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Stabilization of the Quaternary Structure of Transcarboxylase by Cobalt(II) Ions[†]

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ABSTRACT: When dilute solutions of transcarboxylase are incubated at 25 °C in an alkaline 50 mM buffer, the enzyme rapidly loses activity. This loss of activity is accompanied by the dissociation to enzymatically inactive subunits. The inclusion of 2 mM Co²⁺ in the buffer reduces both dissociation and the loss of enzymatic activity. This stabilization does not take place with 2 mM Mg²⁺, Mn²⁺, Fe²⁺, Ni²⁺, Ca²⁺, or Cu²⁺, but there is a slight protection by Zn²⁺. At Co²⁺ concentrations of less than 2 mM, the stabilization decreases. The cobalt

involved in the stabilization is not that required for catalysis as evidenced by the fact that the "catalytic" cobalt does not exchange with added free Co²⁺ under the conditions that prevent loss of enzymatic activity. The stabilizing effects of Co²⁺ were also observed toward inactivation with guanidinium chloride and by heat. It is proposed that Co²⁺ shifts the equilibrium of the dissociation of transcarboxylase toward the associated form and thus enzymatic activity is retained at alkaline pH.

Transcarboxylase (methylmalonyl-CoA:pyruvate carboxyltransferase, EC 2.1.3.1) is a biotin-containing enzyme found in the propionic acid bacteria [for reviews, see Wood & Zwolinski (1976) and Wood (1979)]. An illustration of the quaternary structure of the enzyme and its dissociation to enzymatically inactive subunits is shown in Figure 1. By use of transcarboxylase isolated from cells grown in ⁶⁵Zn or ⁶⁰Co, it has been demonstrated that the outer 5S_E subunit contains Co²⁺ and Zn²⁺ (Northrop & Wood, 1969; Ahmad et al., 1972; Fung et al., 1974). These two metals are tightly bound and dissociate only at a very low pH or in the presence of sodium dodecyl sulfate (Ahmad et al., 1972). Even at pH 9, EDTA¹ fails to remove these metals. These metals will be referred to as the "catalytic" metals.

In this report, we show that the presence of 2 mM exogenously added Co²⁺ ions protects transcarboxylase against dissociation and loss of activity at pH 8 and 9. These Co²⁺ ions are distinct from the catalytic metals and the protective effect is specific for Co²⁺. Co²⁺ also protects against denaturation of the enzyme by guanidinium chloride and treatment at 50 °C.

Materials and Methods

The 26S transcarboxylase was isolated by the method of Wood et al. (1977) and assayed as described by Wood et al. (1969). For the tests of the effects of cobalt on enzymatic activity, the mixtures with and without Co²⁺ were incubated at 25 °C and exposed to air unless otherwise stated, under the

conditions described in the text and figure legends. Aliquots were removed at the stated times and assayed. Unless otherwise stated, the Co²⁺ concentration was 2 mM and the buffer was 50 mM Hepes (Na⁺), pH 8 or 9. Transcarboxylase containing ⁶⁰Co was isolated as described by Ahmad et al. (1972) and the 6S_E subunit by dissociation of 26S transcarboxylase at neutral pH, followed by glycerol gradient centrifugation (Wood et al., 1977). The 12S_H subunit was purified as recently described by Bahler et al. (1981). Reductive methylation was with [¹⁴C]formaldehyde and NaCN-BH₄ as described by Jentoft & Dearborn (1979). ⁶⁰Co was purchased from New England Nuclear and had a specific activity of 123 Ci/mg. Urea and guanidinium chloride were from K & K Laboratories, Rare and Fine Chemicals, Plainview, NY. All other chemicals were reagent grade or better.

Results

Effect of Co²⁺ on the Activity of Transcarboxylase Incubated at pH 8. The results shown in Figure 2 were obtained when transcarboxylase was diluted into 50 mM Hepes buffer, pH 8, at 25 °C, with and without the addition of 2 mM Co²⁺, and was assayed with time. The inclusion of 2 mM Co²⁺ in the incubation buffer results in a pronounced protection of the activity of the enzyme. Similar results are obtained when the experiment is done at pH 9. The effect of Co²⁺ is most pronounced at low protein concentrations (≤1 mg/mL). No protective effect was observed with 2 mM Mg²⁺, Mn²⁺, Fe²⁺, Cu²⁺, Ni²⁺, or Ca²⁺, but a slight protection was noted in the presence of Zn²⁺ (Figure 2). If Co²⁺ is added to samples that have been incubated without Co²⁺ for 5, 10, or 20 min, there is no reactivation of enzyme. Inactivation of transcarboxylase

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¹ Abbreviations: DTNB, 5,5'-dithiobis(2-nitrobenzoic acid); GdmCl, guanidinium chloride; Hepes, *N*-(2-hydroxyethyl)piperazine-*N'*-2-ethanesulfonic acid.

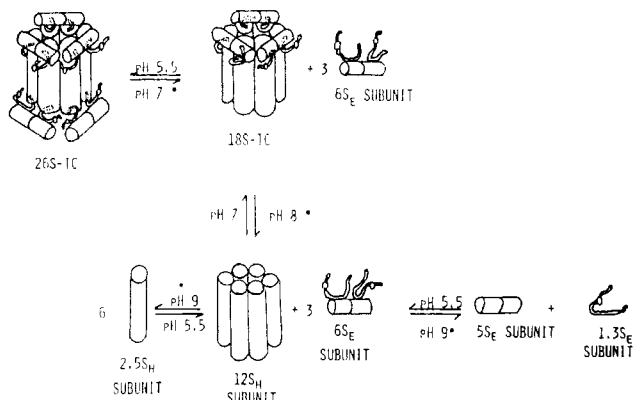


FIGURE 1: Schematic diagram of transcarboxylase and its dissociation to subunits. The undissociated 26S-TC has a sedimentation coefficient of 26 S and is designated 26S-TC (M_r , 1.2×10^6). At neutral pH, there is selective dissociation of outer 6S_E subunits (M_r , 1.44×10^5) from one face of the central 12S_H subunit, giving rise to the 18S form of the enzyme (M_r , 7.9×10^5). At a more alkaline pH, further dissociation occurs, and at about pH 8, no 6S_E subunits remain attached to the central 12S_H subunit. At pH 9, the 6S_E subunits dissociate to the 5S_E subunit (M_r , 1.2×10^5) and two 1.3S_E biotinyl subunits (M_r , 1.2×10^4). The central hexameric 12S_H subunit also dissociates at pH 9 to six 2.5S_H monomeric subunits, but oxygen must be excluded to obtain this reversible dissociation (Bahler et al., 1981). The 12S_H and 6S_E subunits have no enzymatic activity. The 5S_E subunit contains tightly bound Co²⁺ and Zn²⁺, which is required for catalysis. Secondary Co²⁺ atoms bind at pH 8 and 9 and stabilize the quaternary structure of both transcarboxylase and its subunits (see the text). The steps at which stabilization has been demonstrated are indicated by asterisks.

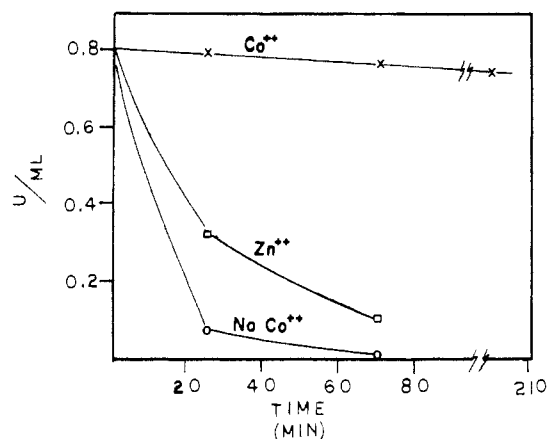


FIGURE 2: Effect of Co²⁺ and Zn²⁺ on enzymatic activity of transcarboxylase at pH 8. Transcarboxylase (0.8 unit/mL) in 50 mM HEPES buffer, pH 8.0, was incubated at 25 °C and assayed with time. (X) With 2 mM Co²⁺; (□) with 2 mM Zn²⁺; (○) with no divalent metal.

in the absence of Co²⁺ is much slower at low temperatures (4 °C).

Effect of Co²⁺ on Inactivation at 50 °C and by Guanidinium Chloride. The loss of activity with time at 50 °C with or without Co²⁺ is shown in Figure 3A. With Co²⁺, the activity decreased rapidly in the first 2 min, but beyond that time there was about 4% residual activity that persisted past 25 min. The sample without Co²⁺ was devoid of activity in 1 min. On the other hand, the sample that was preincubated with Co²⁺ for 20 min at pH 8 at 25 °C and was then warmed to 50 °C lost ~75% of its activity within 4 min, but after that the activity decreased only slowly over a period of 1 h. Preincubation of transcarboxylase with Co²⁺ under conditions where little dissociation occurs (for example, in phosphate buffer, pH 6.3) provides little protection against denaturation at 50 °C at pH 8. Apparently, it is necessary for the conformation of the

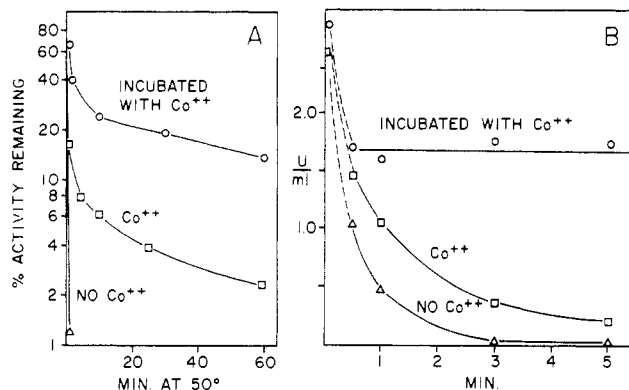


FIGURE 3: Effect of Co²⁺ on inactivation of transcarboxylase at 50 °C or by guanidinium chloride (GdmCl). (A) Transcarboxylase (0.5 mg/mL) (15.2 units/mL) in 150 mM HEPES buffer, pH 8.0, was incubated at 50 °C. (○) Preincubated 20 min at 25 °C in buffer, pH 8, containing 2 mM Co²⁺ prior to heating to 50 °C; (□) 2 mM Co²⁺ without prior preincubation; (Δ) no Co²⁺. (B) Transcarboxylase (0.1 mg/mL, 2.75 units/mL) in 50 mM HEPES, pH 9.0, at 25 °C was incubated 30 min with 2 mM Co²⁺ (the activity then was 2.05 units/mL); then (time zero on the graph) 6 M GdmCl was added to give 0.25 M GdmCl (○); GdmCl and 2 mM Co²⁺ were added without preincubation (□); GdmCl was added in the absence of Co²⁺ (Δ).

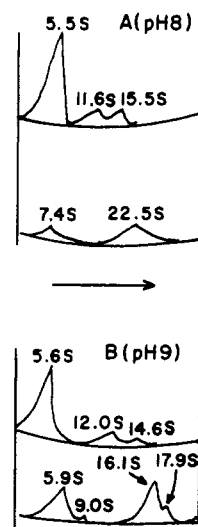


FIGURE 4: Sedimentation velocity profiles of transcarboxylase with and without Co²⁺ at pH 8 and 9. Aliquots of transcarboxylase (4–5 mg/mL) were dialyzed 18 h under N₂ at 5 °C vs. 50 mM HEPES buffer, pH 8 or pH 9, with or without 2 mM Co²⁺. At the end of this time, the activities at pH 8 were 5.5 units/mg without Co²⁺ and 15.5 units/mg with Co²⁺, and at pH 9, they were 5.7 units/mg without Co²⁺ and 16.4 units/mg with Co²⁺. Centrifugation was in a Beckman Model E ultracentrifuge with an AnD rotor at 60000 rpm. (A) pH 8, without Co²⁺ (top) and with Co²⁺ (bottom); (B) pH 9, without Co²⁺ (top) and with Co²⁺ (bottom).

enzyme to be “opened up” at alkaline pH so that the Co²⁺ can occupy sites that stabilize the enzyme.

Transcarboxylase is rapidly inactivated in 0.5 M GdmCl. With 0.026 mg/mL (0.85 unit/mL) transcarboxylase in 50 mM HEPES at pH 8, all activity was lost in 3 min. In the presence of 2 mM Co²⁺, there was loss of activity, but 40–50% of the activity was retained at 1 h. At pH 9, the protection by Co²⁺ against inactivation was not very effective if the GdmCl was added at the same time as the Co²⁺. However, if the enzyme was preincubated with the Co²⁺ at pH 9 for 30 min at 25 °C and then the GdmCl (0.25 M) was added, a pronounced protection resulted with no further loss of activity even after 40 min (Figure 3B). These results again indicate that it is necessary to introduce the Co²⁺ into the sites prior to subjecting the enzyme to destabilization.

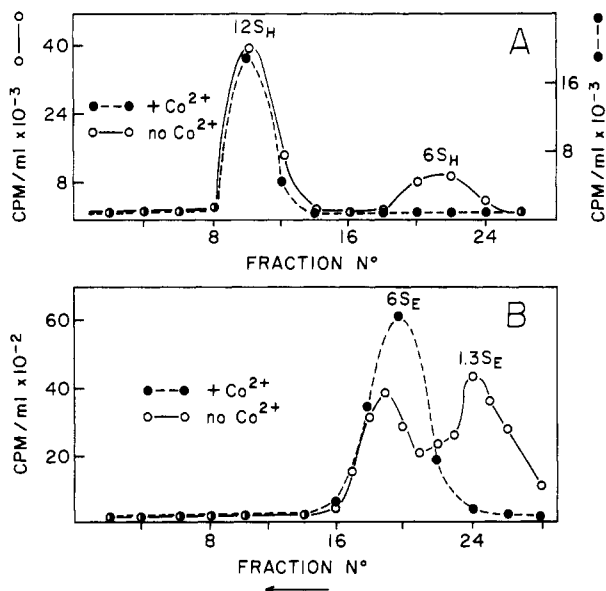


FIGURE 5: Effect of Co²⁺ on dissociation of 12S_H and 6S_E subunits. (A) 12S_H subunit (0.56 mg/mL and 5.9×10^3 cpm/mg) that had been lightly methylated with ¹⁴CH₂O was incubated in 50 mM Hepes, pH 8.0, without Co²⁺ and with 2 mM Co²⁺ for 2 h at 25 °C and then for 16 h at 4 °C. 300 μ L of the sample without Co²⁺ was layered on 11-mL 10–30% glycerol gradients in 50 mM Hepes, pH 8.0, containing no Co²⁺, and that containing Co²⁺ was layered on a gradient containing 2 mM Co²⁺. Centrifugation was at 39000 rpm in an SW 41 rotor for 18 h at 4 °C. Fractions of 0.3 mL were collected and counted. Fractions 8–12 contained undissociated 12S_H subunit while fractions 18–24 contained 6S_H dimer. (B) 6S_E subunit (1.3 mg, 13000 cpm in 0.3 mL) containing [³H]biotinyl 1.3S_E subunit was treated the same as the 12S_H subunit. Fractions 23–28 contained free 1.3S_E biotinyl subunit (M_r 1.2×10^4) whereas fractions 17–21 contained undissociated 6S_E subunit. A control gradient in 0.3 M phosphate buffer, pH 6.3, gave a profile like that with Co²⁺.

Effect of Co²⁺ on Quaternary Structure of Transcarboxylase. Sedimentation velocity profiles of transcarboxylase are shown in Figure 4, with (A) being at pH 8 and (B) at pH 9. In the absence of Co²⁺ (top), the majority of the enzyme was converted to 5.6S material, which presumably is a mixture of 5S_E, 6S_E, and 6S_H subunits (see Figure 1). There was some 12S_H subunit present and not much active enzyme, which has a sedimentation coefficient greater than 12 S. No 2.5S_H monomer was observed. The dissociation was done under N₂, but mercaptoethanol was not added because at alkaline pH with Co²⁺ a colored precipitate forms. Perhaps even under N₂, mercaptoethanol is necessary to prevent formation of disulfide-linked 6S_H dimers as described by Bahler et al. (1981).

In the presence of Co²⁺ (bottom), at pH 8, a peak with an $s_{20,w}$ value of 22.5 S was observed, and at pH 9 there were peaks at $s_{20,w}$ values of 16.1 and 17.9 S. No 12S_H subunit was present. Clearly, the Co²⁺ promoted the binding of the 6S_E subunits to the 12S_H subunit and retarded the dissociation of transcarboxylase into inactive subunits. The dissociation of transcarboxylase is rapidly reversible and the 22.5S peak at pH 8 in the presence of Co²⁺ represents forms of transcarboxylase with an average of about four attached outside subunits (Poto & Wood, 1977).

Effects of Co²⁺ on Dissociation of 12S_H and 6S_E Subunits. Figure 5A shows the results obtained at pH 8 by glycerol gradient centrifugation of 12S_H subunit that had been lightly labeled with [¹⁴C]formaldehyde. Undissociated 12S_H subunit was in fractions 8–12, and the 6S_H subunits were in fractions 18–24. In these experiments, precautions were not taken to keep the solutions anaerobic, and under these conditions, disulfide bonds are formed between the 2.5S_H monomers to yield

6S_H dimers (Bahler et al., 1981). Clearly, there was dissociation in the absence of Co²⁺ (solid line), but in the presence of Co²⁺, dissociation did not occur and only the 12S_H subunit was observed. A control sample centrifuged in phosphate buffer, pH 6.3, in which 12S_H remains intact gave a profile identical with that observed at pH 8 in the presence of Co²⁺. The 12S_H subunit of fraction 10 was tested to determine its competence in forming active enzyme. An excess of 6S_E subunit was added to the fraction in 0.5 M phosphate buffer at pH 6.5 at 4 °C. The high phosphate concentration promotes combination of the subunits and formation of active enzyme (Wood et al., 1975). The reconstituted enzyme had a specific activity of 75 units/mg of 12S_H, which was the same as that obtained with the untreated 12S_H subunit. Fraction 22 (6S_H dimer) did not reconstitute to active enzyme with 6S_E subunit.

A similar experiment was conducted with the 6S_E subunit, which contained [³H]biotin in the 1.3S_E subunit (Figure 5B). The undissociated 6S_E subunit was in fractions 17–21 and dissociated 1.3S_E subunit in fractions 23–28. In the presence of added Co²⁺ (dashed line) there was no dissociation. A control centrifugation in phosphate buffer at pH 6.3 gave results identical with those shown in Figure 5B with Co²⁺. Clearly, Co²⁺ not only retards the dissociation of 6S_E from the 12S_H subunit of transcarboxylase but also retards the dissociation of the 12S_H to monomers and the 6S_E subunit to 5S_E and 1.3S_E subunits.

Capacity of Co²⁺-Treated 6S_E and 12S_H Subunits To Reconstitute to Active Transcarboxylase. The 12S_H or 6S_E subunit was incubated at pH 8 or pH 9 with and without Co²⁺ for 2 h at 30 °C and then was reconstituted to active enzyme by combination with the other subunit, which had not been treated with Co²⁺ or with base. Aliquots were taken with time and assayed. Figure 6A shows the results when the 6S_E subunit was subjected to alkaline conditions and Figure 6B when the 12S_H subunit was subjected to this treatment. It is apparent from Figure 6A that Co²⁺ protected the capacity of the 6S_E subunit to reconstitute with 12S_H subunit. This is most evident at pH 9 where the dissociation of the 6S_E subunit to 5S_E and 1.3S_E subunits is quite complete (Figure 1). Although not shown here, this difference between 6S_E subunit treated with and without Co²⁺ persisted even when the 6S_E subunit was incubated at pH 8 over a longer period of time. It should be noted that, although the dissociation of the 6S_E subunit to 5S_E and 1.3S_E subunits is reversible, an excess of 1.3S_E subunit is required to obtain complete conversion of the 5S_E subunit to 6S_E subunit (Wood et al., 1975).

When the 12S_H subunit was subjected to alkaline treatment, the effect of Co²⁺ was more pronounced (Figure 6B) than with the 6S_E subunit. In part, this may be because the experiments were done in solutions exposed to air, and therefore, during dissociation of 12S_H to the monomer, conversion to the disulfide-linked 6S_H dimer occurred, and this dimer does not reconstitute with the 6S_E subunit to form active transcarboxylase (Bahler et al., 1981). The results with both the 6S_E and 12S_H subunits indicate that Co²⁺ retards dissociation of the subunits, and thus their capacity for reconstitution is retained.

Stability of Co²⁺ Bound to Transcarboxylase and Stoichiometry of Co²⁺ Binding. For these experiments, transcarboxylase was incubated at pH 8 with or without Co²⁺ until the control sample (no Co²⁺) had 25% of the enzymatic activity of the sample with ⁶⁰Co²⁺. Because of the high protein concentration, several hours at 30 °C were required for the inactivation. Samples were then precipitated with ammonium sulfate, dialyzed against either acetate buffer, pH 5.5, or

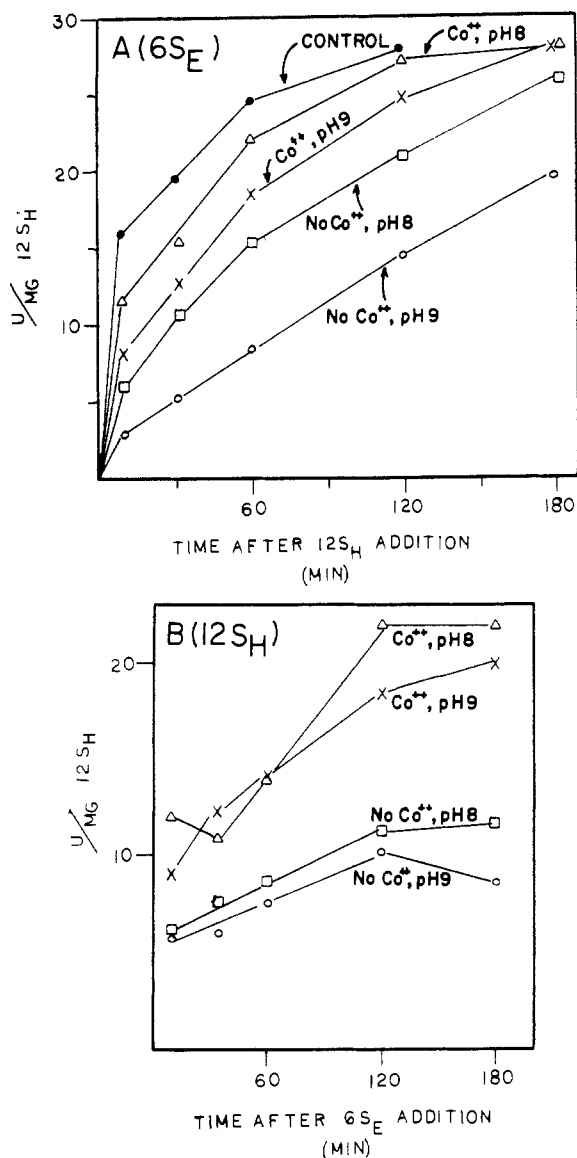


FIGURE 6: Effect of Co^{2+} on capacity of 6S_E and 12S_H subunits to reconstitute to active enzyme following treatment at pH 8 or pH 9. The subunit, 6S_E (10 μg) or 12S_H (1.4 μg), each in 150 μL of 50 mM Hepes buffer, pH 8 or 9, was incubated with or without the addition of Co^{2+} for 2 h at 30 $^\circ\text{C}$. Reconstitution to active enzyme was done by adding to the treated 6S_E subunit 150 μL of 1.5 M phosphate buffer, pH 6.3, at 4 $^\circ\text{C}$ containing 1.4 μg of untreated 12S_H subunit or to the treated 12S_H subunit 150 μL of the phosphate buffer containing 10 μg of untreated 6S_E subunit. The mixtures were then assayed with time for transcarboxylase activity. A control in which neither subunit was treated at pH 8 or pH 9 was also done. (A) Untreated 12S_H subunit plus treated 6S_E subunit: (O) pH 9, no Co^{2+} ; (X) pH 9, plus Co^{2+} ; (□) pH 8, no Co^{2+} ; (▲) pH 8, plus Co^{2+} ; (●) control. (B) Untreated 6S_E subunit plus treated 12S_H subunit: (O) pH 9, no Co^{2+} ; (X) pH 9, plus Co^{2+} ; (□) pH 8, no Co^{2+} ; (▲) pH 8, plus Co^{2+} . All results are expressed in units per milligram of 12S_H since in the reconstitution the 6S_E subunit is in excess.

phosphate buffer, pH 6.8, and then centrifuged in glycerol gradients buffered at pH 5.5 or pH 6.8. At pH 5.5, the 26S form is stable, whereas at pH 6.8, the 18S form, with only three outer 6S_E subunits, is predominant (Figure 1). No Co^{2+} was included in the gradient buffers so that only those Co^{2+} ions that remained bound during ammonium sulfate precipitation and dialysis were associated with the enzyme. Figure 7 shows the results of such an experiment. Fractions 21–25 contained the 6S_E subunit while fractions 3–12 contained the enzyme. In the sample centrifuged at pH 5.5, some 6S_E subunit was present, indicating either that there was some

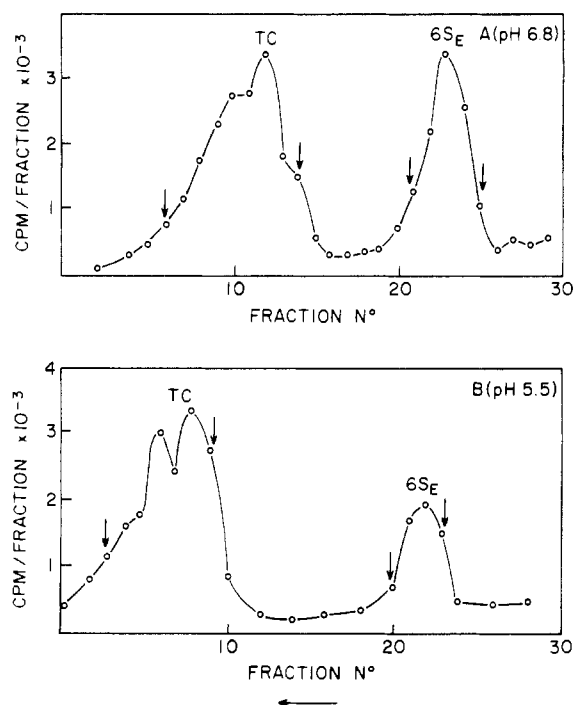


FIGURE 7: Stability and stoichiometry of ^{60}Co bound to transcarboxylase. 26S transcarboxylase was incubated at 25 $^\circ\text{C}$ for 5 h in 50 mM Hepes, pH 8, containing 2 mM $^{60}\text{Co}^{2+}$ (2330 cpm/nmol). It then contained 9.11 mg/mL and had an enzymatic activity of 30.2 units/mg. A control treated similarly without Co^{2+} contained 9.76 mg/mL and an activity of 7.9 units/mg. It was not used in the remainder of the experiment. The sample with $^{60}\text{Co}^{2+}$ was brought to pH 6.7 with 100 μL of 1 M KH_2PO_4 and was precipitated by addition of $(\text{NH}_4)_2\text{SO}_4$ to 60% saturation. The precipitate was dissolved in cold 100 mM acetate buffer, pH 5.5, and half was dialyzed at 4 $^\circ\text{C}$ against this buffer. The other half was dialyzed against 50 mM phosphate buffer, pH 6.8. The resulting sample at pH 5.5 had an enzymatic activity of 42 units/mg and 24 100 cpm/mg and that at pH 6.8 of 23 units/mg and 16 200 cpm/mg. The samples were applied to a 10–30% glycerol gradient buffered with 100 mM acetate buffer, pH 5.5, or 50 mM phosphate buffer, pH 6.8, and centrifuged at 39 000 rpm in an SW 41 rotor for 16 h at 4 $^\circ\text{C}$. Fractions of 0.3 mL were collected, assayed for radioactivity, and pooled (arrows). (A) Profile in pH 6.8 buffer. Fractions 4–12 contained enzyme; fractions 21–25 contained 6S_E subunits. (B) Profile in pH 5.5 buffer. Fractions 3–7 contained enzyme; fractions 21–24 contained 6S_E subunits. The stoichiometry of $^{60}\text{Co}^{2+}$ binding is given in Table I.

dissociation of the 26S enzyme or that after the treatment at pH 8, the reassociation to 26S enzyme was incomplete. Since native 26S enzyme does not dissociate at pH 5.5, we assume the latter was the case.

Pools from the gradients (arrows) were assayed for protein, enzyme activity, and radioactivity, and the results are shown in Table I. The moles of ^{60}Co bound per mole of enzyme and per mole of 6S_E subunit were calculated on the basis of the specific radioactivity of the $^{60}\text{Co}^{2+}$ and the cpm per milligram of protein observed in the fraction. The values are approximations but they show that ^{60}Co was bound to the 6S_E subunit. Since Co^{2+} has a stabilizing effect on the isolated 12S_H subunit (Figures 5A and 6B), it is likely a portion of the Co^{2+} is also bound to the 12S_H subunit in the transcarboxylase.

The experiment of Figure 8 was done to determine if enzyme into which Co^{2+} was introduced at pH 8 is stable in the absence of exogenous Co^{2+} . For this purpose, the labeled enzyme of fractions 4–12 of Figure 7A was used, and it was found that this enzyme was only slightly more resistant to inactivation at pH 8 than transcarboxylase that had not been treated with Co^{2+} . This result indicates either the Co^{2+} dissociated from the enzyme when incubated at pH 8 or, if it remained attached, it is not responsible for the observed stabilizing influence of

Table I: Cobalt Content of Pooled Fractions from Glycerol Gradients of Figure 7

pooled fractions	[protein] (mg/mL)	sp act. (units/mg)	⁶⁰ Co (cpm ^a /mg)	mol of ⁶⁰ Co/mol of enzyme or subunit
7A (pH 6.8)				
4-12 (TC)	0.40	32.7	15 666	5.3 ^b
21-25 (6S _E)	0.29	0	20 800	1.3 ^b
7B (pH 5.5)				
3-7 (TC)	0.52	33.6	13 212	5.7 ^b
21-24 (6S _E)	0.14	0	33 646	2.1 ^b

^a ⁶⁰Co contained 2330 cpm/nmol. ^b For these calculations, it has been assumed that the enzyme obtained at pH 6.8 was 18S transcaboxylase, M_r 7.9×10^5 . Considerable 6S_E subunit was observed in the glycerol gradient at pH 5.5 (Figure 7B). Therefore, it is assumed the enzyme had four outer subunits and an M_r of 9.36×10^5 . The ⁶⁰Co content of the 6S material has been calculated on the basis that it was predominantly 6S_E subunit of M_r 1.44×10^5 .

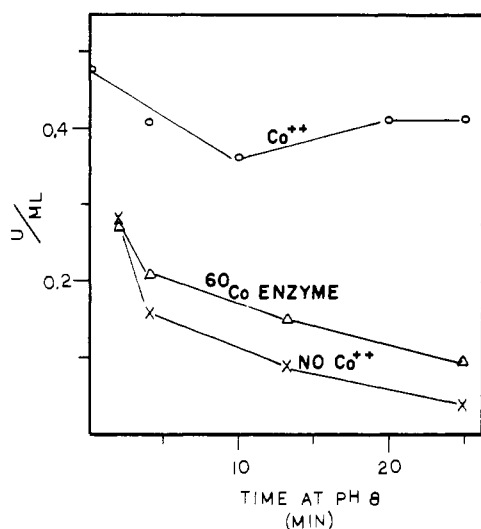


FIGURE 8: Effect of incubation at pH 8 on enzyme in which Co²⁺ had been introduced at pH 8. Aliquots of enzyme were diluted to 0.45 unit/mL into pH 8 buffer (25 °C) and assayed with time. (O) Untreated enzyme diluted in pH 8 buffer containing 2 mM Co²⁺; (X) untreated enzyme at pH 8, no Co²⁺; (Δ) 18S enzyme from fractions 4-12 of Figure 7A containing ⁶⁰Co.

2 mM Co²⁺. To investigate this possibility, we dialyzed the labeled enzyme (fractions 4-12 of Figure 7A) against 50 mM Hepes buffer, pH 8.0, for 7 h at room temperature. Following this treatment, 70% of the ⁶⁰Co remained attached to the enzyme. Passage of the enzyme through a column of Chelex resin at pH 9 likewise failed to remove the radioactivity from the enzyme; half of the ⁶⁰Co emerged from the column still attached to the dissociated enzyme. Since the tightly bound Co²⁺ did not stabilize the enzyme, it appears that weakly bound Co²⁺ ions are responsible for the stabilizing effect. Presumably, the weakly bound Co²⁺ is in equilibrium with unligated Co²⁺ in solution, and 2 mM Co²⁺ is required to maintain the concentration of bound Co²⁺ sufficiently high to stabilize the enzyme.

Comparison of Catalytic Cobalt and Cobalt Incorporated at pH 8. Native transcaboxylase contains cobalt that is considered to be at the keto acid site on the 5S_E subunit (Northrop & Wood, 1969; Ahmad et al., 1972; Fung et al., 1974) and which we have designated catalytic cobalt. An experiment was done with transcaboxylase isolated from cells grown in medium containing ⁶⁰Co (Ahmad et al., 1972). The specific radioactivity of this transcaboxylase was 12 000

cpm/mg. Aliquots of the ⁶⁰Co transcaboxylase were incubated at pH 8 with or without added unlabeled Co²⁺ (2 mM). A third sample of the ⁶⁰Co transcaboxylase was incubated with 2 mM Co²⁺ to which ⁶⁰Co²⁺ was added as a tracer. When the sample without Co²⁺ had 5% of the enzymatic activity of the Co²⁺-containing samples, all the samples were dialyzed extensively vs. 0.25 mM acetate buffer, pH 5.5, and then assayed for specific radioactivity. The samples of ⁶⁰Co transcaboxylase that had been incubated in the presence and absence of unlabeled Co²⁺ both remained at a specific radioactivity of ~12 000 cpm/mg. Hence, no loss or exchange with the catalytic cobalt of the enzyme took place. In addition, the sample incubated at pH 8 in the presence of added ⁶⁰Co²⁺ increased in specific radioactivity to 19 000 cpm/mg, indicating that the incubation led to the binding of additional Co²⁺ ions that are distinct from those that are tightly bound in the 5S_E subunit.

Effect of Co²⁺ on Titratable Sulfhydryl Residues of Transcaboxylase. It has been shown (Bahler et al., 1981) that the reconstitution of dissociated transcaboxylase is most effective if the dissociation and reconstitution are done under anaerobic conditions, which prevent oxidation of sulfhydryl groups. It seemed possible that Co²⁺ protected the sulfhydryl groups at pH 8. We, therefore, tested the effects of Co²⁺ on the total number of titratable thiol groups in the enzyme. Identical aliquots of 26S transcaboxylase (8 mg/mL) were incubated at 25 °C at pH 8, in the presence or absence of 2.5 mM ⁶⁰Co²⁺ (1542 cpm/nmol of Co²⁺). After the incubation, the sample without Co²⁺ had a specific enzymatic activity of 19.5 units/mg whereas that incubated with Co²⁺ had an activity of 28 units/mg. Each sample was then desalted on Sephadex G-25 in 50 mM Hepes buffer, pH 8, and the enzyme in the protein fraction was titrated with DTNB in the presence of sodium dodecyl sulfate and EDTA (Habeeb, 1972). The EDTA was included in the reactions to prevent any metal-catalyzed oxidation of thiol groups. The starting transcaboxylase has 102 cysteines by amino acid composition (Wood & Zwolinski, 1976). The transcaboxylase incubated at pH 8 with Co²⁺ contained 62.3 titratable sulfhydryl groups whereas that incubated without Co²⁺ contained 78 titratable groups. Hence, the sample incubated with Co²⁺ possessed 16 fewer titratable sulfhydryl groups than that incubated without Co²⁺. The specific radioactivity of the transcaboxylase incubated with Co²⁺ was 11 708 cpm/mg. This corresponds to 8.6 mol of ⁶⁰Co/mol of 26S transcaboxylase. There was substantial dissociation to subunits during the desalting at pH 8, and the specific activity of the enzyme without Co²⁺ decreased to 1.8 units/mg and that with Co²⁺ to 4 units/mg. However, all the subunits of transcaboxylase are present in the void volume of a Sephadex G-25 column; thus the total sulfhydryl group determination on the protein peak is valid. When this experiment was repeated with a different sample of 26S transcaboxylase, similar results were obtained. The starting transcaboxylase had 80 thiol groups, that incubated without Co²⁺ had 73.5, and that incubated with Co²⁺ had 55, a net loss of 18.5 thiol groups. Thus, for each Co²⁺ bound, there were approximately two less titratable thiol groups.

It therefore seemed possible that the Co²⁺ was ligated between pairs of thiol groups. Such ligands should have an absorption band at 340 nm due to charge transfer between cobalt and thiol ligands as observed by Sytkowski & Vallee (1976) using cobalt-labeled alcohol dehydrogenase. To test this possibility, we incubated transcaboxylase (~5 mg/mL) at 25 °C at pH 8 in the presence and absence of 2 mM Co²⁺ until the sample without Co²⁺ lost greater than 80% of its

original activity. The samples were then dialyzed against 50 mM sodium acetate buffer, pH 5.5, at 4 °C for 16 h to remove unbound metal ions, and the difference spectra were recorded. On the basis of the extinction coefficient given by Sytkowski & Vallee (1976), a significant difference would have occurred if cobalt–thiol bonds had been formed. None was observed; thus, it is concluded the cobalt ions are not ligated to the enzyme through cysteine linkages. Perhaps the decrease in titratable thiol groups is due to stimulation of oxidation of sulfhydryl by the Co^{2+} per se.²

Discussion

It has been demonstrated that added Co^{2+} (2 mM) protects transcarboxylase from the loss of enzymatic activity that normally occurs at 25 °C in 50 mM Hepes buffer at pH 8. The effect is specific for Co^{2+} . Mg^{2+} , Mn^{2+} , Fe^{2+} , Cu^{2+} , Ni^{2+} , and Zn^{2+} are ineffective, and there is only a slight protection by Zn^{2+} (Figure 2). There also is protection at an alkaline pH against denaturation by guanidinium chloride (0.5 M) and by heat (50 °C) (Figure 3). This protection is most effective if the enzyme is first incubated with the Co^{2+} and then the enzyme is subjected to the guanidinium chloride or heat treatment. If the pretreatment with Co^{2+} is in phosphate buffer at pH 6.5, there is little or no protection. Apparently, an alkaline pH is required to "open" the conformation of the enzyme so that the sites that bind the Co^{2+} are exposed. Such exposure may result from the dissociation of transcarboxylase to subunits at pH 8 as illustrated in Figure 1.

Transcarboxylase is a metalloenzyme and contains very firmly bound cobalt and zinc. The proportion of each metal in the enzyme depends on the relative concentration of Co^{2+} and Zn^{2+} in the medium used to grow the bacteria (Ahmad et al., 1972). The cobalt and zinc have been shown to be near the keto acid site in the outer 5S_E subunit (Northrop & Wood, 1969; Ahmad et al., 1972; Fung et al., 1974). At pH 8, Co^{2+} did not replace ^{60}Co of labeled enzyme, and when the labeled enzyme was treated at pH 8 with $^{60}\text{Co}^{2+}$, there was an increase of radioactivity due to incorporation of additional ^{60}Co . Clearly, the cobalt that is introduced at pH 8 does not enter and replace the cobalt at the catalytic site.

It was shown by ultracentrifugation that in the absence of Co^{2+} at pH 8 or pH 9, the enzyme is largely dissociated to its subunits, whereas in the presence of Co^{2+} , the various forms of the active enzyme still persist (Figure 4). Outer 5S_E subunits are bound to the 12S_H subunits via the biotinyl subunit (see Figure 1), and it has been shown (Kumar & Beegen, 1981) that the binding to the 12S_H subunit involves residues 2–14 of the biotinyl subunit and binding to the 5S_E subunit residues 15–26. However, the effect of Co^{2+} on the dissociation

of transcarboxylase is not confined to these binding sites, since Co^{2+} retards the dissociation at pH 8 of the 12S_H subunit and likewise that of the 6S_E subunit to the 5S_E subunit and the 1.3S_E biotinyl subunit (Figure 5). Thus, the Co^{2+} effects multiple types of binding of the subunits. The effect of Co^{2+} is also evident in maintaining the capability of the 12S_H and 5S_E subunits to reconstitute to active transcarboxylase (Figure 6).

The evidence available suggests that there are three types of cobalt binding to the enzyme. The one is at the catalytic cobalt. The second type is formed when the enzyme is incubated with exogenous Co^{2+} at pH 8, during which some Co^{2+} becomes tightly bound and is not removed by dialysis, ammonium sulfate precipitation, or treatment at alkaline pH or with a Chelex resin. There are approximately six such cobalts per 26S transcarboxylase molecule. These cobalts do not appear to be bound to the enzyme through thiol linkages and, by themselves, have little influence on the dissociation properties of transcarboxylase. A third class of cobalt apparently is more weakly bound and is in equilibrium with the free Co^{2+} , which must be maintained at about 2 mM in order to stabilize the enzyme at alkaline pH. This weakly bound Co^{2+} effects the equilibrium of the association and dissociation of transcarboxylase. At 25 °C in the presence of 2 mM Co^{2+} at pH 8, the dissociation is not complete and partial activity is retained by the enzyme. It is considered that information concerning the sequence of amino acids in the 12S_H and 5S_E subunits and perhaps their quaternary structure will be required before an explanation of this stabilizing effect of exogenous cobalt will be forthcoming.²

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² It has been suggested by one of the reviewers that the Co^{2+} may have been oxidized to Co^{3+} , which forms a more stable complex, and that if the complex was treated with dithionite, the Co^{3+} would be reduced to Co^{2+} and should easily be removed by a Chelex column. We have treated transcarboxylase with $^{60}\text{Co}^{2+}$ at pH 8, removed the unbound $^{60}\text{Co}^{2+}$ by dialysis, and then passed the enzyme with and without treatment with dithionite over Chelex. The treatment with dithionite did not alter the amount of $^{60}\text{Co}^{2+}$ that was tightly bound. This provides proof that the tightly bound cobalt is not Co^{3+} and in addition that the Co^{2+} is not ligated between thiols.