Nature of Butyrate Oxidation by Mycoplasma hominis

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Abstract

VANDEMARK, P. J. (Cornell University, Ithaca, N.Y.), AND P. F. SMITH. Nature of butyrate oxidation by *Mycoplasma hominis*. J. Bacteriol. **89**:373–377. 1965.—Cell-free extracts of butyrate - grown *Mycoplasma hominis* strain O7, though lacking thickinase activity on butyric acid, were found to activate butyrate via an acetyl-butyric thiophorase. These extracts also contained an aceto-coenzyme A (CoA) kinase, a butyryl-CoA dehydrogenase, a crotonase, a reduced nicotinamide adenine dinucleotide-specific β -hydroxybutyryl-CoA dehydrogenase, and a thiolase. Thiolase activity was stimulated by the addition of magnesium ions. The presence of these enzyme activities in this *Mycoplasma* species supports the hypothesis that a fatty acid oxidation represents an energy source for the nonfermentative pleuropneumonia-like organisms.

Recently, workers investigating the problem of Mycoplasma in tissue culture have observed the disappearance of arginine from tissue cultures with these microorganisms contaminated (Schimke and Barile, 1963). On the basis of such observations, these investigators have postulated that the dissimilation of arginine, via the arginine dihydrolase pathway, may represent the main energy source for the nonfermentative Mycoplasma. The existence of the arginine dihydrolase pathway in the nonfermentative Mycoplasma was demonstrated previously by Smith (1960). However, in this latter study it was observed that ornithine accumulated, and its accumulation resulted in the inhibition or repression of further arginine breakdown. In view of these findings it would seem plausible that there exist one or more alternate energyyielding mechanisms in the nonfermentative Mycoplasma.

Lynn (1960) demonstrated that the nonfermentative Mycoplasma oxidize short-chain fatty acids and alcohols. The oxidative dissimilation of such substrates when coupled with a hemeterminated respiratory chain, such as previously reported in the nonfermentative Mycoplasma(VanDemark and Smith, 1964*a*) and involving oxidative phosphorylation, would seem to represent a more efficient energy source for these

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strains than the substrate phosphorylation available from arginine breakdown.

If the oxidation of a fatty acid represents an important metabolic pathway and energy source for the nonfermentative Mycoplasma, one should be able to demonstrate the enzymes involved in fatty acid oxidation including (i) a system of fatty acid activation, (ii) an acyl dehydrogenase, (iii) an enoly hydrase or crotonase, (iv) a β -hydroxyacyl dehydrogenase, and (v) a thiolase, as well as a tricarboxylic acid cycle. Evidence for the latter cycle in the M. hominis strain O7 was previously reported (VanDemark and Smith, 1964b). The present report gives evidence for the presence of those enzymes involved in butyrate oxidation in butyrate-grown cells of this nonfermentative strain.

MATERIALS AND METHODS

Culture. M. hominis strain O7 was grown and harvested from butyrate-containing (0.1%, w/v)PPLO broth (Difco), and resting-cell suspensions were prepared as previously described (VanDemark and Smith, 1964b).

Enzymatic methods. Cell-free extracts were prepared by sonic oscillation of resting-cell suspensions in a 10-kc Raytheon oscillator for 10 min. The cell debris was removed by centrifugation at $30,000 \times g$ for 30 min, and the resulting supernatant fraction was used for enzymatic analysis. The protein level of cell extracts was determined by the trichloroacetic acid method of Stadtman,

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Novelli, and Lipmann (1951) with bovine serum albumin as a standard.

Thiokinase activity was measured as the rate of hydroxamic acid formation by use of the method of Jones et al. (1953). Thiophorase activity was measured spectrophotometrically by the method of Stadtman (1953) as the decrease in optical density at $232 \,\mathrm{m}\mu$ resulting from the arsenolysis of acetyl coenzyme A (CoA) (formed when butyryl-CoA is incubated with acetate), the O7 enzyme preparation, and arsenate. Butyryl-CoA dehydrogenase was measured as the reduction of dichlorophenol-indophenol (DCPIP) at 600 mµ upon incubation of the enzyme with butyryl-CoA, DCPIP, and phenazine methosulfate, as described by Hauge (1956). Crotonase activity was based on the method of Stern (1955) as the decrease in absorption at 263 mµ upon incubation of the enzyme with crotonyl-CoA. β-Hydroxy-butyryl-CoA dehydrogenase was determined by following the oxidation of reduced nicotinamide adenine dinucleotide (NADH₂) with acetoacetyl-CoA and the sonic preparation. Thiolase activity was measured as the decrease in absorption at $305 \text{ m}\mu$ when acetoacetyl-CoA was incubated with the Mycoplasma enzyme.

Materials. Butyryl-CoA and crotonyl-CoA were prepared from their respective acid anhydrides according to the method of Simon and Shemin (1953). Acetoacetyl-CoA was prepared from diketene according to the method of Wieland and Rueff (1953). The diketene was purchased from the K & K Laboratories, Plainview, N.J., and was purified by distillation immediately prior to its use. NADH₂, reduced nicotinamide adenine dinucleotide phosphate (NADPH₂), CoA, and phenazine methosulfate were purchased from the Sigma Chemical Co., St. Louis, Mo. All other chemicals were commercial

 TABLE 1. Thiokinase activity of butyrate-grown Mycoplasma hominis 07

Fatty acid	Hydroxamic acid* (µmoles per hr per mg of enzyme protein)
Formate	< 0.01
Acetate	1.97
Propionate	0.65
Butyrate	0.06
β-Hydroxybutyrate	0.03
Isobutyrate	0.01
Acetoacetate	0.12
Valerate	0.05
Isovalerate	<0.01

* Hydroxamic acid formation was determined by the method of Jones et al. (1953) upon incubation of 20 μ moles of the sodium salt of fatty acid, 60 μ moles of tris(hydroxymethyl)aminomethane buffer (pH 7.4), 20 μ moles of ATP, 10 μ moles of MgCl₂, 20 μ moles of glutathione, 0.13 μ mole of CoA, 600 μ moles of hydroxylamine with 2 mg of enzyme protein.



FIG. 1. Evidence for an acetyl-butyric thiophorase in Mycoplasma hominis O7. The cuvettes contained 100 μ moles of histidine buffer (pH 6.8), 0.5 mg of enzyme protein, 0.5 μ mole of butyryl-CoA, and, where indicated, 20 μ moles of potassium acetate or butyrate.



FIG. 2. Evidence for an acetyl-butyric thiophorase and butyryl-CoA dehydrogenase in the oxidation of butyrate by extracts of Mycoplasma hominis O7. The cuvettes contained 67 µmoles of Tris buffer (pH 7.3), 1 mg of enzyme protein, 2 mg of phenazine methosulfate, 0.05 mg of DCPIP, 0.1 mg of COA, 10 µmoles of ATP, and, where indicated, 20 µmoles of potassium butyrate, 20 µmoles of dilithium acetyl phosphate, 2 µmoles of butyryl-CoA, and 5 units of transacetylase.

preparations of reagent grade. Transacetylase was prepared from a laboratory culture of *Escherichia coli* according to the method of Ochoa, Stern, and Schneider (1951).

Results and Discussion

Nature of butyrate activation. Previous studies of fatty acid activation by acetate-grown M. hominis O7 indicated that the kinase activity of these cells was relatively specific for acetate, with negligible activation of the other fatty acids tested (VanDemark and Smith, 1964b). Butyrate-grown cells showed a similar pattern of fatty acid activation, with thiokinase activity limited to acetate and propionate (Table 1). In the absence of a specific kinase for butyrate, it would seem plausible that butyrate activation is accomplished by means of an acetyl-butyric thiophorase analogous to that reported in Clostridium kluyveri by Stadtman (1953). Butyrate activation via thiophorase activity in the Mycoplasma enzyme preparation is illustrated in Fig.



FIG. 3. Crotonase activity of cell-free extracts of Mycoplasma hominis O7. The assay is based on the method of Stern (1955), with the cuvettes containing 400 μ moles of Tris buffer (pH 7.4), 0.4 mg of egg albumin, 6 μ moles of potassium ethylenediamine-tetraacetate (pH 7.4), 0.002 mg of enzyme protein, with and without 0.3 μ mole of crotonyl-CoA. The blank cuvette contained 0.3 mg of adenylic acid in place of crotonyl-CoA.



FIG. 4. Evidence for a NADH₂-linked β -hydroxybutyric dehydrogenase and its stimulation by Mg^{++} in Mycoplasma hominis O7. Cuvettes contained 67 μ moles of Tris buffer (pH 7.4), 0.5 mg of enzyme protein, 0.2 μ mole of acetoacetyl-CoA, and where indicated 1 mg of NADH₂ or NADPH₂, and 5 μ moles of $MgCl_2$. The blank cuvette contained all components except acetoacetyl-CoA.

1. The decrease in optical density at 232 m μ results from the loss of thiol ester absorption due to the arsenolysis of the acetyl-CoA formed via the transfer of CoA from butyryl-CoA to acetate by the thiophorase of the *M. hominis* O7 enzyme preparation.

Because of the high acetyl-CoA deacylase activity in these enzyme extracts (VanDemark and Smith, 1964b), the reverse reaction, starting with acetyl-CoA and butyrate, could not be demonstrated. For this reason, no attempt has been made at this time to demonstrate the substrate specificity of this transphorase.

Butyryl-CoA dehydrogenase and crotonase activity. Butyryl-CoA dehydrogenase activity, measured as the rate of DCPIP reduction, is illustrated in the upper curve of Fig. 2. These data indicate that dye reduction occurs when butyryl-CoA is incubated with the enzyme preparation and DCPIP. Negligible dye reduction occurs when butyrate, CoA, and adenosine triphosphate (ATP) are incubated with the M. hominis O7 enzyme (Fig. 2). Butyrate serves as a hydrogen donor for dye reduction by this organism only upon the addition of acetyl phosphate, transacetylase, CoA, and ATP, confirming the previous observations that prior to the oxidation of butyryl-CoA, butyrate is not activated via a kinase-type enzyme but by means of a CoA transphorase.

The crotonyl-CoA formed as a result of this oxidation is hydrated to β -hydroxybutyryl-CoA by the action of a crotonase present in this *Mycoplasma* enzyme preparation (Fig. 3). The incubation of the sonic enzyme preparation with crotonyl-CoA results in a rapid decrease in absorption at 263 m μ due to the hydration of the substrate to form β -hydroxybutyryl-CoA (Fig. 3).

 β -Hydroxybutyryl-CoA dehydrogenase and thiolase activities. Evidence for a β -hydroxybutyryl-CoA dehydrogenase is shown in Fig. 4, in which the reduction of acetoacetyl-CoA is measured as rate of NADH₂ oxidation at 340 m μ when acetoacetyl-CoA and NADH₂ are incubated



FIG. 5. Thiolase and β -hydroxybutyric dehydrogenase activity of sonic extracts of Mycoplasma hominis O7. Cuvettes contained 67 µmoles of Tris buffer (pH 7.4), 5 µmoles of MgCl₂, 0.1 mg of enzyme protein, approximately 0.2 µmole of acetoacetyl-CoA, and 1 mg of NADH₂ where indicated. The reaction was followed as the decrease in optical density at $305 \text{ m}\mu$ due to the disappearance of acetoacetyl-CoA. The blank cuvette contained all the reaction components except acetoacetyl-CoA.



FIG. 6. Postulated cycle of butyrate oxidation in Mycoplasma hominis 07.

with the enzyme preparation. The reaction is specific for $NADH_2$ and is markedly stimulated by the addition of Mg ions.

Thiolase activity is illustrated in Fig. 5. The initial portion of the plot represents the decrease in the thiol ester absorption at 305 m μ due to the enzymatic cleavage of acetoacetyl-CoA to yield two molecules of acetyl-CoA. The addition of NADH₂ to the reaction mixture results in an increase in the rate of acetoacetyl-CoA disappearance due to the additive action of the β -hydroxybutyryl-CoA dehydrogenase and thiolase activities present in the crude sonic preparations.

The postulated pathway of butyrate oxidation based on this demonstration of the specific enzymes as described in this paper is illustrated in Fig. 6. In this cyclic mechanism, the resultant acetyl-CoA serves either as the CoA donor for the activation of butyrate by transphorase, or it enters the tricarboxylic and glyoxylic acid cycles which have previously been described in this strain (VanDemark and Smith, 1964b).

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