

Bile acid *N*-acetylglucosaminides

Formation by microsomal *N*-acetylglucosaminyltransferases in human liver and kidney

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Bile acid *N*-acetylglucosaminyltransferase activity has been identified in microsomes from human liver and kidney. In both organs the transferases required UDP-*N*-acetylglucosamine as sugar donor and were mainly active towards ursodeoxycholic acid. Minor activities were observed towards amidated ursodeoxycholic, hyodeoxycholic and β -muricholic acids. No *N*-acetylglucosaminidation was detectable with the major primary and secondary bile acids suggesting a specific requirement of the enzymes for bile acids containing 7 β - or 6 α -hydroxyl groups. Kinetic parameters and other catalytic properties of liver and kidney microsomal *N*-acetylglucosaminyltransferase activities towards ursodeoxycholic acid are described.

Glycosidic conjugation; Bile acid; Glycosyltransferase

1. INTRODUCTION

N-Acetylglucosaminides of bile acids were recently identified in normal human urine [1]. The daily excretion of these novel bile acid conjugates was comparable to the urinary daily excretion of bile acid glucosides [1,2] and bile acid glucuronides [3]. Whereas characteristics of the biosynthesis of bile acid glucosides [4,5] and glucuronides [6–9] could be elucidated from studies of the corresponding glycosyltransferases in man, nothing was known on the enzyme(s) catalyzing bile acid *N*-acetylglucosaminidation. The present report describes a rapid and sensitive assay procedure which enabled the characterization of the biosynthesis of bile acid *N*-acetylglucosaminides.

2. MATERIALS AND METHODS

Sources of chemicals were the same as described in previous papers [4,6]. UDP-*N*-acetyl-D-glucosamine [glucosamine- 14 C (U)], 284 mCi/mmol, and UDP-*N*-acetyl-D-glucosamine [glucosamine-6- 3 H (N)], 19 Ci/mmol were obtained from NEN chemicals, Dreieich. Human tissue specimens were obtained from a similar collective of surgical patients or organ donors as described [5]. Kidney samples were restricted to the cortical region. Only tissues with apparently normal histology were used.

Human tissue specimens could be stored for maximally 4 weeks at -30°C without a marked loss of bile acid *N*-acetylglucosaminyltransferase activity. Homogenization of tissues and preparation of microsomes by differential centrifugation was performed as described in a previous report [6].

Unless otherwise stated, bile acid *N*-acetylglucosaminyltransferase

activity in human kidney or liver towards ursodeoxycholic acid was determined as follows: approximately 10–30 μg of microsomal protein was incubated in a reaction mixture which contained in a total volume of 30 μl 0.1 mM ursodeoxycholic acid, 50 μM UDP-*N*-acetyl-D- 14 H]glucosamine (0.05 μCi), 0.05 mM ADP for inhibition of nucleotide pyrophosphatase and the following additions for incubations with renal or hepatic microsomes, respectively: 0.1 M sodium acetate (pH 6.2), 2 mM MgCl_2 and 0.002% (w/v) Brij 58 with renal microsomes; 0.1 M 4-morpholineethanesulfonic acid (Mes)-KOH (pH 6.4) and 1 mM MgCl_2 with hepatic microsomes. ADP (0.05 mM) had no inhibitory effect on bile acid *N*-acetylglucosaminidation and maintained UDP-*N*-acetylglucosamine levels in 15-min incubations at about 95% and 99% of the initial values as compared to 77% and 93% without ADP in kidney and liver microsomes, respectively, as determined by thin layer chromatography (cf. below). After 15 min at 37°C the reactions were terminated with 30 μl 0.7 M Glycine-HCl (pH 2.7) containing 0.8 M sodium chloride. The water phase was extracted for 5 min with 300 μl water-saturated ethyl acetate resulting in transfer of about 95% of the reaction product into the organic phase. The unreacted UDP-*N*-acetylglucosamine remained in the water phase (> 99.9%). A 200 μl aliquot of the organic phase was counted for radioactivity in 5 ml Optifluor (Packard) scintillation cocktail.

For identification of bile acid *N*-acetylglucosaminides the reaction products were purified on a Sep-Pak C_{18} cartridge (see legend to Table I) and were incubated with or without *N*-acetylglucosaminidase as described [1]. Ethanol extracts of these incubation mixtures were applied to silica gel 60 thin-layer plates which were developed with *n*-butanol/acetic acid/water (50/5/10, v/v) and analyzed in a thin-layer chromatogram scanner (Berthold, Wildbad). When incubations were performed in the absence of *N*-acetylglucosaminidase single spots of the following *N*-acetylglucosaminides could be detected: *N*-acetylglucosaminides of ursodeoxycholic, R_F 0.609; glycooursodeoxycholic, R_F 0.235; tauroursodeoxycholic, R_F 0.158; hyodeoxycholic, R_F 0.591 and β -muricholic acids, R_F 0.535, respectively. After incubation of reaction products in the presence of *N*-acetylglucosaminidase these spots were shifted to a position identical to that of authentic *N*-acetyl- 14 C]glucosamine (R_F 0.266).

Protein was determined as described in a previous report [9].

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3. RESULTS AND DISCUSSION

3.1. Substrate specificity and tissue distribution

Analysis of reaction mixtures after incubation of renal or hepatic microsomes with various bile acids as acceptor substrates and UDP-*N*-acetylglucosamine as donor substrate showed that formation of *N*-acetylglucosaminides mainly occurred in the presence of ursodeoxycholic acid (Table I). Amidation of ursodeoxycholic acid with glycine or taurine, change of the position and configuration of the hydroxyl group from 7 β to 6 α (hyodeoxycholic acid versus ursodeoxycholic acid) or the presence of a third hydroxyl group in the bile acid molecule as in β -muricholic acid led to a marked decrease in the rate of *N*-acetylglucosaminidation (Table I). No reaction was detectable (< 1 pmol/min/mg protein) with the primary bile acids cholic and chenodeoxycholic acids, the major secondary bile acids lithocholic and deoxycholic acids and the following other bile acids: 3 α , 6 β -di-hydroxy-5 β -cholanoic, ursocholic (3 α , 7 β , 12 α -trihydroxy-5 β -cholanoic) and hyocholic (3 α , 6 α , 7 α -trihydroxy-5 β -cholanoic) acids. These results suggest a specificity of the enzymes from liver and kidney towards the 7 β , or 6 α -hydroxyl group of preferentially dihydroxy as compared to trihydroxy bile acids. The restricted bile acid specificity of *N*-acetylglucosaminyltransferases in vitro assays is in accordance with in vivo findings since ursodeoxycholic acid was the only bile acid that could be identified as *N*-acetylglucosaminide from human urine [1].

In contrast to the activities observed in kidney and liver (Table I), no *N*-acetylglucosaminyl transfer from UDP-*N*-acetylglucosamine (50 μ M or 6 μ M) to ursodeoxycholic acid (0.1 mM) was detectable with

homogenate or microsomes from different segments of human intestinal mucosa (ileum, colon, sigmoid and rectum).

3.2. Characteristics of *N*-acetylglucosaminyltransferases in liver and kidney

The influence of pH on the formation of ursodeoxycholic acid *N*-acetylglucosaminide was similar for the enzymes from liver and kidney. Enzyme activities were optimal between pH 6.1–6.4 in renal microsomes and pH 6.1–6.7 in hepatic microsomes and showed a steep decline of activity towards pH 5.0 and 8.0 (not shown).

Divalent metal ions exhibited a weak stimulatory effect on enzyme activities towards ursodeoxycholic acid. Compared to the control without metal ion and containing 10 mM EDTA, the kidney enzyme was maximally stimulated to about 140% in the presence of 2 mM Mg²⁺, 2 mM Mn²⁺ or 10 mM Ca²⁺. The liver enzyme showed maximal activation to about 160% in the presence of 1 mM Mg²⁺, 0.1 mM Mn²⁺ or 0.5 mM Ca²⁺. In contrast to the enzyme from renal microsomes, the metal ion dependence of the liver enzyme was only observed at a low concentration of UDP-*N*-acetylglucosamine (6 μ M). No influence of these metal ions on hepatic enzyme activity was detectable at 50 μ M UDP-*N*-acetylglucosamine. Zn²⁺ was not stimulatory but inhibitory to the enzymes from liver and kidney in concentrations >0.01 mM.

Since membrane-bound enzymes may be activated by the addition of detergents [11] the influence of the non-ionic detergents Brij 58 and Triton X-100 (0.001–0.2%, w/v or v/v, respectively) was studied on ursodeoxycholic acid *N*-acetylglucosaminidation in liver and kidney microsomes. Whereas the effect of Triton X-100 on both hepatic and renal enzymes and Brij 58 on the liver enzyme was not activatory but inhibitory with increasing concentration ($> 0.002\%$), Brij 58 led to a 1.5-fold activation of the renal enzyme at a concentration of 0.002%.

Double-reciprocal plots of *N*-acetylglucosaminyltransferase activities towards ursodeoxycholic acid with UDP-*N*-acetylglucosamine as varied substrate yielded a straight line with renal microsomes whereas with hepatic microsomes plots were concave downwards (Fig. 1). The same curved double-reciprocal plots were obtained with liver microsomes in the presence or absence of the detergent Brij 58 (0.002%, w/v). The data in Fig. 1 indicate that rates of ursodeoxycholic acid *N*-acetylglucosaminidation in hepatic microsomes are higher at low concentrations of UDP-*N*-acetylglucosamine than calculated from extrapolation of the plots at high concentrations of the sugar donor. Therefore, destruction of UDP-*N*-acetylglucosamine in other metabolic microsomal pathways cannot explain these results. Whether or not nonlinearity of double-reciprocal plots could be explained by a negatively cooperative binding mechanism [12] or UDP-*N*-

Table I

Bile acid substrate specificity of <i>N</i> -acetylglucosaminyltransferases		
Bile acid	Relative activity	
	Kidney (%)	Liver (%)
Ursodeoxycholic acid (3 α , 7 β)	100 ^a	61
Glyoursodeoxycholic acid (3 α , 7 β)	19	3.1
Taoursodeoxycholic acid (3 α , 7 β)	35	12
Hyodeoxycholic acid (3 α , 6 α)	9.4	13
β -Muricholic acid (3 α , 6 β , 7 β)	7.4	4.0

Enzyme activities were determined as described in Materials and Methods at bile acid concentrations of 0.1 mM with the following modifications: microsomal protein (100 μ g) was incubated in a total volume of 180 μ l; the sugar donor was UDP-*N*-acetyl-D-[¹⁴C]glucosamine (6 μ M, 0.3 μ Ci); the incubations were terminated with 800 μ l ethanol and protein was removed by centrifugation; the reaction products were purified from the supernatants by adsorption to a Sep-Pak C₁₈ cartridge [10] and analyzed by thin-layer chromatography as described in Materials and Methods. Position and configuration of hydroxyl groups in the bile acid skeleton are given in parentheses.

^a Definition of 100% value: 13.5 pmol of product formed per min per mg of protein

Table II

Kinetic constants of bile acid-*N*-acetylglucosaminyltransferases in kidney and liver

Kinetic constants	Kidney	Liver	
		Low UDPGNac ^a	High UDPGNac ^a
K_m for ursodeoxycholic acid	93.0 ± 23.5	27.4 ± 5.3	31.8 ± 8.9
K_m for UDP- <i>N</i> -acetylglucosamine	409.8 ± 138.9	2.2 ± 1.3	51.2 ± 26.2
V_{max}	295.6 ± 95.5	1.0 ± 0.5	3.9 ± 0.8

Values are given as mean ± SD which were obtained from the following number of organ samples: kidney, $n = 5$; liver, $n = 4$. The kinetic parameters for the kidney enzyme were calculated according to Florini and Vestling [13] from double-reciprocal plots of initial rates as a function of the concentration of ursodeoxycholic acid (0.01–0.1 mM) at various fixed concentrations of UDP-*N*-acetylglucosamine (25 μM, 35 μM, 50 μM, 100 μM) yielding an intersecting pattern of straight lines. The kinetic data for the liver enzyme were determined from Lineweaver-Burk plots as follows: with ursodeoxycholic acid (0.01–0.1 mM) as variable parameter at either low (6 μM) or high (50 μM) concentration of UDP-*N*-acetylglucosamine yielding the K_m -values for ursodeoxycholic acid; with UDP-*N*-acetylglucosamine as variable parameter by extrapolating the regions in double reciprocal plots tending towards linearity at low or high concentrations of UDP-*N*-acetylglucosamine (see Fig. 1) yielding the K_m -values for UDP-*N*-acetylglucosamine and the V_{max} -values. ^aUDPGNac, UDP-*N*-acetylglucosamine; K_m , μM; V_{max} , pmol/min/mg protein.

acetylglucosamine to the liver enzyme or by the existence of two hepatic isoenzymes for the synthesis of ursodeoxycholic acid *N*-acetylglucosaminide should be analyzed with a purified enzyme preparation and not with a heterogenous microsomal fraction.

The apparent kinetic constants for microsomal *N*-acetylglucosaminyltransferase activities from kidney and liver towards ursodeoxycholic acid are summarized in Table II. Due to the nonlinearity of double-reciprocal plots with liver *N*-acetylglucosaminyltransferase (Fig. 1) the kinetic parameters for this enzyme were calculated at low and high concentrations of UDP-*N*-acetylglucosamine (see legend to Table II). As may be seen from Table II the enzymes from kidney and

liver show marked differences in kinetic parameters especially with respect to the K_m -values for UDP-*N*-acetylglucosamine and the V_{max} -values. From these kinetic parameters, however, a similar efficiency of ursodeoxycholic acid *N*-acetylglucosaminidation in kidney and liver can be calculated based on the ratio of the V_{max} - to K_m -values for UDP-*N*-acetylglucosamine at saturating concentrations of ursodeoxycholic acid: 0.7 for the renal enzyme and 0.45 for the hepatic enzyme at low concentrations of UDP-*N*-acetylglucosamine.

The results of the present study show that the occurrence of bile acid *N*-acetylglucosaminides in urine of man [1] can be explained by the existence of bile acid *N*-acetylglucosaminyltransferases in liver and kidney. The bile acid specificity of these enzymes is interesting because ursodeoxycholic acid was the preferred substrate in *N*-acetylglucosamine conjugation, and this bile acid was recently shown to improve liver function in patients with cholestatic liver disease [14]. Further studies are needed to elucidate whether or not a correlation exists between *N*-acetylglucosaminidation and the pathophysiological effects of ursodeoxycholic acid in liver disease.

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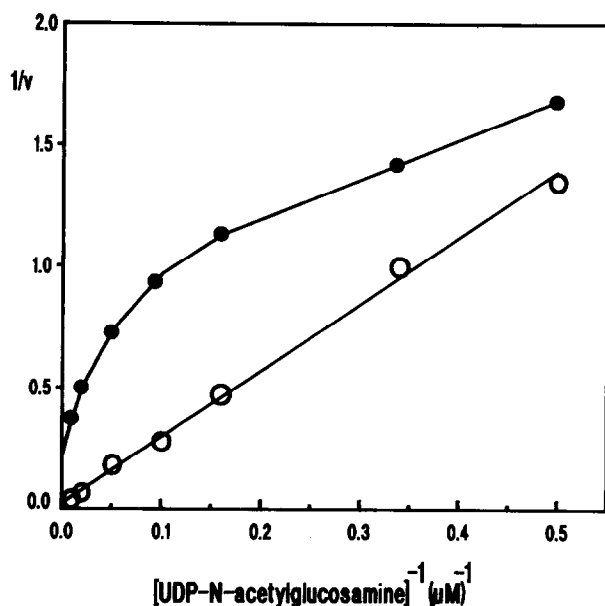


Fig. 1. Double reciprocal plots of initial rates of *N*-acetylglucosaminidation as a function of the concentration of UDP-*N*-acetylglucosamine at an ursodeoxycholic acid concentration of 0.1 mM with microsomes from liver (●) and kidney (○). The depicted plots show typical results from a single microsomal preparation of each organ; v , pmol/min/mg protein.

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