control.
- **14.** Place 200 μl of leftover cell suspension into a labeled microcentrifuge tube. This will be used to measure protein concentration by Bradford assay.
- **15.** Crimp and label vials. Forceps can be helpful in placing stoppers and crimps onto each vial.
- **16.** Remove the vials from the chamber and place in a 35 °C incubator for 5 min before placing vials in the GC autosampler.
- **17.** Do six runs of a seven-vial sequence using the "Methane" method on Agilent GC (Table 1). Each vial should be run in order before repeating the sequence for the next measurement, resulting in 20–40 min for methane to accumulate in each vial between measurements.

- **18.** To measure protein concentration of the cell suspensions, spin down the 200 μl cell suspension saved in step 14 at 1,500 $x g$ for 3 min. Remove supernatant and resuspend with 200 μl ddH₂O. Lyse cells and perform a Bradford with Coomassie reagent and 2 mg BSA standard. Methanogen cells grown in HS medium are easily lysed by resuspension in $ddH₂O$ by osmotic shock, but organisms grown in low-osmolarity medium or that have cell walls may require boiling and/or freeze-thaw and vortexing or sonicating to fully lyse.
- **19.** Graph the methane peak area vs time for each sample (Figure 6). Use the slopes to calculate the amount of methane produced or consumed with time (Table S1.).

Representative data

Figure 1. Crimper, gastight autosampler vials and gas-tight Hamilton syringes A. Crimpers are used to seal autosampler vials using aluminum crimps and rubber stoppers. Hamilton syringes showing open (B) and closed (C) luer fittings.

Figure 2. Dual-sided custom Coy anaerobic chamber showing Medlite clinical centrifuge (blue lid)

Figure 3. Methane gas tank fitted with a septa

Figure 5. Method for preparing gas standards and sampling gas headspace

A. An 18 G needle is pushed into the stopper about three-quarters of the way through. This acts as a guide for the Hamilton needle to push through the stopper and not bend. B. A Hamilton syringe is inserted into an 18 G needle and pushed through the rest of the stopper. Once the end of the Hamilton syringe needle is through the end of the stopper (arrow), headspace can be extracted. Do not push the Hamilton syringe so far into the stopper that the 18 G needle is pushed through the stopper, as this will allow headspace gas to quickly escape.

Figure 6. Example kinetic assay results

Recipes

Notes:

- different osmolarity (low-salt, LS, or high-salt, HS) culture medium (Sowers et al., 1993). If growing other methanogens or methanotrophs, use the appropriate medium recipe for the organism(s) of interest.
- **1.** 200x mupirocin stock

3.5 mg/ml (14 mM).

e. Use vacuum-vortex technique to make the mupirocin stock solution anaerobic (Wolfe and Metcalf, 2010).

- **f.** Store at 4 °C for up to a month.
- **2.** Plain medium (no C source)

Follow the same culture medium recipe as usual, but omit the carbon source. For methanogens that cannot produce methane from $CO₂$ such as Methanosarcina acetivorans, the normal HS medium is prepared. For methanogens and autotrophs that can fix $CO₂$, carbonate, bicarbonate, and $CO₂$ gas should also be eliminated from the recipe and replaced with a buffer at the appropriate pH. For example, for Methanosarcina species that can produce methane from $CO₂$, the bicarbonate in the normal HS medium recipe is replaced with 50 mM 3-N-(morpholino) propanesulfonate (MOPS) (pH 6.8), and the medium is sparged and dispensed into Balch tubes under 100% nitrogen.

3. 2x C medium

Add twice the concentration of carbon source to Plain medium (no C source). For example, when assaying *Methanosarcina* grown on HS medium, if the desired final concentration in the assay is 50 mM methanol, add 100 mM methanol to Plain HS medium to make 2x C HS medium.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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References

- Berndt ME, Allen DE, Seyfried WE. Reduction of $CO₂$ during serpentinization of olivine at 300 °C and 500 bar. Geology. 1996; 24:351–354.
- Catlett JL, Ortiz AM, Buan NR. Rerouting cellular electron Flux to increase the rate of biological methane production. Appl Environ Microbiol. 2015; 81(19):6528–6537. [PubMed: 26162885]
- Dry ME. The fischer-tropsch process: 1950–2000. Catal Today. 2002; 71:227–241.
- Horita J, Berndt ME. Abiogenic methane formation and isotopic fractionation under hydrothermal conditions. Science. 1999; 285(5430):1055–1057. [PubMed: 10446049]

- Metcalf WW, Griffin BM, Cicchillo RM, Gao J, Janga SC, Cooke HA, Circello BT, Evans BS, Martens-Habbena W, Stahl DA, van der Donk WA. Synthesis of methylphosphonic acid by marine microbes: a source for methane in the aerobic ocean. Science. 2012; 337(6098):1104–1107. [PubMed: 22936780]
- Sowers KR, Boone JE, Gunsalus RP. Disaggregation of methanosarcina spp. and growth as single cells at elevated osmolarity. Appl Environ Microbiol. 1993; 59(11):3832–3839. [PubMed: 16349092]

Thauer RK. Biochemistry of methanogenesis: a tribute to Marjory Stephenson. 1998 Marjory Stephenson Prize Lecture. Microbiology. 1998; 144(Pt 9):2377–2406. [PubMed: 9782487]

Wolfe RS. Biochemistry of methanogenesis. Experientia. 1982; 38:198–201.

Wolfe RS, Metcalf WW. A vacuum-vortex technique for preparation of anoxic solutions or liquid culture media in small volumes for cultivating methanogens or other strict anaerobes. Anaerobe. 2010; 16(3):216–219. [PubMed: 20004732]

Table 1

Gas chromatograph "Methane" method settings

Table 2

Kinetic assay medium volumes and controls

 $\frac{k}{1}$ In a 1.5 ml vial with an actual volume of 1.99 cm³