

Purification and characterization of 7 β -hydroxysteroid dehydrogenase from rabbit liver microsomes

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Abstract

7 β -Hydroxysteroid dehydrogenase (7 β -HSD), a specific enzyme active in the metabolization of 7 β -hydroxycholesterol, was purified about 300-fold from male rabbit liver microsomes using ion exchange, hydroxylapatite, 2'5'ADP Sepharose 4B, and high-performance liquid chromatography on the basis of its catalytic activity.

The specific activity of the purified enzyme was 276 nmol/min/mg protein. The molecular weight of the purified enzyme was 34,000. The preferred coenzyme was β -NADP⁺. The optimum pH for oxidation was around 7.7 in potassium phosphate buffer, and 11.0 in glycine–NaOH buffer. The purified enzyme catalyzed the synthesis of not only 7 β -hydroxycholesterol but also corticosterone and hydrocortisone. Enzyme activities toward these three substrates accompanied all purification steps of 7 β -HSD. The amino acid sequence of the N-terminal of the purified enzyme showed that 7 β -HSD had sequence similarity to rabbit type I 11 β -hydroxysteroid dehydrogenase (11 β -HSD), indicating that 7 β -HSD may belong to the rabbit type I 11 β -HSD family and may play the same role in the metabolism of 11-hydroxysteroids and 7-hydroxysterols.

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1. Introduction

7-Oxycholesterol is thought to be an autooxidation product of cholesterol in mammals [1,2]. This sterol is found in human aorta, brain, kidney, liver and serum. Recently, many effects of 7-oxycholesterol in experimental animals and humans have been demonstrated [3–7]. However, its biological importance is still under investigation.

7 α - and 7 β -hydroxycholesterol are products of cholesterol. 7 α -Hydroxycholesterol is an intermediate of the bile acid biosynthetic pathway and metabolized into bile acids [8–10]. On the other hand, the role and metabolic fate of 7 β -hydroxycholesterol are not well defined. Björkhem et al. [11] have preliminarily reported the presence of 7 β -hydroxysteroid dehydrogenase (7 β -HSD) in a 20,000 g supernatant of rat liver that converted 7 β -hydroxycholesterol to 7-ketocholesterol. Song et al. [12,13] reported purification of an enzyme from hamster liver microsomes that converts 7 α -hydroxycholesterol into 7-ketocholesterol. They called the enzyme 7 α -hydroxycholesterol dehydrogenase and reported that the purified enzyme catalyzed not only

7 α -hydroxycholesterol but also 7 β -hydroxycholesterol, corticosterone and hydrocortisone with almost the same catalytic activity. They also reported that a partial N-terminal amino acid sequencing showed that this enzyme had a sequence similar to that of human 11 β -hydroxysteroid dehydrogenase (11 β -HSD). Recently, we reported a comparative study of the conversion of 7 α - and 7 β -hydroxycholesterol in various animals and demonstrated species differences and possibility of interconversion between these sterols [14]. While 7 α -hydroxysteroid dehydrogenase (7 α -HSD) was observed in hamster and chicken, 7 β -HSD was observed in all vertebrates tested.

To clarify further the biological importance of the 7 β -dehydrogenation of 7 β -hydroxycholesterol, we here describe the purification and characterization of the enzyme from rabbit liver microsomes.

2. Materials and methods

2.1. Materials

7 β -Hydroxycholesterol and 7-ketocholesterol were purchased from Steraloid (Wilton, NH). 7 α -Hydroxy-4-cho-

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lesten-3-one was synthesized according to the method described by Shimasue [15]. DEAE Sepharose and 2/5' ADP Sepharose 4B were purchased from Pharmacia (Uppsala, Sweden). Hexane, isopropanol and methanol of high-performance liquid chromatography (HPLC) grade, β -NADP⁺ and glycerol were purchased from Wako Pure Chemical (Osaka, Japan). Other reagents were of the highest grade commercially available. Hydroxylapatite was obtained from Bio-Rad (Hercules, CA).

2.2. Purification of the enzyme

A male New Zealand white rabbit weighing 2.0–2.5 kg was sacrificed under diethyl ether anesthesia. The liver was quickly removed, perfused with 0.9% NaCl and then homogenized with 4 volumes of 0.25 M sucrose containing 5 mM Tris-HCl and 1 mM EDTA, pH 7.4. The homogenate was centrifuged at $9000 \times g$ for 15 min in the cold. The supernatant was centrifuged at $105,000 \times g$ for 60 min. The precipitate was washed and recentrifuged at $105,000 \times g$ for 60 min.

The microsomal precipitate was suspended in 100 mM Tris-HCl buffer, pH 7.4, containing 20% glycerol, 1 mM EDTA, and 0.5 mM dithiothreitol (DTT). The following procedures were carried out at 4 °C except for HPLC, which was performed at room temperature. All buffer solutions used in the following purification procedures contain 20% glycerol, 1 mM EDTA, and 0.5 mM DTT unless otherwise stated.

The microsomal pellet was solubilized with 1% (w/v) sodium deoxycholate. The solubilized preparation was fractionated with 50% polyethylene glycol (PEG). A fraction precipitated with 8%–16% PEG was re-dissolved in the 100 mM Tris-HCl buffer (pH 7.4), containing 0.5% sodium deoxycholate. The re-dissolved solution was dialyzed against 100 mM Tris-HCl (pH 7.4) containing 0.2% sodium deoxycholate. The dialysate was subjected to Phenyl Sepharose CL4B column chromatography (2.5 cm \times 10 cm). The column was washed with the same buffer and eluted with 100 mM Tris-HCl (pH 7.4), containing 1% Lubrol PX. The fractions showing enzyme activity were collected and dialyzed against 20 mM Tris-HCl (pH 8.0) containing 0.02% Lubrol PX. The dialysate was subjected to a DEAE Sepharose CL6B column (2.5 cm \times 5 cm) equilibrated with the same buffer used for dialysis. The column was washed with the equilibration buffer and eluted with a linear gradient of 0–0.5M sodium chloride in the equilibrated buffer. The fraction showing enzyme activity were pooled and dialyzed against 20 mM potassium phosphate buffer (pH 7.4) containing 0.02% Lubrol PX. The dialysate was applied to a hydroxylapatite column (2.5 cm \times 2.0 cm) equilibrated with the same buffer used for dialysis and a stepwise elution was performed with 40, 100, 200 and 400 mM potassium phosphate buffer (pH 7.4). The fraction showing enzyme activity was pooled and dialyzed against 20 mM potassium phosphate buffer (pH 7.4), con-

taining 0.02% Lubrol PX. The dialysate was subjected to column chromatography (1.5 cm \times 1.5 cm). The column was washed with the equilibration buffer and eluted with 2 M sodium chloride in the equilibrated buffer. The fraction with the enzyme activity was collected and dialyzed against 20 mM Tris-HCl (pH 8.0) containing 0.4% Lubrol PX. Aliquots were injected into a TSK-gel DEAE-5PW column (0.75 cm \times 7.5 cm) equilibrated with the same buffer used for dialysis. The column was eluted with a linear gradient of 0–0.5M sodium chloride in the equilibrated buffer. The flow rate of the anion exchange column chromatography was 0.5 ml/min, and effluents were monitored at 280 nm. The fractions with enzyme activity were collected and concentrated with a centricon. After concentration, the samples were stored at –20 °C.

The research protocol was approved by the animal study committees at Miyazaki Medical College (2002–16).

2.3. Enzyme assay

7 β -Hydroxysteroid dehydrogenase activity was measured as described previously [14]. A typical assay mixture contained 0.1 M of potassium phosphate buffer (pH 7.4), 1 mM EDTA, 0.5 μ mol of NADP⁺ and 20 nmol of 7 β -hydroxycholesterol dissolved in 10 μ l of isopropanol in a final volume of 1.0 ml. Incubation was carried out at 37 °C for 5 min and stopped by addition of 1 ml of methanol. Steroids were extracted with 5 ml of *n*-hexane. An aliquot of the extract was evaporated, dissolved in isopropanol and injected into a high-performance liquid chromatograph produced by Shimadzu (Kyoto, Japan, Model LC-8A) and equipped with an integration system (Shimadzu; Model Chromatopack C-R3A). The HPLC column (Wakosil 5SIL, 4.6 mm \times 250 mm) was obtained from Wako Pure Chemical Industries.

Oxidation enzyme activities at C-11 of hydrocortisone and corticosterone were determined by measuring the rate of conversion in the presence of β -NADP⁺. A typical incubation mixture contained 0.1 M of potassium phosphate buffer (pH 7.4), 1 mM EDTA, 0.5 μ mol of β -NADP⁺ and 20 nmol of hydrocortisone or corticosterone dissolved in 10 μ l of methanol in a final volume of 1.0 ml. Incubation was carried out at 37 °C for 5 min and steroids were extracted with 5 ml of ethyl acetate. An aliquot of the extract was evaporated, dissolved in methanol and injected into the HPLC. The column used was a C18 column (Wako I, 4.6 mm \times 150 mm), and the solvent mixture was methanol/water (50/50, v/v). Effluents were monitored at 240 nm.

2.4. Other methods

Protein concentration was determined by the method of Lowry et al. [16]. SDS-polyacrylamide gel electrophoresis (SDS-PAGE) was performed according to the method of Laemmli [17], and proteins were visualized by silver staining.

2.5. Amino acid sequence analysis

Partial N-terminal amino acid sequence analysis was performed. The purified enzyme was subjected to SDS-PAGE and transferred to a polyvinylidene difluoride membrane using transfer buffer, containing 10 mM CAPS and 10% (v/v) methanol. Transferred protein was stained with 0.02% Coomassie brilliant blue. The band of protein was cut out and sequenced using an Hewlett–Packard G1005A Protein Sequencer System.

3. Results

Table 1 summarizes the purification of 7 β -HSD from male rabbit liver microsomes. The activity of the enzyme was decreased significantly after solubilization of microsomes with sodium cholate (data not shown). On the other hand, a significantly increased activity (approximately three times that of

the microsomal fraction) was observed when sodium deoxycholate was used as the detergent. Over 60% of the total enzyme activity was retained in the fraction eluted between 8% and 16% of PEG 6000. The enzyme activity was observed in the non-bound fractions of the DEAE chromatography. The fractions eluted by 200 and 400 mM of potassium phosphate buffer (pH 7.4) had high enzyme activities. The fractions of a HA column eluted with 400 mM potassium phosphate buffer were applied to a 2'5' ADP Sepharose 4B column. The enzyme was eluted with the 20 mM potassium phosphate buffer (pH 7.4), containing 2 M NaCl. The enzyme was further purified using HPLC equipped with an anion-exchange column (DEAE-5PW).

SDS-polyacrylamide gel electrophoretograms of the fractions separated by HPLC are shown in Fig. 1. The molecular weight of the purified enzyme from rabbits was 34,000. Table 2 shows the substrate specificity of the purified enzyme. Cholesterol, 7 α -hydroxycholesterol, 25-hydroxycholesterol, pregnenolone, dehydroepiandrosterone,

Table 1
Purification steps for rabbit liver microsomal 7 β -hydroxysteroid dehydrogenase

Fraction	Total protein (mg)	Total activity (nmol/min)	Yield (%)	Specific activity (nmol/min/mg protein)
Microsome	2520	2368.8	100	0.94
Solubilization	2544	7248.5	306.6	2.885
PEG (8%–16%)	936	5083.4	214.6	5.43
Phenyl Sepharose				
Non-bound	203.4	2359.3	99.6	11.60
DEAE				
Non-bound	82.7	2096.4	88.5	25.3
HA				
Elution with 400 mM	6.2	824.6	34.8	133.0
HPLC (DEAE 5PW)	1.7	469.4	19.8	276.1

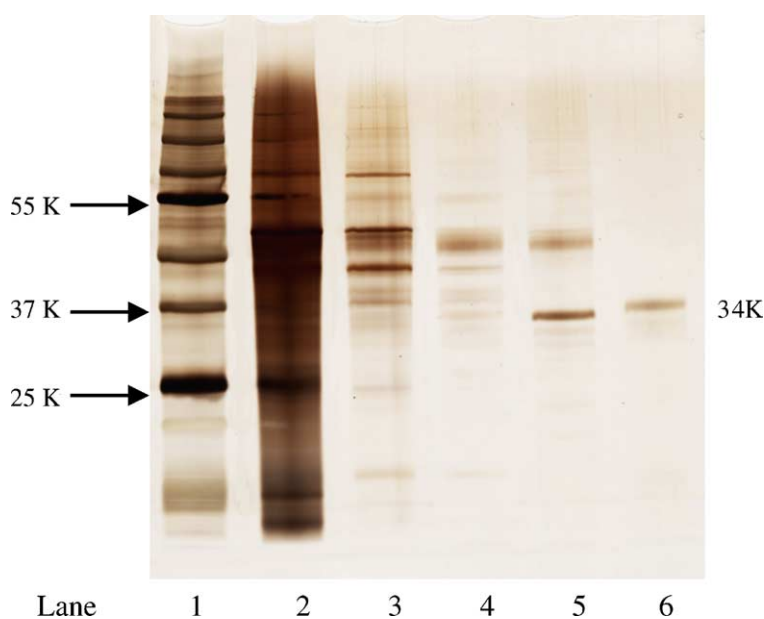


Fig. 1. SDS-PAGE of the purification steps of rabbit liver microsomal 7 β -HSD. Lane 1: standard; lane 2: protein profile of rabbit liver microsomes; lane 3: PEG 8%–16% fraction; lane 4: after DEAE; lane 5: after HA; lane 6: after DEAE 5PW. Protein bands were visualized by silver staining.

Table 2
Substrate specificity of 7 β -hydroxysteroid dehydrogenase

Substrate	Product formed (nmol of product/min/mg protein)
5-Cholesten-3 β -ol	ND ^a
5-Cholesten-3 β ,7 α -diol	ND
5-Cholesten-3 β ,7 β -diol	267.9
3 α ,7 α -Dihydroxycholanoic acid	ND
3 α ,7 β -Dihydroxycholanoic acid	ND
Cortisol	534.4
Corticosterone	477.7

^a Not detected.

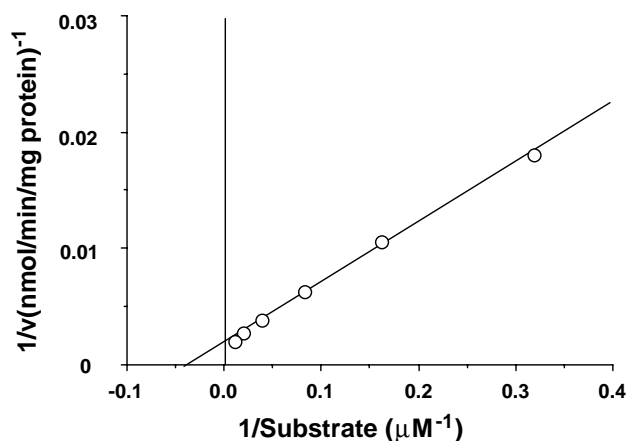


Fig. 2. Lineweaver–Burk plot. One microgram of purified enzyme was incubated with various amounts of substrate and the products were determined by HPLC as described in Section 2.

Table 3
N-terminal amino acid sequence of purified rabbit hepatic 7 β -HSD

Rabbit 7 β -HSD	AFMKKYLPLLLGLFLAYYY
Rabbit 11 β -HSD	AFMKKYLPLLLGLFLAYYYYSA (Ozols, 1995)

chenodoxycholic acid and ursodeoxycholic acid were not oxidized by the purified enzyme. A Lineweaver-Burk plot is shown in Fig. 2. The V_{\max} and K_m were 512.8 nmol/min/mg of protein and 2.63 μ M, respectively. Enzyme activities of each purification step are shown in Fig. 3. Enzyme activities toward the three substrates accompanied each other at all purification steps.

The amino acid sequence of the N-terminal of the purified enzyme is shown in Table 3. As shown in the Table 3, the sequences of the enzyme were similar to those of the rabbit type I 11 β -hydroxysteroid dehydrogenase reported.

4. Discussion

7 β -Hydroxysteroid dehydrogenase is a hepatic microsomal enzyme that catalyzes the conversion of 7 β -hydroxycholesterol to 7-ketocholesterol. This reaction has been described in the literature since 1968 [11], but details of

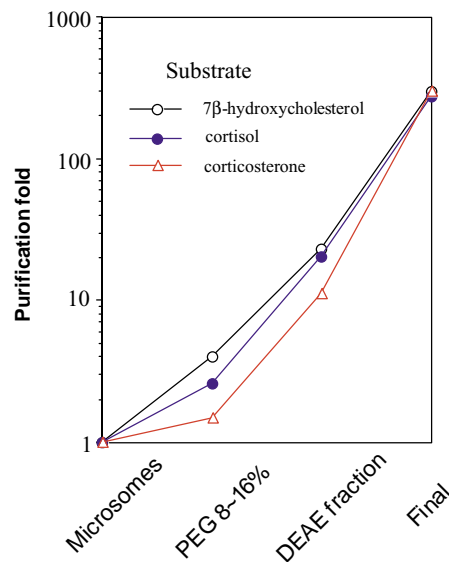


Fig. 3. Yield of each purification step. Proteins of each step were incubated with three different substrates and determined by HPLC as described in Section 2.

the enzyme have not been investigated to date. Recently, we studied 7-hydroxysteroid dehydration in several animals and demonstrated that 7 β -dehydration occurs in many vertebrates, including humans [14]. On the other hand, 7 α -dehydration occurs in hamster and chicken liver microsomes. In 1998, Song et al. [13] reported the purification of an enzyme from hamster liver microsomes that catalyzed the conversion of 7 α -hydroxycholesterol to 7-ketocholesterol and that the apparent molecular weight of the purified enzyme was 34,000. They described the enzyme as a 7 α -hydroxycholesterol dehydrogenase and demonstrated that the purified enzyme catalyzed the conversion of not only 7 α -hydroxycholesterol but also 7 β -hydroxycholesterol. On the other hand, according to our results, microsomes prepared from rabbit, guinea pig and rat catalyzed the conversion of 7 β -hydroxycholesterol, but not 7 α -hydroxycholesterol. From their reports regarding the partial N-terminal amino acid sequence [13], 7 α -hydroxycholesterol dehydrogenase has a sequence similar to that of human 11 β -HSD.

In this experiment, we purified rabbit liver microsomes and demonstrated that the purified enzyme catalyzed the formation of 7 β -hydroxycholesterol, hydrocortisone and corticosterone but not 7 α -hydroxycholesterol. The result suggests a species difference between rabbit and hamster in the substrate specificity of this enzyme and also suggests that the purified enzyme is different from the enzymes that catalyze the oxidoreduction of bile acids [18,19]. During the enzyme purification procedure, we determined that the specific enzyme activities toward these three substrates (7 β -hydroxycholesterol, hydrocortisone and hydroxycorticosterone) were elevated simultaneously. In the present study, we did not test conversion between 7 α -

7 β -dehydroepiandrosterone (DHEA) and 7-oxo-DHEA. However, recently, Robinzon et al. reported that interconversion of 7-oxo-DHEA and 7-OH-DHEA isomers is catalyzed by 11 β -HSDI in both liver and kidney [20]. A partial N-terminal amino acid sequence of 7 β -HSD was similar to that of rabbit type I 11 β -HSD. These results suggest that rabbit 7 β -HSD may belong to the family of rabbit type I 11 β -HSD. To clarify its clinical importance, further study (for example, serum or tissue concentrations of those steroids) is needed.

11 β -Hydroxysteroid dehydrogenase is a microsomal enzyme that is widely distributed in various tissues and responsible for the oxidation of 11-hydroxy- to 11-ketosteroids. Two tissue-specific isoforms with different functions have been described [21]. Type I 11 β -HSD is widely distributed, with high levels in liver and kidney [22,23], while type II 11 β -HSD is mainly present in kidney and placenta [24–26]. Recently, it was shown that the type I 11 β -HSD plays an important role in xenobiotic carbonyl reduction [27–29]. In our experiment, reduction of 7 β -hydroxycholesterol was demonstrated. What is this microsomal 7 β -HSD and how does it function physiologically? Corticosterone and hydrocortisone are important endogenous hormones essential for water and electrolytes metabolism. 7 β -Hydroxycholesterol is not a very active metabolite in the body, but it is a member of the oxysterols. Therefore, it should be clarified which 7-oxocholesterol (7-ketocholesterol or 7 β -hydroxycholesterol) is the dominant product and which is less toxic (or harmful) in the lipid metabolism in the body.

In conclusion, we purified 7 β -hydroxysteroid dehydrogenase from rabbit liver microsomes and demonstrated an amino acid sequence similarity to that of the rabbit type I 11 β -hydroxysteroid dehydrogenase.

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