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Articles

Nitrite, a New Substrate for Nitrogenase[†]

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Received April 1, 1988; Revised Manuscript Received August 24, 1988

ABSTRACT: We have examined the reactivity of the purified component proteins of Azotobacter vinelandii nitrogenase (Av1 and Av2) toward nitrate and nitrite. Nitrate has no effect on H_2 evolution or C_2H_2 reduction by nitrogenase and thus is neither a substrate nor an inhibitor. Nitrite dramatically inhibits H_2 evolution. This inhibition has two components, one irreversible and one reversible upon addition of CO. The irreversible inhibition is due to nitrite inactivation of the Fe protein. The rate of this inactivation is greatly enhanced by addition of MgATP, suggesting the [4Fe-4S] cluster is the site of nitrite attack. The reversible inhibition does not represent an inhibition of electron flow but rather a diversion of electrons away from H_2 evolution and into the six-electron reduction of nitrite to ammonia. Thus, nitrogenase functions as a nitrite reductase.

ritrogenase is composed of two easily separated component proteins, called the molybdenum—iron protein (MoFe protein) and the iron protein (Fe protein). The physical properties of these two proteins have been recently reviewed (Orme-Johnson, 1985), and great similarity among proteins from different bacterial sources is evident (Emerich & Burris, 1976a,b). Nitrogen fixation and all other reductions catalyzed by nitrogenase require both component proteins, a source of reducing

equivalents, MgATP, protons, and an anaerobic environment (Bulen & LeComte, 1966). The MoFe protein is believed to contain the site of substrate reduction (Shah et al., 1973; Hageman & Burris, 1978; Hawkes et al., 1984), while the Fe protein is generally accepted as the specific one-electron donor for the MoFe protein (Hageman & Burris, 1978a,b; Thorneley & Lowe, 1985). In addition to N_2 , nitrogenase catalyzes the reduction of protons and a number of apparently nonphysiological substrates, many of which contain triple bonds (Burgess, 1985). Such alternative substrates have been studied as probes for the number and nature of sites of substrate interaction on nitrogenase and the types of intermediates that might be formed during N_2 reduction. This paper will describe recent studies on the reactivity of nitrogenase toward two N-O bond

[†]This study was supported by Grant 85-CRCR-1-1635 from the U.S. Department of Agriculture.

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[‡]Supported by a National Institutes of Health Pre-Doctoral Training Grant and a Corporate University Partnership Fellowship.

containing molecules, nitrate and nitrite.

MATERIALS AND METHODS

Reagents and Chemicals. ATP, creatine phosphokinase, creatine phosphate, Tes, NaCl, iodine (crystalline), soluble potato starch, NaNO₂, and NaNO₃ were from Sigma Chemical Co. The specialty gases 99.997% Ar, 99.997% CO, and 1.05% H₂ in Ar (gravimetric standard) and 10% C₂H₂ in Ar were provided by Liquid Carbonic. Purified NO was kindly provided by Dr. William Evans from the UC Irvine Chemistry Department. The 1000 ppm C₂H₄ standard was provided by Alltech. Na₂S₂O₄ was from E-M Chemicals (a division of E. Merck).

Nitrogenase Assay. Azotobacter vinelandii MoFe and Fe proteins, designated Av1 and Av2, respectively, were purified and analyzed as described elsewhere (Burgess et al., 1980). Specific activities of the proteins were ca. 1800 nmol of H₂ \min^{-1} (mg of Av2)⁻¹ and 2800 nmol of H₂ \min^{-1} (mg of Av1)⁻¹. Unless otherwise indicated, all assays were performed at 30 °C in 9.5-mL calibrated vials, fitted with butyl rubber serum caps, containing the appropriate gas mixture. The 1.0-mL reaction mixture contained 38 mM Tes-KOH (pH 7.4), 2.5 mM ATP, 5.0 mM MgCl₂, 30 mM creatine phosphate, 20 mM neutralized Na₂S₂O₄ (except for dithionite utilization studies see below), and 2.5 units of creatine phosphokinase. The vessel containing the reaction mixture was degassed and filled with the appropriate gas mixture by using a published procedure (Corbin, 1978), Na₂S₂O₄ was added, Av1 was added, and the mixture was incubated at 30 °C for 5 min. Then, NaNO₃, NaNO₂, or NaCl was added when appropriate, and the reaction was started 10 s later by adding Av2 to give 1 mg/mL total protein at the Av2/Av1 molar ratio of 5. For assays involving salt addition, a 1.2 M stock solution in 0.1 M Tes-KOH, pH 7.4, was used. Because nitrite causes irreversible inactivation of the Fe protein (see Results), the reaction is not linear with time. Consequently, assays were run for the shortest practical time period for the measurements in question. The exact times for each assay are found in the figure legends. The assays were run with shaking at 30 °C and were quenched at the appropriate time with 0.4 M EDTA, pH 7.4 (0.2 mL), unless dithionite was to be determined (see below)

Product Analysis. With the exception of dithionite utilization studies (see below), all products were measured on the same reaction vial. All data represent means of, typically, triplicate determinations with standard deviations less than 10%. Gas samples, 200 μ L at bottle pressure, were taken with a pressure-lock syringe (Precision Sampling, Baton Rouge, LA) and were analyzed with a Varian 3700 gas chromatograph with a Poropack N columm (He) and a flame ionization detector for C_2H_4 analysis and a thermal conductivity detector with a molecular sieve 13X column (Ar) for H_2 and NO analysis. Ammonia was determined by an HPLC fluorescence method described elsewhere (Corbin, 1984). Data calculations for H_2 , NH_3 , and C_2H_4 were performed as described previously (Rubinson et al., 1985; Robinson et al., 1986).

For dithionite utilization experiments H_2 and $S_2O_4^{2-}$ utilization were measured on the same reaction vial. These assay vials contained 3 mM $Na_2S_2O_4$. The assays were performed and data analyzed as described by Li et al. (1982) with the following modifications. The reaction vials were quenched by

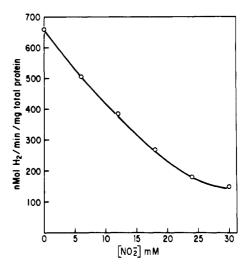


FIGURE 1: Nitrite inhibits H_2 evolution by nitrogenase. Plot of nmol of H_2 formed min⁻¹ (mg of total protein)⁻¹ for 5-min assay under Ar. All assays contained 1 mg/mL total protein at an Av2/Av1 molar ratio of 5. For all data points, total salt concentration was adjusted to 30 mM by addition of NaCl.

rapid chilling in an ice-ethanol bath. This procedure was added because the dithionite-HCHO adduct is somewhat unstable toward oxidation by NO₂⁻. (Dithionite itself was insensitive to oxidation by NO₂⁻ over the time course of our assays.) Shortly after chilling, 0.1 mL of 40% HCHO in water was added to the vials followed by 0.1 mL of a 2% soluble potato starch solution and 0.1 mL of a 2.0 M sodium acetate solution in 0.1 M Tris, pH 7.4. This procedure differs from that of Li et al. (1982), which used a 2.0 M sodium acetate solution at pH 4.0. This modification was necessary to prevent the oxidation of the dithionite-HCHO adduct by HNO₂, which is a much stronger oxidizing agent than NO₂⁻.

RESULTS

Reactivity of Nitrogenase toward Nitrate. H_2 evolution and C_2H_2 reduction by nitrogenase were monitored over the concentration range 0–30 mM nitrate. Total salt concentration was kept at 30 mM for all vials by addition of Cl^- . Both activities were constant regardless of NO_3^- concentration, giving values of 531 ± 12 nmol of $H_2 \min^{-1}$ (mg of total protein)⁻¹ and 433 ± 19 nmol of $C_2H_2 \min^{-1}$ (mg of total protein)⁻¹. Thus, the N–O bond containing molecule NO_3^- does not appear to be either a substrate or an inhibitor of the enzyme.

Nitrite Is an Inhibitor. Figure 1 demonstrates that, in the same concentration range where nitrate has no effect on nitrogenase, nitrite is an inhibitor of H₂ evolution under Ar.

Nitrite Inhibition Has both CO-Irreversible and CO-Reversible Components. The nitrite inhibition shown in Figure 1 could have been due to (a) irreversible inactivation of Av1, Av2, or some other component of the assay system, (b) reversible inhibition of the total electron flow of the type caused by CN⁻ (Li et al., 1982) or CH₃NC (Rubinson et al., 1983), or (c) a redirection of electrons away from H₂ evolution and into some NO₂⁻ reduction product that was not being measured. CO is known to relieve CN⁻-type inhibition and to inhibit the reduction of all known nitrogenase substrates except protons (Burgess, 1985). Thus, in order to distinguish irreversible from reversible inhibition, we have monitored the effects of CO on this NO₂⁻ inhibition.

Figure 2, panel A, shows that 60 mM NO_2^- causes $\sim 85\%$ inhibition of H_2 evolution under an atmosphere of 100% Ar but only 45% inhibition under an atmosphere of 5% CO/95%

¹ Abbreviations: ATP, adenosine 5'-triphosphate; Tes, 2-[[tris(hydroxymethyl)methyl]amino]ethanesulfonic acid; HPLC, high-performance liquid chromatography.

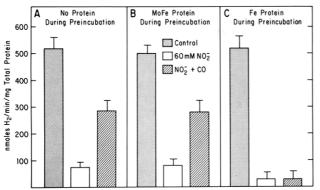


FIGURE 2: Nitrite inhibition has both irreversible and reversible components. Control vials show nmol of H_2 min⁻¹ (mg of total protein)⁻¹ under an atmosphere of 100% Ar and 5% CO/95% Ar (three vials each) for a 2-min assay with 60 mM NaCl present during both the assay and a 5-min preincubation period. Open boxes are identical conditions except that 60 mM NaNO2 replaces NaCl and the atmosphere throughout is 100% Ar. The diagonally striped boxes are identical with open boxes except that the atmosphere throughout is 5% CO/95% Ar. Preincubations were as described under Materials and Methods except that the salt was added at the beginning of the preincubation period and the proteins were added as follows. In panel A neither component protein was present during preincubation, and the assay was started by adding both components. In panel B only the Av1 was present, and during preincubation the assay was started by adding Av2, while in panel C only Av2 was present during preincubation and the assay was started by adding Av1. All assays contained 1 mg/mL total protein at an Av2/Av1 molar ratio of 5.

Ar. Thus, CO is able to partially protect the enzyme from NO_2^- inhibition. The simplest explanation for this result is that NO_2^- inhibition has two forms, one CO irreversible and one reversible by CO.

CO-Irreversible NO_2^- Inhibition Is Due to Inactivation of the Fe Protein. Figure 2, panel B, shows that the extent of NO_2^- inhibition under Ar and the ability of CO to protect the H_2 evolution reaction from NO_2^- inhibition are unaffected by preincubation of Av1 with NO_2^- relative to preincubation with no protein present (panel A). Panel C, however, shows that a 5-min preincubation of 60 mM NO_2^- with Av2 causes almost complete inhibition of the enzyme in the presence or absence of CO such that once the 2-min assay is started almost no product is formed. These data indicate that NO_2^- is causing CO-irreversible inhibition of the Fe protein component of nitrogenase.

For the experiments of Figure 2, MgATP was also present during the preincubation period. The binding of MgATP to the Fe protein is known to have a significant influence on the structure of the protein, apparently causing its previously protected [4Fe-4S] cluster to be exposed to the solvent (Mortenson et al., 1976; Ljones & Burris, 1978). The time dependence of the inhibition reaction shown in Figure 3 is characteristic of an irreversible inactivation reaction. (In control experiments we observed no time dependence for preincubation of the reaction mixture alone or of the MoFe protein alone.) Figure 3 shows that the rate of this Fe protein inactivation by NO₂⁻ is dramatically increased by having MgATP present. These data indicate that the site of NO₂⁻ inactivation is likely to be on the [4Fe-4S] cluster.

Inactivation of the Fe protein by NO₂⁻ was previously recognized by Meyer (1981), who also concluded that the [4Fe-4S] cluster was the likely site of NO₂⁻ attack. He suggested, however, that the inactivation of the Fe protein was not actually caused by nitrite but rather by NO which was produced nonenzymatically in the assay system. Under the conditions for the detection of H₂, our gas chromatograph registers NO eluting from the molecular sieve at 0.61 min, well

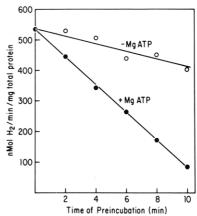


FIGURE 3: Rate of Fe protein inactivation by nitrite is enhanced by addition of MgATP. A 4.83 mg/mL Fe protein solution was preincubated for the time periods indicated with 60 mM NaNO₂ with (•) or without (O) 2.5 mM MgATP. At the end of the preincubation period, an aliquot of the Fe protein solution was used to initiate a standard H₂ evolution under Ar nitrogenase assay. The final [NO₂⁻] in the actual assay vial was 7.2 mM, and the assay was run for 2 min. The assay contained 1 mg/mL total protein at an Av2/Av1 molar ratio of 5

Table I: Effect of Added NO on H ₂ Evolution		
gas-phase [NO] (atm) ^a	H ₂ evolution ^b	5% CO present
0	548 ± 53	
6.7×10^{-4}	540 ± 9	, -
6.7×10^{-4}	545 ± 23	+
2.0×10^{-2}	291 ± 38	_
2.0×10^{-2}	295 ± 24	+

^aTo obtain a concentration of 6.7 × 10⁻⁴ atm, 10 μ L of NO was added to a 15-mL assay vial filled with Ar. This corresponds to 30 μ M NO in the gas phase or about 1.4 μ M NO in the 1-mL liquid phase after correction for solubility. To obtain a concentration of 2.0 × 10⁻² atm, 300 μ L of NO was added to a 15-mL assay vial filled with Ar. This corresponds to 892 μ M NO in the gas phase or 41 μ M in the liquid phase after correction for solubility. ^bIn nmol of H₂ min⁻¹ (mg of total protein)⁻¹.

separated from H_2 at 0.35 min, N_2 at 0.48 min, O_2 at 0.52 min, and CO at 0.85 min. No peaks were detected at 0.61 min during measurement of H_2 evolution for the experiments in Figures 2 and 3 or for other standard NO_2^- assays. Thus, if NO was being produced nonenzymatically in our vials, it was being produced at levels below our threshold of detectability.

To test if such low NO concentrations could be responsible for the inhibition of H₂ evolution we observed after addition of NO₂, we monitored H₂ evolution by nitrogenase following the addition of easily detectable concentrations of NO. These data are shown in Table I. The NO concentration of $6.7 \times$ 10⁻⁴ atm was 10-fold above our detection limits and was easily detected in the assay vials during H₂ evolution measurements made after the assays were run. The data in Table I show that this level of NO had no effect on H2 evolution under Ar. We therefore conclude that the inhibition illustrated in Figures 2 and 3 could not have been caused by even lower levels of nonenzymatically produced NO and was being caused instead by NO₂ itself. It should be noted that we did observe significant inhibition of H2 evolution by NO at higher concentrations (Table I), which is in agreement with previous reports of NO inhibition (Rivera-Ortiz & Burris, 1975; Meyer, 1981; Trinchant & Rigaud, 1982). The inhibition of H₂ evolution observed at the higher concentrations of NO was not reversed by the addition of CO (Table I).

Nitrite Is a New Substrate for Nitrogenase: Electron Balance Studies. The CO reversible NO₂⁻ inhibition of H₂

FIGURE 4: Nitrite reduction by nitrogenase. (Left) $\rm H_2$ evolution and $\rm S_2O_4^-$ utilization data for 30 mM $\rm NO_2^-$ under CO. Assays were run for 8 min. (Right) $\rm H_2$ evolution, $\rm NH_3$ production, and $\rm S_2O_4^{2-}$ utilization data for nitrogenase in the presence of 30 mM $\rm NO_2^-$ under Ar. Controls without either the Fe or the MoFe protein under Ar gave no $\rm H_2$ or $\rm NH_3$. Controls without $\rm NO_2^-$ (replaced with Cl⁻) under Ar also gave no $\rm NH_3$. The assays contained 1 mg/mL total protein at an $\rm Av2/Av1$ molar ratio of 5.

evolution shown in Figure 2 could be the result of NO₂ inhibiting total electron flow through the enzyme in the same way that CN⁻ inhibits electron flow (Li et al., 1982). Alternatively, NO₂ could be a substrate for the enzyme, redirecting electrons away from H₂ evolution and into its reduction. To distinguish these possibilities, we have directly measured the rate of electron flow by monitoring dithionite utilization. Figure 4 shows that under an atmosphere of CO with 30 mM nitrite present the rate of electron flow into nitrogenase (dithionite utilization) equals the rate of electron flow out of nitrogenase as H₂ gas. Figure 4 also demonstrates that under an atmosphere of Ar the rate of electron flow into nitrogenase in the presence of 30 mM nitrite is identical with the rate observed under CO. It is only the rate of H₂ evolution that is dramatically decreased. The finding of identical dithionite utilization rates under Ar and CO in the presence of 30 mM $NO_2^- \, demonstrates \, that \, NO_2^- \, is \, not \, a \, \, CN^- \! - \! type \, inhibitor \, of$ total electron flow.

In order to account for the difference in electron flow into nitrogenase as dithionite and out of the enzyme as H₂ gas (Figure 4, right panel), we searched for possible products of NO₂⁻ reduction. Figure 4 shows that ammonia is a major product of this reaction. The NH₃ is not formed in the absence of added NO₂⁻, or if S₂O₄², MgATP, or either of the two component proteins is left out of the assay. The formation of NH₃, like the reduction of all N₂ase substrates, is inhibited by CO (Figure 4). The reduction of NO₂⁻ to NH₃ requires six electron. Figure 4 shows that the sum of electron pairs that flow out of nitrogenase as H₂ and NH₃ is equal, within the limits of the measurements, to the amount of dithionite consumed. Thus, nitrogenase functions as a nitrite reductase.

DISCUSSION

Of the N-O bond containing molecules that might be expected to react with nitrogenase only N₂O has been studied in any detail. N₂O was first implicated as a nitrogenase substrate from in vivo studies which showed that it inhibited N₂ reduction (Hoch et al., 1960; Mozen & Burris, 1954). Subsequent in vitro experiments, using Av and Clostridium pasteurianum cell-free extracts, established N₂O as an alternative nitrogenase substrate by demonstrating that the reaction required reductant and MgATP and was inhibited by CO (Lockshin & Burris, 1965; Hardy & Knight, 1966). More recently, Jensen and Burris (1986) have studied N₂O reduction using purified nitrogenase and have shown that N₂O

is a competitive inhibitor of N_2 reduction and that it is not an inhibitor of electron flow. As N_2 is the major product of N_2O reduction, the reaction presumably involves cleavage of the N-O bond to yield N_2 and H_2O and suggests that the single N-O bonded resonance form $O^{-1}N \equiv N$ is the most appropriate representation of the substrate.

Interaction with Nitrate. The observation that nitrogenase appears to be reactive toward the N–O single bond in N_2O led us to consider the enzyme's reactivity toward the other N–O single bond containing molecules NO_3^- and NO_2^- . The interaction of purified nitrogenase with NO_3^- had not been previously investigated in spite of the fact that NO_3^- is reduced by two electrons and two protons to NO_2^- and H_2O by another (albeit dissimilar) Mo-containing enzyme, nitrate reductase. The data presented here show that NO_3^- has no effect on H_2 evolution or C_2H_2 reduction by nitrogenase and thus is neither a substrate nor an inhibitor of the enzyme.

Irreversible Inactivation of the Fe Protein by NO_2^- . The interaction of NO₂ with nitrogenase has received more attention because this naturally occurring highly reactive anion was thought to have a possible role in vivo (Trinchant & Rigaud, 1980, 1982; Meyer, 1981). The data presented here clearly show that NO2- has two very different effects on nitrogenase, one CO irreversible and the other reversible by CO. The irreversible effect was previously recognized by Meyer (1981), who concluded that NO₂⁻ irreversibly inactivated the Fe protein component of nitrogenase probably by interaction with its [4Fe-4S] center. The data shown in Figures 2 and 3 support these conclusions of Meyer (1981) and are in conflict with the prior report by Trinchant and Rigaud (1980) that NO₂⁻ had no effect on the Fe protein component. Meyer attributed the irreversible effect of NO₂not to that anion specifically but rather to NO which was produced nonenzymatically in the assay by reaction of NO₂ with dithionite. Because we were unable to detect NO in our assay vials (they were run for much shorter time periods) and because NO did not inhibit H₂ evolution in our assays when we added it to 10-fold above detectable levels, we conclude that NO₂ itself is the molecule causing irreversible inactivation of the Fe protein. These data do not preclude the possibility, however, that the mechanism of inactivation involves the one-electron oxidation of the [4Fe-4S] cluster by NO₂⁻ to yield a nitrosyl adduct.

 NO_2^- Is a New Substrate for Nitrogenase. Trinchant and Rigaud (1980, 1982) reported that NO₂- was a reversible, competitive inhibitor of C₂H₂ reduction by partially purified Rhizobium nitrogenase although they concluded that it was not a substrate for the enzyme. That conclusion was based on their attempts to monitor the disappearance of NO2 during the course of an assay. This procedure may have been in error, however, because dithionite interferes with the NO₂⁻ colorimetric assay (Trinchant & Rigaud, 1982). As shown in Figure 4, we have now established that nitrite is a substrate for nitrogenase and is reduced by six electrons to ammonia. Unfortunately, the continuous inactivation of the Fe protein caused by NO₂- during the course of the assay prevents meaningful calculation of kinetic constants for this substrate. The data in Figure 4 do suggest that NO₂ has a fairly high $K_{\rm m}$ (on the order of 10-20 mM) when compared to other alternative substrates (Burgess, 1985). However, the p K_a of NO_2^- is ~3.8, and the possibility that HNO_2 could be the actual substrate (or irreversible inhibitor) cannot be eliminated at this time.

Nitrogenase Interaction with N-O Bond Containing Molecules: Mechanistic Considerations. In nature there are only

three known six-electron reduction reactions: the reduction of dinitrogen to ammonia catalyzed by nitrogenase, the reduction of sulfite to sulfide catalyzed by sulfite reductase, and the reduction of nitrite to ammonia catalyzed by assimilatory nitrite reductase (Guerrero et al., 1981). We now know that nitrogenase can catalyze two of these reactions. It is not known if it can also catalyze the reduction of sulfite. It is interesting to note, however, that sulfite is always present in nitrogenase assays as an oxidation product of dithionite and its reduction could pass unnoticed.

Currently there is no direct evidence to suggest how or where the N-O bond containing molecule N₂O binds to nitrogenase or to indicate the chemical mechanism of its reduction. On the basis of what is known about N₂O reduction (Jensen & Burris, 1985) and the apparently analogous reduction of N_3 (Rubinson et al., 1985), we suggest that nitrogenase binds N_2O and catalyzes its reduction as in eq 1. By binding to the active

$$E + {}^{-}O - {}^{+}N = N \rightarrow E - O - N = N: \xrightarrow{2e^{-}, 2H^{+}} E + H_{2}O + N_{2}$$
 (1)

site with its anionically polarized oxygen atom, it could simply serve as a scavenger of the two electrons and two protons to form H₂O, leaving N₂ as an easily lost fragment.

The six-electron reduction of NO₂⁻ to NH₃ by the sirohemeand [Fe-S]-containing enzyme assimilatory nitrite reductase does not yield any partially reduced products, and the nature of nitrite binding or intermediate formation is unknown (Guerrero et al., 1981). The bent molecule NO₂ can bind to metals in two distinct ways: the N-bound nitro form and the O-bound nitrito form (Burmeister, 1968). N-Bound forms are typically thermodynamically favored and in the present case may help explain why reduction proceeds all the way to NH₃. Electron balance studies in the present work (Figure 4) are consistent with the complete reduction of NO₂⁻ although they do not eliminate the possibility of small quantities of some lesser reduced product being formed. On the basis of what is known from other substrates, it is likely that the nitrogenase reaction involves a series of one- or two-electron reduction steps with enzyme-bound intermediates being formed along the way. A one-electron/two-proton reduction to give a nitrosyl intermediate is quite probably as shown in eq 2. The enzyme-

$$E - : N = 0$$
 $\frac{1e^{-}.2H^{+}}{}$ $E - N = 0 + H_{2}O$ (2)

bound nitrosyl intermediate would then go on to be reduced by five additional electrons and five protons to yield NH₃ + H₂O. Although we have not detected NO as a product of NO₂ reduction, future experiments will be directed toward identification of this possible enzyme-bound intermediate. If an enzyme-bound NO intermediate does occur during NO₂⁻ reduction, then it is probable that NO itself would be a substrate reducible to ammonia. The reactivity of nitrogenase toward NO has not been extensively studied, but it is interesting to note that like NO₂ it is known to have both irreversible and reversible inhibitory effects (Rivera-Ortiz & Burris, 1975; Meyer, 1981; Trinchant & Rigaud, 1982).

The other N-O bond containing molecule we have considered here is NO₃⁻. It is neither a substrate nor an inhibitor for the enzyme. There are two possible reasons why N_2O and NO₂⁻ are substrates while NO₃⁻ is not. First, N₂O and NO₂⁻ are softer, more polarizable ligands, and they may bind more readily to the active site. Second, NO₃⁻ is an anion of high thermodynamic stability, and there may simply be a much greater activation energy barrier to the reduction of its first nitrogen-oxygen bond.

Conclusions

In summary, our studies have shown the following: (1) NO₃⁻ is neither a substrate nor an inhibitor of nitrogenase, (2) NO₂ causes irreversible inactivation of the Fe protein probably by reaction with its [4Fe-4S] cluster, and (3) NO₂⁻ is a new substrate for nitrogenase and is reduced by six electrons to ammonia.

ACKNOWLEDGMENTS

We are pleased to acknowledge the talents of Charles Miller and Teresa Chun in cell growth and protein purification.

Registry No. NO₂⁻, 14797-65-0; NO₃⁻, 14797-55-8; NO, 10102-43-9; nitrogenase, 9013-04-1.

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Purification and Characterization of an Isoform of Protein Kinase C from Bovine Neutrophils[†]

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Received April 6, 1988; Revised Manuscript Received August 5, 1988

ABSTRACT: Protein kinase C (PKC) from bovine neutrophils was purified 1420-fold. Subcellular fractionation analysis of bovine neutrophil homogenate in the presence of EGTA indicated that more than 95% of the PKC activity was present in the soluble fraction. The purification procedure from cytosol involved sequential chromatographic steps on DE-52 cellulose, Mono Q, and phenyl-Sepharose. Whereas bovine brain PKC could be resolved into four isoenzymatic forms by chromatography on a hydroxylapatite column, bovine neutrophil PKC was eluted in a single peak, suggesting that it corresponded to a single isoform. The apparent molecular weight of bovine neutrophil PKC was 82 000, as judged by sodium dodecyl sulfate-polyacrylamide gel electrophoresis. By filtration on Sephadex G-150, a molecular weight of 85 000 was calculated, indicating that bovine neutrophil PKC in solution is monomeric. Its isoelectric point was 5.9 ± 0.1 . Bovine neutrophil PKC was autophosphorylated in the presence of $[\gamma^{-32}P]ATP$, provided that the medium was supplemented with Mg²⁺, Ca²⁺, phosphatidylserine, and diacylglycerol; phorbol myristate acetate could substitute for diacylglycerol. Autophosphorylated PKC could be cleaved by trypsin to generate two radiolabeled peptides of M, 48 000 and 39 000. The labeled amino acids were serine and threonine. During the course of the purification procedure of bovine neutrophil PKC, a protein of M_r 23 000, which was abundant in the cytosolic fraction of the homogenate, was found to exhibit a strong propensity to PKC-dependent phosphorylation in the presence of $[\gamma^{-32}P]ATP$, Mg^{2+} , Ca^{2+} , phosphatidylserine, and diacylglycerol. This protein was recovered together with PKC in one of the two active peaks eluted from the Mono Q column at the second step of PKC purification. It is suggested that the M_r 23 000 protein might be a natural substrate for bovine neutrophil PKC.

Protein kinase C, a Ca²⁺- and phospholipid-dependent kinase $(PKC)^1$ of $M_r \simeq 80\,000$, has been found in a large number of tissues and appears to play a central role in cellular economy [for review, see Nishizuka (1984)]. Recently, the existence of a family of PKC genes has been reported (Knopf et al., 1986; Parker et al., 1986; Coussens et al., 1986; Ono et al., 1986; Ohno et al., 1987). Three cDNA clones were obtained from rat, bovine, human, and rabbit libraries, and two sequences from rat brain were shown to derive from alternative splicing of a single clone (Kubo et al., 1987). Isoforms of PKC have been identified after separation on a hydroxylapatite (HTP) column from rat brain (Huang et al., 1986a; Kikkawa et al., 1987) and rabbit brain (Jaken & Kiley, 1987). PKC isoforms have also been reported in rat liver (Azhar et al., 1987), in murine fibroblasts (McCaffrey et al., 1987), in rat spleen (Brandt et al., 1987), and in bovine adrenocortical tissue (Pelosin et al., 1987).

It is a current view that PKC in neutrophils accomplishes a number of strategic functions, following exposure of these cells to specific stimuli or to phorbol myristate acetate (PMA), in particular the activation of the respiratory burst [for reviews, see Rossi (1986) and Tauber (1987)]. Our group is interested in the activation of the respiratory burst in bovine neutrophils (Ligeti et al., 1988). We therefore decided to explore in more detail the nature, properties, and function of PKC from bovine neutrophils. The present paper describes a simple and rapid method of purification of PKC from bovine neutrophils and some properties of this enzyme. The search for isoforms of neutrophil PKC is also reported, together with the demonstration of a PKC-dependent phosphorylation of a small mo-

[†]This work was supported by grants from the Centre National de la Recherche Scientifique (CNRS/U.A. 1130 alliée à l'INSERM) and from the Faculté de Médecine, Université Joseph Fourier de Grenoble.

¹ Abbreviations: PKC, protein kinase C; EDTA, ethylenediaminetetraacetic acid; EGTA, ethylene glycol bis(β -aminoethyl ether)-N,N,N',N'-tetraacetic acid; PMSF, phenylmethane sulfonyl fluoride; TPCK, N-tosyl-L-phenylalanine chloromethyl ketone; TLCK, N*-p-tosyl-L-lysine chloromethyl ketone; Mops, 3-(N-morpholino)propanesulfonic acid; Tris-HCl, tris(hydroxymethyl)aminomethane hydrochloride; PMA, 4 β -phorbol 12-myristate 13-acetate; ATP, adenosine 5'-triphosphate; HTP, hydroxylapatite; M_n, molecular weight; SDS, sodium dodecyl sulfate.