

The interactions of cobalt(II) with mitochondria from rat liver

Marcantonio Bragadin · Antonio Toninello ·
Mario Mancon · Sabrina Manente

Received: 10 October 2006 / Accepted: 22 January 2007
© SBIC 2007

Abstract The interactions of Co^{2+} with mitochondria have been investigated. The results indicate that Co^{2+} inhibits ATP synthesis. Further investigations into ATP synthesis mechanisms indicated that inhibition is due to the opening of a transmembrane pore. The opening of this pore causes the collapse of the high-energy intermediate where, under a pH and a potential gradient, the energy is stored and subsequently utilized to form ATP from ADP.

Keywords Cobalt · Mitochondria · Energy synthesis

Introduction

Cobalt has both beneficial and harmful effects on human health. It is beneficial for humans because it is a part of vitamin B_{12} , which is essential for the maintenance of human health. When too much cobalt is taken into the body, it can produce harmful effects, which have been the subject of many *in vivo* studies. *In vitro* investigations have also been performed, in order to establish the molecular mechanism responsible for these effects on whole organisms. However, the exact molecular mechanism that Co^{2+} exerts on cells has still not been identified, although a number of potential mechanisms have been proposed [1–3]. The aim of the work presented here was not to carry out

a toxicological investigation, but rather a study of the molecular mechanisms of the interaction of Co^{2+} with biological structures, which can subsequently be utilized for toxicological studies.

Since, in many cases, Co^{2+} causes apoptosis in isolated cells [4, 5], and since the apoptosis can be correlated with mitochondrial function, we investigated the interactions of Co^{2+} with mitochondria. The results indicated that Co^{2+} actually does inhibit ATP synthesis, and further investigations were then carried out to identify the step responsible for ATP synthesis inhibition. This step appears to be the opening of a membrane pore which reduces the energy accumulation in the mitochondria.

Materials and methods

The mitochondria were prepared from the livers of fasted albino Wistar rats, weighing about 300 g [7]. Mitochondrial protein was determined using the Lowry procedure [8].

The mitochondrial oxygen consumption was measured using a Clark oxygen electrode (Yellow Springs Instruments, Yellow Springs, OH, USA) fitted in a thermostat-controlled, closed chamber with a magnetic stirrer. The reaction medium (2 ml) was maintained at 25 °C throughout all the experiments (i.e., under standard *in vitro* conditions).

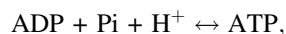
The following reducing substrates were utilized: succinate, glutamate/malate and ascorbate/*N,N,N',N'*-tetramethyl-*p*-phenylenediamine (TMPD).

To follow the absorbance decrease, the swelling experiments were performed at 540 nm, using a Jenway 6400 (Felsted, UK) spectrophotometer, equipped with stirring apparatus. After the addition of the mitochondria to the resuspending medium (2.5 ml), the instrument was adjusted to zero absorbance.

M. Bragadin (✉) · S. Manente
Dipartimento di Scienze Ambientali,
Università Cà Foscari,
Venezia DD 2137, 30123 Venice, Italy
e-mail: bragadin@unive.it

A. Toninello · M. Mancon
Dipartimento di Chimica Biologica,
Università di Padova e Centro Studio delle Biomembrane,
C.N.R. Padova, Viale G. Colombo 3, 35121 Padua, Italy

The ATP synthesis/hydrolysis experiments were performed in a low-buffered medium, using a pH electrode connected to an 84 PHM Radiometer (pH meter) (Radiometer, Copenhagen, Denmark) in order to monitor the pH changes which accompany the reaction



where Pi is sodium phosphate, under stirring conditions, at room temperature.

All of the reagents were of analytical grade. 2,4-Dinitrophenol (DNP), sodium succinate, sodium glutamate, sodium malate, oligomycin, and TMPD were obtained from Sigma (Milan, Italy).

Results

In mitochondria, the substrates arising from the Krebs cycle are oxidized by molecular oxygen, and ATP is produced. This oxidation occurs by means of a sequence of redox couples, called the mitochondrial respiratory chain (RC). As envisaged in the chemiosmotic hypothesis [9], the electron flow in the RC is coupled to a proton extrusion. Since the (inner) mitochondrial membrane is not permeable to protons, their extrusion gives rise to a ΔpH and a $\Delta\Psi$. The sum of these two components is called the proton motive force (pmf):

$$\text{pmf} = \Delta\text{pH} + \Delta\Psi.$$

Under this form, the free energy arising from the oxidation of the substrates is stored and subsequently utilized to synthesize ATP from ADP.

As a consequence of the mechanism discussed above, inhibition of the ATP synthesis can be due to:

- The inhibition of ATPase (the enzyme that catalyzes the ATP synthesis)
- Inhibition of the RC
- Enhancement of the membrane proton permeability

Therefore, as it has been proven that Co^{2+} inhibits ATP synthesis (Fig. 1) (although this inhibition, in the presence of phosphate, could be due to the formation of the CoHPO_4 complex [10]), each step involved in ATP synthesis was analyzed separately, in order to identify the step that requires the lowest dose of Co^{2+} needed to inhibit the ATP synthesis.

ATPase

As ATPase is the enzyme which catalyzes ATP synthesis, its inhibition implies a corresponding inhibition of ATP synthesis.

As indicated in Fig. 1, the inhibition of ATP synthesis by Co^{2+} is not accompanied by an inhibition of ATP hydrolysis following the production of ATP. This type of behavior was further confirmed by means of direct experiments concerning ATP hydrolysis which can be seen in Fig. 2. The addition of ATP to the uncoupled mitochondria gives rise to a rapid acidification in the (low-buffered) medium. This acidification is a consequence of the following reaction:



The addition of oligomycin (Fig. 2, reaction b), which is an ATPase inhibitor, stops the process of ATP hydrolysis, while the addition of Co^{2+} , up to a concentration of 300 μM , does not induce a cessation in the acidification (Fig. 2, reaction a). This leads to the conclusion that Co^{2+} is not an ATPase inhibitor.

The RC

As the RC is correlated to the proton extrusion, an inhibition of the RC implies an equivalent inhibition of ATP synthesis. The interactions of Co^{2+} with the RC were investigated by stimulating respiring mitochondria by adding ADP and Pi (this was possible because the previous experiments had demonstrated that Co^{2+} is not an ATPase

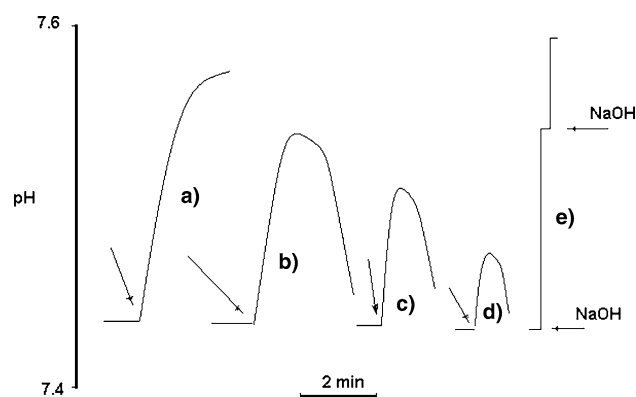


Fig. 1 Co^{2+} inhibits ATP synthesis in mitochondria. The medium composition was 0.25 M sucrose, 0.1 mM *N*-(2-hydroxyethyl)piperazine-*N'*-ethanesulfonic acid (Hepes)/3-(*N*-morpholino)propanesulfonic acid (Mops) pH 7.4, 1 mM succinate, 1 mM MgCl_2 , 0.5 mM sodium phosphate (Pi), and 0.5 mM ADP. The final concentration of mitochondria was 0.5 mg/ml. Since the ATP synthesis induces alkalization, the reaction was followed using a pH meter in a low-buffered medium. The addition of 0 μM (a), 5 μM (b), 10 μM (c), and 15 μM (d) Co^{2+} induces a subsequent inhibition of the alkalization, which demonstrates that Co^{2+} inhibits the ATP synthesis. The arrows indicate the addition of succinate (or NaOH in e). In e, each arrow indicates the addition of 5 nmol NaOH to the medium containing the mitochondria

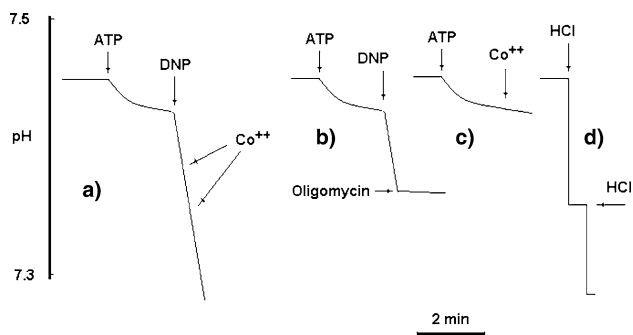


Fig. 2 Co²⁺ does not inhibit ATP hydrolysis in mitochondria. The medium composition was 0.25 M sucrose, 0.1 mM HEPES/MOPS pH 7.4, 1 μg/ml cyclosporine, 1 mM MgCl₂, 1 mM ATP, and 0.1 mM Pi. The final concentration of mitochondria was 0.5 mg/ml. ATP hydrolysis is facilitated by the addition of an uncoupler (0.1 mM 2,4-dinitrophenol). Under these conditions, *a* the rapid ATP hydrolysis is not inhibited by successive additions of Co²⁺ up to 0.3 mM. The effects of the addition of oligomycin (1 μg/ml), a potent ATPase inhibitor, are shown for comparison purposes in *b*. In *c* the addition of 0.2 mM Co²⁺ clearly shows that Co²⁺ is not an uncoupler. In *d*, each *arrow* indicates the addition of 2 nmol HCl to the medium containing the mitochondria

inhibitor). Under these conditions (Fig. 3 shows a typical experiment), the addition of Co²⁺ induces an inhibition of the respiratory rate when succinate is used as a reducing substrate. The graph in Fig. 3 shows the quantitative inhibition of the RC by Co²⁺, but it is evident from the graph that the doses necessary to induce inhibition of the RC are much higher than those needed to inhibit ATP synthesis. When the reducing substrate is either glutamate/malate or ascorbate/TMPD, no RC inhibition occurs (not shown). Therefore, the possibility that the inhibition of the RC by Co²⁺ produces inhibition of ATP synthesis can be excluded.

The enhancement of membrane permeability

The crucial point of the chemiosmotic hypothesis is that the mitochondrial membrane is not permeable to protons; therefore, any mechanism which enhances proton permeability gives rise to an equivalent inhibition of ATP synthesis.

An enhancement of the membrane permeability to protons is called an uncoupling, or a protonophore, effect. A protonophore must be a weak acid (or base). Take, for example, DNP, which was the first protonophore studied [9]. It enters into the matrix as an undissociated compound, the driving force being the internal alkaline pH. Once inside, DNP is extruded as a phenate anion by means of the negative internal potential. The resulting cyclic process implies the entry of a proton at each cycle.

Such a mechanism is, in theory, possible with Co²⁺, as it is a weak acid, but the experiments reported in Fig. 2,

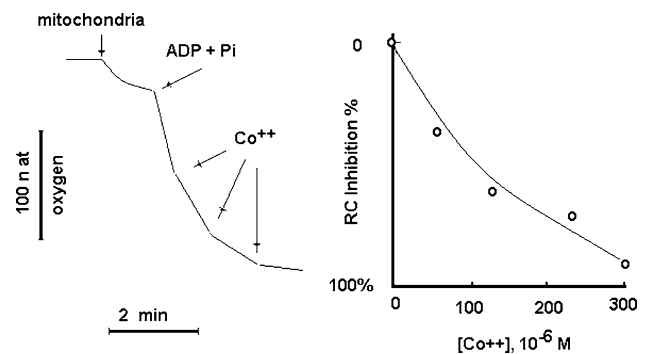


Fig. 3 Co²⁺ inhibits the respiratory chain (RC). The medium composition was 0.25 M sucrose, 10 mM HEPES/MOPS pH 7.4, 1 mM MgCl₂, and 1 mM succinate. The final concentration of mitochondria was 0.5 mg/ml. The addition of 0.5 mM Pi and ADP induces a respiratory rate increase. Under these conditions, successive additions of Co²⁺ induce an inhibition (typical experiment). The *graph* shows the percentage of respiratory rate inhibition as a function of the amounts of Co²⁺ added

reaction *c* exclude this possibility, as it was observed that the mitochondria were energized by ATP. Under these conditions, a steady state occurs between the ATP and the pmf,



and, for this reason, in the absence of an uncoupler, the rate of ATP hydrolysis is very slow and, consequently, the acidification rate is also very slow.

In the presence of an uncoupler, both ΔpH and $\Delta\Psi$ collapse, and a rapid ATP hydrolysis takes place. This hydrolysis can be easily monitored by a pH change in a low-buffered medium (in the same way as in the experiments reported in Fig. 1). In contrast to DNP, Co²⁺ does not stimulate ATP hydrolysis (Fig. 2, reaction *c*). This leads to the conclusion that Co²⁺ is neither an uncoupler nor, for similar reasons, a detergent, since a detergent compound renders permeable the membrane to all solutes, including protons.

Another mechanism which enhances the membrane permeability is the opening of a membrane pore (MTP pore).

Many chemical and toxic compounds induce the opening of a membrane pore. The size of this pore allows the passage of large molecules [10, 11]. If the mitochondria are resuspended in a sucrose medium, the opening of the pore allows the entry of sucrose. The latter occurs in conjunction with the entry of water and, as a consequence of a colloid-osmotic effect, membrane swelling occurs [10]. This swelling, which is inhibited by cyclosporine [10, 11], can be identified by means of absorbance quenching at 540 nm. Figure 4 shows both the swelling induced by Co²⁺ and its inhibition by cyclosporine. These examples demonstrate that Co²⁺ induces the opening of the MTP pore. In

addition, the doses necessary to induce this phenomenon are very similar to those needed to inhibit ATP synthesis (Fig. 1).

Therefore, we have concluded that the inhibition of ATP synthesis by Co^{2+} is a consequence of the opening of the MTP pore, which reduces the energy in the mitochondria, stored in the form of a proton gradient ($\Delta\text{pH} + \Delta\Psi$). It should be noted, however, that it is possible that the effective inducer of the opening of the MTP pore could be the CoHPO_4 complex.

Discussion

As far as the mechanism of action of Co^{2+} is concerned, the problem is not so easily solved. The mechanism of the opening of the MTP pore has been widely studied. It is very complicated, as it depends on many concomitant factors. An exhaustive explanation of the mechanism has still not been given [12]. For this reason, some proposals regarding the mechanism of action of Co^{2+} can only be advanced when making analogies with other metals, other proposals can be excluded.

The involvement of Co^{2+} as a trigger for apoptotic factors, such as the Bcl proteins, normally involved in the opening of the pore during the cell apoptosis would appear to be excluded [13–15].

Recently [16, 17], in this regard, it was demonstrated that, in isolated mitochondria, the effect of Bax proteins (one of the components of the Bcl family), relative to the opening of the MTP pore, occurs with a protein dose (200 nM) very much higher than that normally present in the (outer) mitochondrial membrane. The authors concluded that, under normal conditions, Bax proteins are not

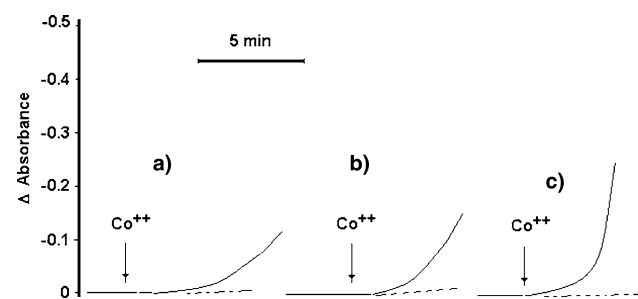


Fig. 4 Co^{2+} induces swelling in mitochondria and this phenomenon is inhibited by cyclosporine. The medium composition was 0.25 M sucrose, 10 mM Hepes/Mops pH 7.4, 1 mM MgCl_2 , 1 mM Pi, and 1 mM succinate. The final concentration of mitochondria was 0.5 mg/ml. Under these conditions, at 540 nm, the addition of 5 μM (a), 10 μM (b), and 15 μM (c) Co^{2+} induces absorbance quenching, which is inhibited by cyclosporine (1 $\mu\text{g}/\text{ml}$; dotted line), thus indicating that the quenching is due to the opening of a pore

involved in the opening of the MTP pore in isolated mitochondria.

Analogously, the experiments of Li et al. [18], regarding the interactions of Cd^{2+} with mitochondria, exclude interactions between Cd^{2+} and Bcl- x_L proteins (proteins of the Bcl family) and suggest (see below) interaction of Cd^{2+} with adenine nucleotide translocase (ANT).

All these conclusions are in agreement with those of other research groups that have studied the interactions of metals with isolated mitochondria: in all cases the proposed mechanism for the opening of the MTP pore excludes the involvement of Bcl proteins. This fact, together with the fact that Bcl proteins are hydrophobic membrane proteins, leads us to analogous conclusions.

Regarding the opening of the MTP pore, then behavior observed in the experiments is similar for many metals, such as Zn^{2+} , Pb^{2+} , Cd^{2+} , Hg^{2+} , and Al^{3+} , and phenylarsine oxide. The experiments performed on isolated mitochondria and cells show that all metals induce the opening of the MTP pore and that this opening is inhibited by cyclosporine (in the case of Cd^{2+} , however, not all authors are in agreement concerning the cyclosporine sensitivity, as will be discussed below).

With regard to Zn^{2+} [19] and Pb^{2+} [20], the opening of the MTP pore has been observed in the presence of micromolar amounts of Ca^{2+} . In our experiments (Fig. 4), Ca^{2+} was not added, but, by operating under conditions similar to those reported in the literature [19, 20] (i.e., 0.25 M sucrose, 10 mM Hepes/Mops pH 7.4, 5 μM CaCl_2), the addition of small amounts of Co^{2+} (around 10 μM) induces a swelling, and the swelling is cyclosporine-sensitive (not shown). The above mentioned evidence appears to indicate similar targets for Co^{2+} , Zn^{2+} , and Pb^{2+} , but in the cases of Zn^{2+} and Pb^{2+} [19, 20] the authors have not proposed a specific action site for these two metals.

Analogously, in the case of Cd^{2+} , a swelling was observed on isolated mitochondria [18, 21–24] and, also in this case the swelling was facilitated by the presence of Ca^{2+} . However, there is no agreement between the authors concerning the cyclosporine sensitivity. In particular, Pourahmad et al. [21] and Dorta et al. [23], found cyclosporine sensitivity, while Belyaeva et al. [24] and Li et al. [18] do not find cyclosporine sensitivity. As a consequence, different mechanisms for Cd^{2+} have been proposed; Li et al. [18] proposed binding of Co^{2+} to the thiol groups in ANT.

Both the swelling and the cyclosporine sensitivity were verified in the case of Al^{3+} and Hg^{2+} [11, 21]. In the case of Al^{3+} “binding of aluminum to the inner mitochondrial membrane, most likely at the level of adenine translocase (ANT)” has been proposed [11].

In the case of phenylarsine oxide, the cyclosporine sensitivity has been verified [25], and it has been proposed

that phenylarsine (and diamide) cause an intramolecular cross-linking between the thiol groups, i.e., Cys¹⁶⁰ and Cys²⁵⁷ [25] in ANT. This cross-linking stabilizes the “c” conformation of ANT. This stabilization should enhance the sensitivity of the MTP pore to the endogenous Ca²⁺ [25].

The strong analogy between Co²⁺, Al³⁺, Cd²⁺, and phenylarsine oxide could suggest similar molecular behavior, i.e., interaction of Co²⁺ with the thiol groups in ANT.

References

- Hervouet E, Pecina P, Demont J, Vojtiskova A, Simonnet H, Houstek J, Godinot C (2006) *Biochem Biophys Res Commun* 344:1086–1093
- Fleury C, Petit A, Mwale F, Antoniu J, Zukor DJ, Tabrizian M, Huk OL (2006) *Biomaterials* 27:3351–3360
- Coddou C, Lorca RA, Acuna-Castillo C, Grauso M, Rassendren F, Huidobro-Toro JP (2005) *Toxicol Appl Pharmacol* 202:121–131
- Catelas I, Petit A, Vali H, Fragiskatos C, Meilleur R, Zukor DJ, Antoniu J, Huk OL (2005) *Biomaterials* 26:2441–2453
- Huk OL, Catelas I, Mwale F, Antoniu J, Zukor DJ, Petit A (2004) *J Arthroplasty* 19:84–87
- Robertson JD, Orrenius S (2000) *Crit Rev Toxicol* 30:609–627
- Azzone GF, Pozzan T, Bragadin M, Miconi V (1979) *J Biol Chem* 254:10213–10219
- Lowry OH, Rosebrough NJ, Farr AL, Randall RJ (1951) *J Biol Chem* 193:265–275
- Mitchell P (1979) *Science* 206:1148–1159
- Smith RM, Martell AE (1976) In: Smith RM, Martell AE (eds) *Critical stability constants*, vol 4. Plenum, New York, p 54
- Toninello A, Clari G, Mancon M, Tognon G, Zatta P (2000) *J Biol Inorg Chem* 5:612–623
- Zoratti M, Szabò I, De Marchi U (2005) *Biochim Biophys Acta* 1706:40–52
- Kaufmann T, Schinzel A, Borner C (2004) *Trends Cell Biol* 14:8–12
- Desagher S, Martinou JC (2000) *Trends Cell Biol* 10:369–377
- Sharpe JC, Arnoult D, Youle RJ (2004) *Biochim Biophys Acta* 1644:107–113
- Campello S, De Marchi U, Szabo I, Tombola F, Martinou JC, Zoratti M (2005) *FEBS Lett* 579:3695–3700
- De Marchi U, Campello S, Szabo I, Tombola F, Martinou JC, Zoratti M (2004) *J Biol Chem* 279: 37415–37422
- Li M, Xia T, Jiang CS, Li LJ, Fu JL, Zhou ZC (2003) *Toxicology* 194:19–33
- Jang D, Sullivan PG, Sensi SL, Steward O, Weiss JH (2001) *J Biol Chem* 276:47524–47529
- He L, Poblens AT, Medrano CJ, Fox DA (2000) *J Biol Chem* 275:12175–12184
- Pourahmad J, Mihailovic A, O’Brien PJ (2001) *Adv Exp Med Biol* 500:249–252
- Belyaeva EA, Glazunov VV, Nikitina ER, Korotkov SM (2001) *J Bioenerg Biomembr* 33:303–318
- Dorta DJ, Leite S, De Marco KC, Prado IM, Rodriguez T, Mingatto FE, Uyemura SA, Santos AC, Curti C (2003) *J Inorg Biochem* 97:251–257
- Belyaeva EA, Glazunov VV, Korotkov SM (2004) *Acta Biochim Pol* 51:19–33
- McStay GP, Clarke S, Halestrap A (2002) *Biochem J* 367:541–548