

Effect of trace metals on ethanol production from synthesis gas by the ethanologenic acetogen, *Clostridium ragsdalei*

Jyotisna Saxena · Ralph S. Tanner

Received: 28 April 2010/Accepted: 21 July 2010/Published online: 10 August 2010
© Society for Industrial Microbiology 2010

Abstract The effect of trace metal ions (Co^{2+} , Cu^{2+} , Fe^{2+} , Mn^{2+} , Mo^{6+} , Ni^{2+} , Zn^{2+} , SeO_4^- and WO_4^-) on growth and ethanol production by an ethanologenic acetogen, *Clostridium ragsdalei* was investigated in CO:CO_2 -grown cells. A standard acetogen medium (ATCC medium no. 1754) was manipulated by varying the concentrations of trace metals in the media. Increasing the individual concentrations of Ni^{2+} , Zn^{2+} , SeO_4^- and WO_4^- from 0.84, 6.96, 1.06, and 0.68 μM in the standard trace metals solution to 8.4, 34.8, 5.3, and 6.8 μM , respectively, increased ethanol production from 35.73 mM under standard metals concentration to 176.5, 187.8, 54.4, and 72.3 mM, respectively. Nickel was necessary for growth of *C. ragsdalei*. Growth rate (μ) of *C. ragsdalei* improved from 0.34 to 0.49 (day^{-1}), and carbon monoxide dehydrogenase (CODH) and hydrogenase (H_2ase)-specific activities improved from 38.45 and 0.35 to 48.5 and 1.66 U/mg protein, respectively, at optimum concentration of Ni^{2+} . At optimum concentrations of WO_4^- and SeO_4^- , formate dehydrogenase (FDH) activity improved from 32.3 to 42.6 and 45.4 U/mg protein, respectively. Ethanol production and the activity of FDH reduced from 35 mM and 32.3 U/mg protein to 1.14 mM and 8.79 U/mg protein, respectively, upon elimination of WO_4^- from the medium. Although increased concentration of Zn^{2+} enhanced growth and ethanol production, the activities of CODH, FDH, H_2ase and alcohol dehydrogenase (ADH) were not

affected by varying the Zn^{2+} concentration. Omitting Fe^{2+} from the medium decreased ethanol production from 35.7 to 6.30 mM and decreased activities of CODH, FDH, H_2ase and ADH from 38.5, 32.3, 0.35, and 0.68 U/mg protein to 9.07, 7.01, 0.10, and 0.24 U/mg protein, respectively. Ethanol production improved from 35 to 54 mM when Cu^{2+} was removed from the medium. The optimization of trace metals concentration in the fermentation medium improved enzyme activities (CODH, FDH, and H_2ase), growth and ethanol production by *C. ragsdalei*.

Keywords Bioethanol · Gasification · Synthesis gas · Acetogens · *C. ragsdalei* · Metalloenzymes · Trace metals

Introduction

The environmental concerns over the use of petroleum-based fuels and their limited supplies have generated a great interest in producing ethanol as an alternative fuel. The use of bioethanol (ethanol from biomass) reduces the dependence on imported oil, supports agriculture, and limits greenhouse gas emissions. In the United States, bioethanol is primarily produced from the fermentation of corn starch, which is presently facing critical issues such as potential diversion of food and fodder-grade feedstocks towards biofuel production, its indirect effects on land-use change, and its limited supply [24]. Hence, to make the bioethanol production processes sustainable, rigorous criteria must be considered and best practices in terms of choice of feedstocks and its direct and indirect environmental and socio-economic effects should be utilized [11]. Bioethanol produced from lignocellulosic biomass grown on agriculturally marginal lands (e.g., switchgrass) and waste biomass (e.g., agricultural and forestry wastes) could

J. Saxena · R. S. Tanner
Department of Botany and Microbiology,
University of Oklahoma, Norman, OK 73019, USA

J. Saxena (✉)
Coskata, Inc., 4575 Weaver Parkway, Suite 100,
Warrenville, IL 60555, USA
e-mail: Jsaxena@coskata.com

compete with present supplies [21] and provide significant environmental and economic benefits over corn starch-based ethanol [11].

Lignocellulosic biomass can be converted to ethanol by direct fermentation of sugars obtained by acidic/enzymatic hydrolysis of the biomass (cellulosic ethanol) [19]. Although this process has tremendous potential, and over the past several years great advances have been made in this area [17], there are several disadvantages associated with this technology, such as the difficulties in handling and converting diverse sources of biomass to ethanol, and the lignin component of biomass is left unutilized. A promising approach for conversion of biomass to ethanol is the stepwise process of gasification and microbial fermentation [30]. In gasification, all biomass components including lignin are gasified to synthesis gas (syngas: a mixture of primarily CO, CO₂, and H₂) [30]. Syngas is fermented to several commodity chemicals, including ethanol by a group of obligatory anaerobic bacteria known as acetogens [30]. As a feed-stock flexible process, syngas fermentation can utilize a wide variety of biomass sources that are combustible and non-food based, such as municipal solid waste, tires, and residual agricultural biomass [30].

Acetogens utilize the acetyl-CoA/Wood-Ljungdahl pathway for production of acetyl-CoA, a precursor of cellular biomass, acetate, and ethanol. Key enzymes of this pathway include formate dehydrogenase (FDH), bifunctional carbon monoxide dehydrogenase/acetyl CoA synthase (CODH/ACS), and hydrogenase (H₂ase), all of which are metalloenzymes [9]. In addition, solvent-producing clostridia also have iron- and zinc-containing alcohol dehydrogenase (ADH) [15, 26] that plays a key role by catalyzing the reduction of acetyl CoA to ethanol [6].

Considering the key role of metalloenzymes in the metabolism of acetogens, a study was undertaken to investigate the effect of trace metals (Co²⁺, Cu²⁺, Fe²⁺, Mn²⁺, Mo⁶⁺, Ni²⁺, Zn²⁺, SeO₄²⁻, and WO₄²⁻) on growth, production of ethanol, production of acetate and enzyme activities (CODH, FDH, H₂ase, and ADH) by a recently isolated ethanologenic acetogen, *Clostridium ragsdalei* using syngas components. *C. ragsdalei* was isolated from duck pond sediment, and produces ethanol and acetate from syngas components [12] (US Pat. No. US 2008/0057554 A1). It was hypothesized that the optimization of trace metal concentrations in the culture medium might enhance the activity of metalloenzymes, thus improving growth and ethanol production by *C. ragsdalei*.

Materials and methods

Bacterial strain, media, and growth conditions

The bacterium used in this study was *Clostridium ragsdalei* (ATCC PTA 7826), which was isolated in our laboratory

from duck pond sediment in Brandt Park, Norman, OK, USA. The standard acetogen medium (ATCC medium no. 1754) for cultivation of *C. ragsdalei* was prepared using strict anaerobic techniques [2] under an atmosphere of N₂:CO₂ (80:20). The medium (per liter) contained: mineral solution, 25 ml [29]; trace metal solution, 10 ml [29]; vitamin solution, 10 ml [29]; 2-(N-morpholino)ethanesulfonic acid (MES), 20 g; yeast extract (Difco, Becton Dickinson, Sparks, M.D.), 0.5 g; 0.1% resazurin, 0.1 ml and cysteine-sulfide reducing agent, 2.5 ml [29]. The pH of the medium was adjusted to 6.1 using NaOH. The trace metal solution (per liter) contained MnSO₄·H₂O (1.0 g), Fe(NH₄)₂(SO₄)₂·6H₂O (0.8 g), CoCl₂·6H₂O (0.2 g), ZnSO₄·7H₂O (0.2 g), CuCl₂·2H₂O (0.02 g), NiCl₂·6H₂O (0.02 g), Na₂MoO₄·2H₂O (0.02 g), Na₂SeO₄ (0.02 g), and Na₂WO₄ (0.02 g). The concentrations of individual trace metals in the media are presented in Table 1. Metals present in the yeast extract were removed as metal ion phosphates using a procedure derived from dephosphorylating media components [20].

Clostridium ragsdalei was grown in anaerobic culture tubes (Bellco Glass, Inc., Vineland, NJ, USA) containing 5 ml of medium and a gas mixture of CO:N₂:CO₂ (70:24:6 pressurized to 230 kPa gauge) in the head space. Since CO is sparingly soluble in water, the tubes were incubated on their sides at 36°C on a rotary shaker (100–120 rpm) to maximize gas–liquid mixing.

Fermentation experiment

Batch fermentation experiments were conducted to examine the effect of trace metals (Co²⁺, Cu²⁺, Fe²⁺, Mn²⁺, Mo⁶⁺, Ni²⁺, Zn²⁺, SeO₄²⁻, and WO₄²⁻) on growth and ethanol production by *C. ragsdalei*. The concentration of trace metals in standard trace metals solution (Table 1) is denoted as 1X throughout the study. The concentrations of each trace metal was either increased ten-fold (10X) with respect to standard metal concentration (1X) (Table 1) or eliminated (0X) from the media. Once effective metals for growth and ethanol production were identified, different

Table 1 Concentration of trace metals in standard growth medium, ATCC 1754 (1X)

Trace metal	(μM)
Co	8.40
Cu	1.17
Fe	20.40
Mn	59.17
Mo	0.83
Ni	0.84
Se	1.06
W	0.68
Zn	6.96

concentrations (1X, 5X, 10X, and 20X) of each metal were titrated to find their optimal concentrations for enhanced ethanol production. Fermentation experiments were performed in triplicate in 160-ml Wheaton serum vials containing 10 ml of medium. *C. ragsdalei* cells were transferred three times in the media with each trace metal concentrations to minimize the metal ion carry over via inoculum into the experimental media. A 1% inoculum was used in all experiments. The bottles were fed with CO (440-kPa gauge) and incubated at 36°C on a rotary shaker. Once the cells were grown, the headspace gases were exchanged with CO (440-kPa gauge) every 24 h for 10 days. At the end of the experiments, final pH and growth were recorded and the samples were analyzed for acetate and ethanol.

A time course experiment was conducted on the optimized concentrations metals and to study their effects on growth and fermentation products formed over a period of 10 days. The time course experiment was conducted in 500-ml Wheaton serum bottles with 50 ml of medium. Experiments were conducted according to the procedure described above, except that samples (1.5 ml) were collected every 24 h for 10 days. The samples were analyzed for pH, growth, and fermentation products.

Analytical methods

Growth was monitored by measuring optical density (OD) at 600 nm with a Beckman Coulter DU 640 spectrophotometer (Bausch and Lomb, Rochester, NY). Growth rate (μ) was derived from OD data as described elsewhere [4]. The fermentation products (i.e., acetate and ethanol) were analyzed quantitatively using a Shimadzu GC-8A gas chromatograph (Shimadzu, Maryland, USA) equipped with a flame ionization detector (FID) and helium was used as the carrier gas. The column and the injector/detector temperatures were maintained at 155 and 200°C, respectively. The glass column (2.6 m) was packed with Carbopack BDA (80/120 mesh) and 4% Carbowax 20 M resin (Supelco Inc., Bellefonte, PA). Data were analyzed with a Shimadzu C-R8A integrator.

Enzyme assays

Enzyme activities (FDH, CODH, H₂ase, and ADH) were determined spectrophotometrically as the substrate-dependent reduction of methyl viologen (MV) in 2-ml cuvettes as described previously [18]. The cultures in the enzyme assays were grown in 500-ml Wheaton serum bottles with 50 ml of media as stated above. The exponential phase cultures were used for enzyme assays. Protein concentrations were estimated using bicinchoninic acid method [28] with bovine serum albumin as a standard.

Results

Screening of trace metals for their effect on ethanol production

The effect of different trace metals on growth rate and production of ethanol and acetate by *C. ragsdalei* is presented in Table 2. *C. ragsdalei* produced 35 mM of ethanol and 9 mM of acetate under the standard medium (1X) condition. At higher concentrations (10X) of Ni²⁺, Zn²⁺, SeO₄⁻ and WO₄⁻ ethanol production increased to 176, 190, 75, and 55 mM, respectively. Ethanol production decreased to 27.01, 6.30, 22.02, and 1.55 mM when Co²⁺, Fe²⁺, Mo²⁺ and WO₄⁻ were eliminated (0X) from the media, indicating that these metals were required for ethanol production. An increase in ethanol production to 54 mM was noticed when Cu²⁺ was removed (0X) from the media. Ethanol production did not change considerably at higher concentration (10X) of Cu²⁺, Co²⁺, Fe²⁺, and Mn²⁺, and also when Mn²⁺, Zn²⁺, and SeO₄⁻ were omitted (0X) from the media. A decrease in ethanol production to 23.64 mM was observed at higher concentration (10X) of Mo²⁺.

Acetate production increased to 15 and 18.6 mM at increased concentrations (10X) of Ni²⁺ and Zn²⁺, respectively. An increase in acetate production to 15.34 and 18.64 mM was also observed when Fe²⁺ and WO₄⁻ were eliminated (0X) from the media. There were no appreciable changes in acetate production at greater concentration (10X) of Co²⁺, Cu²⁺, Fe²⁺, Mn²⁺, Mo⁶⁺, SeO₄⁻, and WO₄⁻, and also when Co²⁺, Cu²⁺, Mn²⁺, Mo⁶⁺, Zn²⁺, and SeO₄⁻ were omitted (0X) from the media.

The growth rate of *C. ragsdalei* improved at increased concentrations (10X) of Ni²⁺, Zn²⁺, and WO₄⁻. Nickel was necessary for growth of *C. ragsdalei*. The growth rate of *C. ragsdalei* for remaining metals (Co²⁺, Cu²⁺, Fe²⁺, Mn²⁺, Mo⁶⁺, and SeO₄⁻) was similar to that with standard metals (1X) in growth medium. The final pH of the media was similar under the various metal concentrations.

Effect of optimized concentrations of Ni²⁺, Zn²⁺, WO₄⁻, and SeO₄⁻ on growth rate, ethanol production, and acetate production

The optimal concentrations of Ni²⁺, Zn²⁺, SeO₄⁻, and WO₄⁻ for ethanol production were 8.5, 35, 5, and 7 μM and the concentrations of ethanol produced at these concentrations were 180, 200, 60, and 90 mM, respectively (Fig. 1). Ethanol production at the standard metal concentrations (1X) was 35 mM. The optimum concentrations of Ni²⁺, Zn²⁺, SeO₄⁻, and WO₄⁻ increased cell growth, which became more pronounced by the addition of Ni²⁺ and Zn²⁺. Acetate production at optimum concentrations

Table 2 Effect of trace metals on growth rate (μ), final pH (pH_f), production of ethanol, and production of acetate by *C. ragsdalei* after 10 days of growth

Trace metals	Growth rate (day $^{-1}$)	pH _f	Ethanol (mM)	Acetate (mM)
1X ^a	0.34 ± 0.006	5.9	35.73 ± 2.05	8.87 ± 0.31
Ni	0X	No growth	6.1	No growth
	10X	0.49 ± 0.002	5.8	176.52 ± 5.62
Zn	0X	0.34 ± 0.018	5.9	33.48 ± 1.42
	10X	0.48 ± 0.002	5.6	187.80 ± 8.56
Se	0X	0.30 ± 0.012	5.9	38.32 ± 2.12
	10X	0.30 ± 0.012	5.9	54.35 ± 3.23
W	0X	0.30 ± 0.015	5.9	1.14 ± 0.04
	10X	0.44 ± 0.003	5.9	72.29 ± 1.80
Cu	0X	0.30 ± 0.002	5.9	54.09 ± 2.62
	10X	0.30 ± 0.008	5.9	36.63 ± 1.57
Fe	0X	0.33 ± 0.004	5.9	6.30 ± 0.92
	10X	0.34 ± 0.007	5.9	35.87 ± 1.72
Co	0X	0.31 ± 0.001	5.9	27.01 ± 1.31
	10X	0.34 ± 0.001	5.9	37.34 ± 0.69
Mo	0X	0.31 ± 0.004	5.9	22.02 ± 0.90
	10X	0.31 ± 0.001	5.9	23.64 ± 1.42
Mn	0X	0.31 ± 0.001	5.9	33.11 ± 1.46
	10X	0.31 ± 0.001	5.9	36.78 ± 0.24

^a 1X standard concentrations of all the trace metals, 0X elimination of trace metal, 10X tenfold higher metal concentration than 1X

of Ni²⁺, Zn²⁺, and SeO₄²⁻ was similar to that produced under control conditions.

Time course experiment

Cultures of *C. ragsdalei* demonstrated improved growth under optimized concentration of Ni²⁺ and Zn²⁺ compared to the control (1X) (Fig. 2) and remained in stationary phase of growth for 7–8 days before cell lysis was observed. However, cells lysed in the standard medium (1X) after 3–4 days of growth, which was also observed for optimum concentrations of WO₄²⁻ and SeO₄²⁻ (Fig. 3). When Ni⁺, Zn⁺, SeO₄²⁻, and WO₄²⁻ were used at their optimum concentrations, and Cu²⁺ was eliminated from the trace metals solution (CuNiZnSeW), a significant increase in growth was observed, and cells remained in the stationary phase until the end of the experiment (10 days). An increase in ethanol production was observed once the cultures entered into the exponential phase, which continued to increase until the stationary phase (Figs. 2, 3). An increase in acetate production was observed during the exponential phase of growth in all metals

concentrations tested, which started decreasing once *C. ragsdalei* cells entered into the stationary phase of growth. The transient reduction in acetate concentration was associated with an increase in ethanol concentration (Figs. 2, 3).

Effect of Ni²⁺, Zn²⁺, SeO₄²⁻, and WO₄²⁻ on enzyme activities in *Clostridium ragsdalei*

The optimum concentration of Ni²⁺ stimulated the activities of CODH and H₂ase; whereas, the activity of FDH was enhanced at optimum concentrations of SeO₄²⁻ and WO₄²⁻ (Table 3). Inclusion of optimum concentrations of SeO₄²⁻ and WO₄²⁻ together in the medium had the same effect on FDH activity as the individual additions of SeO₄²⁻ and WO₄²⁻ (Table 3). Enzyme activity of CODH, FDH, H₂ase, and ADH were reduced when Fe²⁺ was removed from the medium, whereas at increased concentration of Fe²⁺ (10X), there was no change in enzyme activities as compared to those under the standard metal concentration (1X) (Table 3). Although the optimum concentration of Zn²⁺ increased the growth and ethanol production (Fig. 1),

Fig. 1 Effect of different concentrations of nickel (Ni), zinc (Zn), selenite (Se), and tungstate (W), on growth rate (/day), production of ethanol, and production of acetate by *C. ragsdalei*

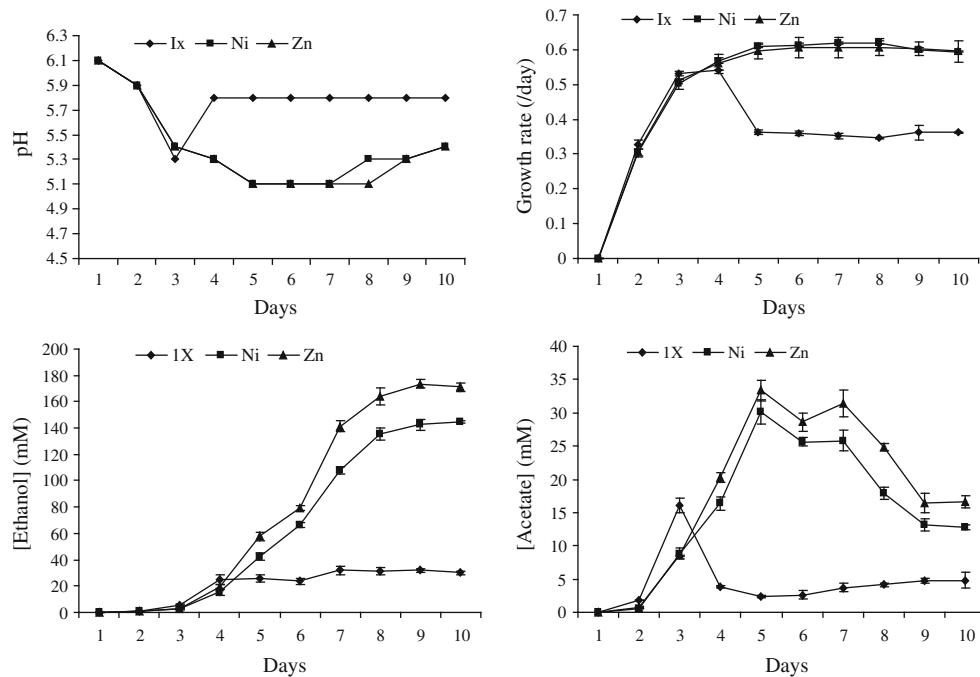
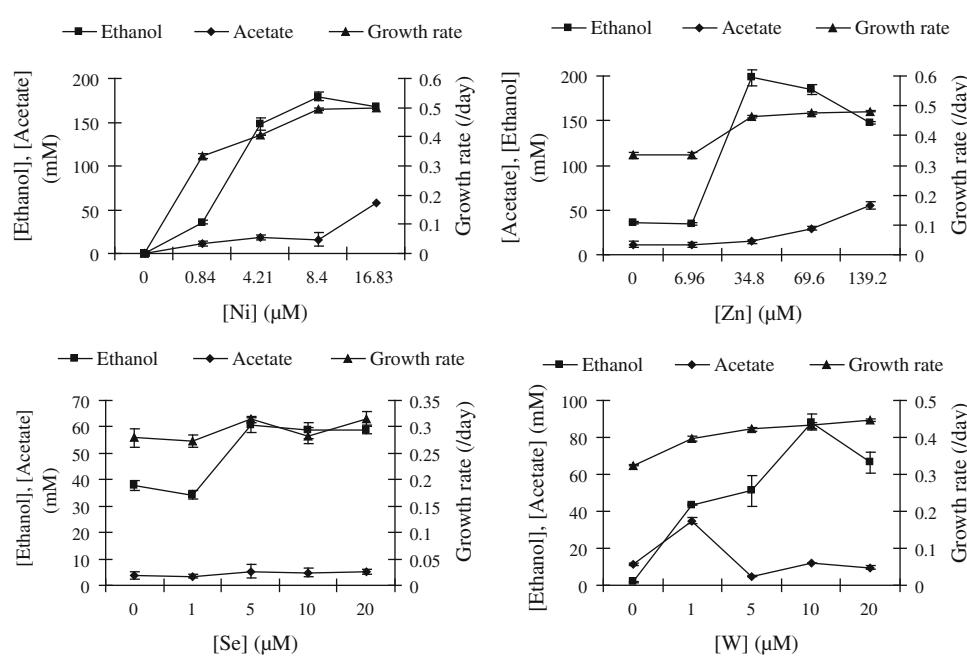


Fig. 2 Effect of optimal concentrations of nickel (Ni) and zinc (Zn) on final pH, growth rate (/day), production of ethanol, and production of acetate by *C. ragsdalei*

enzyme activities were not affected by the concentration of Zn^{2+} (Table 3).

Discussion

Acetogens utilize the acetyl-CoA/Wood-Ljungdahl pathway for production of acetyl-CoA, a precursor of cell

biomass, acetate, and ethanol [9]. Hence, to improve the growth and fermentation products, it is necessary to improve the carbon flow to acetyl-CoA, which requires upstream manipulations in this pathway. In this study, the activities of the key metalloenzymes of this pathway were enhanced by optimizing the concentrations of trace metals in the growth medium, which improved growth and ethanol production by *C. ragsdalei*.

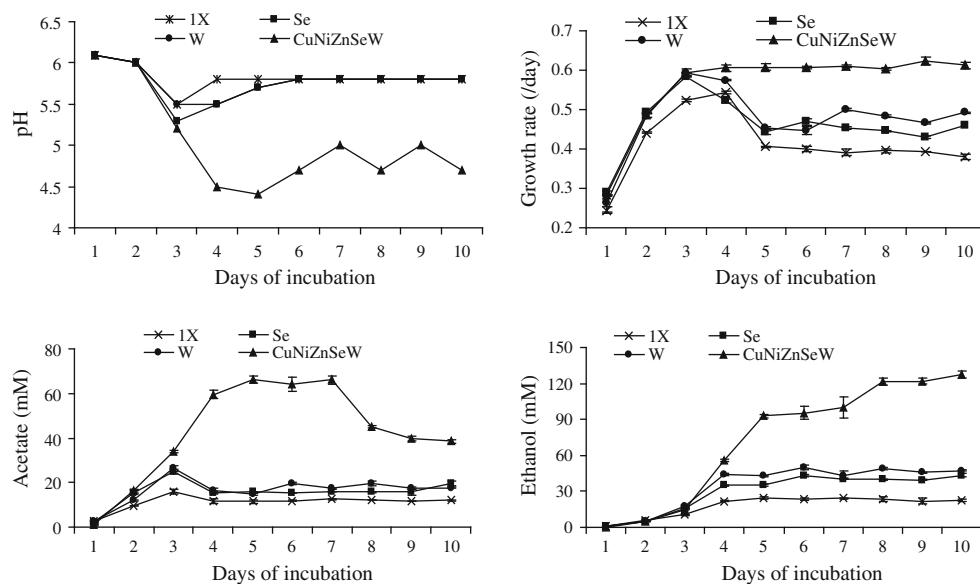


Fig. 3 Effect of optimal concentrations of selenate (Se), tungstate (W), and optimal concentrations of copper-nickel-zinc-selenite-tungstate combined together (CuNiZnSeW) on final pH, growth rate (/day), production of ethanol, and production of acetate by *C. ragsdalei*

Table 3 Effect of trace metals on activities (U/mg protein) of carbon monoxide dehydrogenase (CODH), formate dehydrogenase (FDH), hydrogenase (H₂ase), and alcohol dehydrogenase (ADH) by *C. ragsdalei*

Specific activity = 1 μmol of substrate reduced.min⁻¹ mg⁻¹ of protein

^a 1X standard concentrations of all the trace metals, 0X elimination of trace metals, 5X fivefold higher concentration than 1X, 10X tenfold higher concentration than 1X

	Growth conditions ^a	FDH	CODH	H ₂ ase	ADH
1X		32.25 ± 1.41	38.45 ± 1.68	0.35 ± 0.01	0.68 ± 0.01
Ni					
0X		No growth	No growth	No growth	No growth
10X		34.6 ± 1.13	48.46 ± 1.57	1.66 ± 0.06	0.65 ± 0.12
Se					
0X		31.97 ± 0.92	38.01 ± 0.49	0.32 ± 0.03	0.83 ± 0.06
5X		42.62 ± 0.73	37.07 ± 3.35	0.28 ± 0.01	0.61 ± 0.03
W					
0X		8.79 ± 0.16	36.24 ± 0.29	0.33 ± 0.03	0.77 ± 0.01
10X		45.44 ± 0.21	33.57 ± 0.79	0.41 ± 0.08	0.68 ± 0.05
Se (5X) + W (10X)		35.46 ± 0.34	36.25 ± 0.41	0.31 ± 0.02	0.71 ± 0.13
Zn					
0X		31.96 ± 1.21	35.64 ± 2.50	0.32 ± 0.03	0.65 ± 0.03
5X		34.02 ± 0.08	36.81 ± 2.39	0.44 ± 0.04	0.75 ± 0.03
Fe					
0X		7.01 ± 0.62	9.07 ± 0.04	0.10 ± 0.04	0.24 ± 0.02
10X		35.65 ± 0.33	37.51 ± 0.50	0.39 ± 0.05	0.63 ± 0.01

The ethanol production and enzyme activities (CODH, FDH, H₂ase, and ADH) of *C. ragsdalei* were reduced when Fe²⁺ was removed (0X) from the medium. All the key metalloenzymes of acetyl-CoA pathway (CODH, FDH, and H₂ase) are iron-sulfur proteins [5, 10, 32]. Hence, a reduction in enzyme activities (CODH, FDH, H₂ase, and ADH) and ethanol production by *C. ragsdalei* upon elimination of Fe²⁺ from the growth media was anticipated. However, increasing Fe²⁺ concentration (10X) had no effect on the growth, production of ethanol, or production of acetate, which indicates that Fe²⁺ concentration in the

growth medium became saturating at a concentration exceeding 20.4 μM (1X) for the growth and metabolism of *C. ragsdalei*.

FDH catalyzes the first reaction on the methyl branch of the acetyl Co-A pathway, where it reduces CO₂ to formate [9]. The metal co-factors of FDH in clostridia are reported to be variable, where iron, tungsten, selenium, and/or molybdenum have been shown to be the components of FDH [16, 31, 32]. FDH in acetogens contain selenium and tungsten [14, 31, 32]; whereas, in non-acetogenic clostridia such *C. formicoaceticum*, molybdenum-containing FDH is

present [16]. In acetate-producing acetogens, the presence of tungsten and selenium in FDH has been suggested to function as cofactors, promoting the FDH-mediated reduction of CO₂ [32]. *C. ragsdalei* produced acetate and ethanol as the fermentation products when grown on CO:CO₂. Whole-cell FDH activity of *C. ragsdalei* was enhanced at the optimum concentration of tungstate and selenite, and reduced upon elimination of tungstate and iron, suggesting that FDH of *C. ragsdalei* may be a tungsten-selenium-iron protein.

At the optimum concentration of Ni²⁺, an increase in ethanol production by *C. ragsdalei* was observed that was associated with increased CODH activity. CODH in acetogenic bacteria such as *M. thermoacetica* occurs as a bifunctional CODH/ACS enzyme where CODH generates CO from CO₂, and the ACS combines the CO with CoA and a methyl group to form acetyl-CoA [10]. Both CODH and ACS contain iron-nickel-sulfur metal clusters (CODH: Fe-[NiFe₃S₄] and ACS: [Fe₄S₄]-Cys-Gly-Cys-Ni-Cys-Gly-Cys-Ni) at their active sites [10]. Therefore, CODH/ACS and the associated metals (Fe and Ni) play a key role in the metabolism of acetogens.

In addition to Fe²⁺ and Ni²⁺, Cu²⁺ also regulates the activity of ACS in acetogens, where Cu²⁺ has a negative effect on ACS activity [3, 10, 27]. The metal cluster of ACS contains binuclear metal sites: a proximal metal site (M_p) and a distal metal site (M_d), both of which contain Ni²⁺ under the condition where no Cu²⁺ is present in the growth medium. However, in the presence of Cu²⁺, the proximal metal site in the A-cluster can undergo a metal substitution resulting in a binuclear site composition of Cu–Ni and Ni–Ni [10, 27]. The Ni–Ni form of ACS is active whereas Cu–Ni form of enzyme is inactive, which eliminates ACS activity (measured by ability to generate the NiFeC EPR signal) [27]. Although ACS activity in the present study was not measured, an increase in ethanol production was observed upon removing Cu²⁺ from the growth medium, which could be due to inhibition of ACS activity by Cu²⁺. Increasing the concentration of Cu²⁺ had no change on growth and product formation, suggesting Cu²⁺ concentration in standard trace metals solution (1X) was enough to cause ACS inhibition, probably due to the fact that the cells of acetogenic bacteria can concentrate Cu²⁺ over 85-fold and Ni²⁺ at least tenfold from the growth medium [27].

Hydrogenases catalyze the reversible oxidation of molecular hydrogen, and are of two types in bacteria: NiFe and Fe-only H₂ase [5]. The NiFe-H₂ase is typically involved in oxidation of hydrogen; whereas Fe-only H₂ase catalyzes the reduction of protons as terminal electron acceptors to yield hydrogen [5]. Hydrogenase plays a key role in the metabolism of acetogens, most of which can grow autotrophically, as well as heterotrophically [9]. During autotrophic

growth on H₂ and CO₂, H₂ase-mediated oxidation of H₂ provides reducing equivalents for fixation of CO₂ into acetate and cell biomass [9]. In contrast, H₂ase in acetogens growing on carbohydrates dispose off the excess reducing equivalents by reducing protons into H₂ [1]. In addition, existence of multiple H₂ases that account for different roles under different growth conditions have also been reported in acetogens [9]. H₂ase of *C. pasteurianum*, a Fe-only H₂ase has been most extensively studied both biochemically and spectroscopically [22]. H₂ase purified from fructose grown cells of *Acetobacterium woodii* also does not contain nickel, and exhibits spectroscopic properties similar to that of Fe-only H₂ase of *C. pasteurianum* [22, 25]. To our knowledge, all purification and spectroscopic studies of H₂ase in acetogens have been done from heterotrophically grown acetogens, which has decreased the probability of isolating Fe–Ni hydrogenase, and have ruled out the presence of nickel in the enzyme. However, a genome sequence analysis of *M. thermoacetica* has revealed a gene cluster homologue of genes that make up an operon required for maturation of the Ni–Fe H₂ase [23]. In the present study, H₂ase activity was stimulated by increased concentrations of Ni²⁺, and reduced by the elimination of Fe²⁺ from the media. Thus, H₂ases present in CO:CO₂-grown *C. ragsdalei* may be Fe-only and/or NiFe H₂ases. Since, CO:CO₂ was used as growth substrate in present study, low H₂ase activities obtained were probably due to inhibition of H₂ase by CO [25].

In solventogenic clostridia, ADH catalyzes the reduction of acetyl CoA to ethanol [6]. The biochemical and genetic studies on ADH from different clostridia [6, 13] have indicated the role of ADH in solvent production. Clostridial ADHs have been shown to contain zinc [15] and iron [26]. A reduction in ADH activity of *C. ragsdalei* was observed when iron was removed from the media, suggesting that ADH of *C. ragsdalei* may be an iron-containing enzyme. Furthermore, supplementing the growth medium of *C. ragsdalei* with optimized levels of Zn²⁺ resulted in increased ethanol production and improved cell growth; however, no changes in enzyme activities (CODH, FDH, H₂ase, and ADH) were observed upon varying Zn²⁺ concentration. These results indicate that both types of ADH may be present in *C. ragsdalei*.

The effects of trace metals on growth, homoacetogenesis, and activities of enzymes of the acetyl-CoA pathway (CODH, FDH, and H₂ase) [8, 14] have been mostly studied in heterotrophically grown acetogens. However, the effects of trace metals on product formation under chemolithotrophic growth conditions have not been measured. The activities of key enzymes of Wood-Ljungdahl pathway (CODH, FDH, and H₂ase) may vary under heterotrophic and chemolithotrophic growth conditions [7, 18] with higher enzyme activities obtained when cultures were grown on CO:CO₂ and/or H₂:CO₂ than when grown on

sugars [7, 18]. In the present study, the effects of trace metals on growth, enzyme activities, and ethanol production by an acetogen (*C. ragsdalei*) grown on CO:CO₂ was studied, which can provide further insights about medium manipulation for increased ethanol production by solventogenic acetogens via syngas fermentation.

Conclusions

Bioethanol production from fermentation of lignocellulosic biomass-derived syngas is a promising alternative for biofuel production. This process offers advantages such as the utilization of whole biomass, including lignin, irrespective of the biomass source and the elimination of the extensive pretreatment steps to obtain fermentable sugars. One of the major limitations of syngas fermentation is the low ethanol yield of biocatalysts. In the present study, ethanol production by *C. ragsdalei* was improved four-fold by optimizing the trace metal concentrations in the growth medium, emphasizing the importance of medium optimization for improving the ethanol fermentation from syngas. The increase in ethanol production by *C. ragsdalei* was also substantiated by increased cell growth and metalloenzyme activities of the acetyl-CoA pathway. The presence of Cu²⁺ in the medium negatively affected the ethanol production by *C. ragsdalei*. The optimum medium concentrations of Cu²⁺, Ni²⁺, Zn²⁺, SeO₄²⁻, and WO₄²⁻ for ethanol production were 0, 8.5, 35, 7, and 5 μM, respectively. Evaluation of other media components (minerals and vitamins) is warranted for further improvement of the growth and ethanol production by *C. ragsdalei* from syngas.

Acknowledgments This research was supported in part by USDA-CSREES Special Grant awards 2005-34447-15711 and 2006-34447-16939. We would like to thank Dr. James A. Zahn for his constructive suggestions in this manuscript.

References

1. Adams MWW, Mortenson LE, Chen J-S (1981) Hydrogenase. *Biochem Biophys Acta* 594:105–176
2. Balch WE, Wolfe RS (1976) New approach to the cultivation of methanogenic bacteria: 2-mercaptopethanesulfonic acid (HS-CoM)-dependent growth of *Methanobacterium ruminantium* in a pressurized atmosphere. *Appl Environ Microbiol* 32:781–791
3. Bramlett MR, Tan X, Lindahl PA (2003) Inactivation of acetyl CoA synthase/carbon monoxide dehydrogenase by copper. *J Am Chem Soc* 125:9316–9317
4. Burdette DS, Jung S-H, Shen G-J, Hollingsworth RI, Zeikus JG (2002) Physiological function of alcohol dehydrogenase and long-chain (C30) fatty acids in alcohol tolerance of *Thermoanaerobacter ethanolicus*. *Appl Environ Microbiol* 68:1914–1918
5. Cammack R (1999) Hydrogenase sophistication. *Nature* 397: 214–215
6. Chen JS (1995) Alcohol dehydrogenase: multiplicity and relatedness in the solvent-producing clostridia. *FEMS Microbiol Rev* 17:263–273
7. Clark JE, Ragsdale SW, Ljungdahl LG, Wiegel J (1982) Levels of enzymes involved in the synthesis of acetate from carbon dioxide in *Clostridium thermoautotrophicum*. *J Bacteriol* 151:507–509
8. Diekert GB, Thauer RK (1980) The effect of nickel on carbon monoxide dehydrogenase formation in *Clostridium thermoaceticum* and *Clostridium formicoaceticum*. *FEMS Microbiol Lett* 7:187–189
9. Drake HL, Kusel K, Matthies C (2006) Acetogenic prokaryotes. In: Dworkin M, Falkow S, Rosenberg E, Schleifer K-H, Stachebrandt E (eds) *The prokaryotes*, 3rd edn. Springer, Berlin Heidelberg New York, pp 354–420
10. Drennan CL, Doukov TI, Ragsdale SW (2004) The metalloclusters of carbon monoxide dehydrogenase/acetyl-CoA synthase: a story in pictures. *J Biol Inorg Chem* 9:511–515
11. Hill J, Nelson E, Tilman D, Polasky S, Tiffany D (2006) Environmental, economic, and energetic costs and benefits of biodiesel and ethanol biofuels. *Proc Natl Acad Sci* 103:11206–11210
12. Huhnke R, Lewis R, Tanner RS (2008) Isolation and characterization of novel clostridial species. US patent application. Publication no. US 2008/0057554 A1
13. Ismaiel AA, Zhu CX, Colby GD, Chen JS (1993) Purification and characterization of a primary-secondary alcohol dehydrogenase from two strains of *Clostridium beijerinckii*. *J Bacteriol* 175:5097–5105
14. Koesnandar NN, Nagai S (1991) Effects of trace metal ions on the growth, homoacetogenesis and corrinoid production by *Clostridium aceticum*. *J Ferment Bioeng* 71:181–185
15. Korkhin Y, Kalb (Gilboa) AJ, Peretz M, Bogin O, Burstein Y, Frolow F (1998) NADP-dependent bacterial alcohol dehydrogenases: crystal structure, cofactor-binding and cofactor specificity of the ADHs of *Clostridium beijerinckii* and *Thermoanaerobacter brockii*. *J Mol Biol* 278:967–981
16. Liu C-L, Mortenson LE (1984) Formate dehydrogenase of *Clostridium pasteurianum*. *J Bacteriol* 159:375–380
17. Margeot A, Hahn-Hagerdal B, Edlund M, Slade R, Monot F (2009) New improvements for lignocellulosic ethanol. *Curr Opin Biotechnol* 20:372–380
18. Mehta MD, Saxena J, Tanner RS (2004) Enzyme activities in clostridia producing ethanol from carbon monoxide. *Abstr 104th Annu Meet Am Soc Microbiol*, O-81, p 474
19. Mielenz JR (2001) Ethanol production from biomass: technology and commercialization status. *Curr Opin Microbiol* 4:324–329
20. Peckman KR (1976) Investigation of the phylogenetic relationship of *Sporomusa ureae* to members of the *Bacillaceae* using primary structural characterization of 16S ribosomal ribonucleic acids. PhD thesis, University of Illinois, Urbana
21. Perlack RD, Wright LL, Turhollow AF, Graham RL, Stokes BJ, Erbach DC (2005) Biomass as feedstock for a bioenergy and bioproducts industry: the technical feasibility of a billion-ton annual supply. DOE/GO-102005-2135, Oak Ridge National Laboratory, Oak Ridge. <http://www.osti.gov/bridge>
22. Peters JW, Lanzilotta WN, Lemon BJ, Seefeldt LC (1998) X-ray crystal structure of the Fe-only hydrogenase (CpI) from *Clostridium pasteurianum* to 1.8 angstrom resolution. *Science* 5395:1853–1858
23. Pierce E, Xie G, Barabote RD, Saunders E, Han CS, Detter JC, Richardson P, Brettin TS, Das A, Ljungdahl LG, Ragsdale SW (2008) The complete genome sequence of *Moorella thermoacetica* (f. *Clostridium thermoaceticum*). *Appl Environ Microbiol* 10:2550–2573

24. Pimentel D (2003) Ethanol fuels: energy balance, economics, and environment impacts are negative. *Nat Resour Res* 12:127–134
25. Ragsdale SW, Ljungdahl LG (1984) Hydrogenase from *Acetobacterium woodii*. *Arch Microbiol* 139:361–365
26. Scopes RK (1983) An iron-activated alcohol dehydrogenase. *FEBS Lett* 156:303–306
27. Seravalli J, Xiao Y, Gu W, Cramer SP, Antholine WE, Krymov V, Gerfen GJ, Ragsdale SW (2004) Evidence that NiNi acetyl-CoA synthase is active and that the CuNi enzyme is not. *Biochem* 43:3944–3955
28. Smith PK, Krohn RI, Hermanson GT, Mallia AK, Garter FH, Provenzano MD, Fujimoto EK, Goeke MN, Olson BJ, Klenk DC (1985) Measurement of protein using bicinchoninic acid. *Anal Biochem* 150:76–85
29. Tanner RS (2007) Cultivation of bacteria and fungi. In: Hurst CJ, Crawford RL, Mills AL, Garland JL, Stettenbach LD, Lipson DA (eds) *Manual of environmental microbiology*, 3rd edn. ASM Press, Washington, DC, pp 69–78
30. Tanner RS (2008) Production of ethanol from synthesis gas. In: Wall J, Harwood CJ, Demain AL (eds) *Bioenergy*. ASM Press, Washington, DC, pp 147–151
31. Wagner R, Andreesen JR (1987) Accumulation and incorporation of ^{185}W -tungsten into proteins of *Clostridium acidiurici* and *Clostridium cylindrosporum*. *Arch Microbiol* 147:295–299
32. Yamamoto I, Saiki T, Liu SM, Ljungdahl LG (1983) Purification and properties of NADP-dependent formate dehydrogenase from *Clostridium thermoaceticum*, a tungsten-selenium-iron protein. *J Biol Chem* 258:1826–1832