

Pharmacokinetics of bambuterol in subjects homozygous for the atypical gene for plasma cholinesterase

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Aims It has been assumed that both plasma cholinesterase (EC 3.1.1.8) and oxidative enzymes are needed for optimum formation of the bronchodilator terbutaline from its biscarbamate prodrug bambuterol. The present study aimed at investigating the fate of bambuterol in subjects with deficient plasma cholinesterase but with normal oxidative (CYP2D6) capability.

Methods The pharmacokinetics of bambuterol and terbutaline were studied in four healthy subjects (two men and two women) being homozygous for the atypical gene for plasma cholinesterase. Their oxidative metabolism was apparently good as they were all rapid metabolizers of debrisoquine. Bambuterol hydrochloride 20 mg was given orally once daily for 10 days, and plasma and urine samples were taken for 1.5 days (plasma) and 4.5 days (urine) after administration of the last dose.

Results The pharmacokinetic parameters in the present study were grossly similar to those found in a study of bambuterol in subjects with normal plasma cholinesterase activity (N). However, subjects with atypical cholinesterase had a shorter terminal half-life of bambuterol (a measure of uptake rate), 4.8–12.6 h *vs* 8.3–22.3 h in N, and slightly higher plasma concentrations of bambuterol (average concentrations 1.9–3.7 nmol l⁻¹ *vs* 1.5–3.1 nmol l⁻¹ in N). Peak/trough terbutaline plasma concentrations ratios (2.1–3.2) were somewhat increased, but average plasma concentrations (8.3–14.5 nmol l⁻¹) and terminal half-life (16.5–21.8 h) of terbutaline did not differ.

Conclusions In Caucasian populations, one subject out of 2500 is homozygous for the atypical gene for plasma cholinesterase. The atypical enzyme has a much lower affinity for bambuterol than the normal enzyme. Nevertheless, the subjects with atypical cholinesterase were able to produce terbutaline as efficiently as normal subjects. This might be explained by an altered uptake and metabolism in the absence of plasma cholinesterase, or the importance of this enzyme for the formation of terbutaline from bambuterol *in vivo* may have been overestimated.

Keywords: Bambuterol, terbutaline, pharmacokinetics, atypical plasma cholinesterase

Introduction

Bambuterol is the inactive bis-*N,N*-dimethylcarbamate prodrug to the β_2 -adrenoceptor agonist terbutaline. The active moiety is formed through hydrolytic and/or oxidative pathways (Figure 1). The hydrolytic reactions are catalysed mainly by plasma cholinesterase (EC 3.1.1.8) and the oxidative reactions most likely by cytochrome P450 (CYP) enzymes [1].

Plasma cholinesterase (pChE) is produced by the liver and exhibits polymorphism [2]. The activity of pChE may be reduced for several reasons such as physiological variation, disease, drugs, and genetic factors. This does not seem to interfere with normal physiology but may influence the duration of action of substances dependent on pChE for their elimination, such as suxamethonium and mivacurium [2–6].

Bambuterol has a very high affinity for pChE from

subjects with normal genotype (UU) or from those heterozygous for the atypical and the normal gene (UA). Much lower affinity of bambuterol is seen towards the plasma cholinesterase produced in subjects homozygous for the atypical gene (AA) and hydrolysis of bambuterol is slow in plasma from these subjects [7]. Homozygotes for the atypical gene are rare in Caucasian populations, the frequency being about 1/2500 in the Danish population [3].

In healthy subjects with normal cholinesterase genotype, the administration of a normal oral dose (about 20 mg) of bambuterol hydrochloride once daily resulted in sustained plasma concentration profiles of terbutaline with a mean peak/trough ratio of slightly less than 2.0 [8]. The pChE activities just before dosing were between 55 and 77% (mean 67%) of non-inhibited activities. One hour after dosing, the activities had dropped to between 17 and 60% (mean 40%), but 48 h after the last dose they were all at least 85% (mean 95%). Two weeks after the last dose, the subjects had regained their normal pChE activities.

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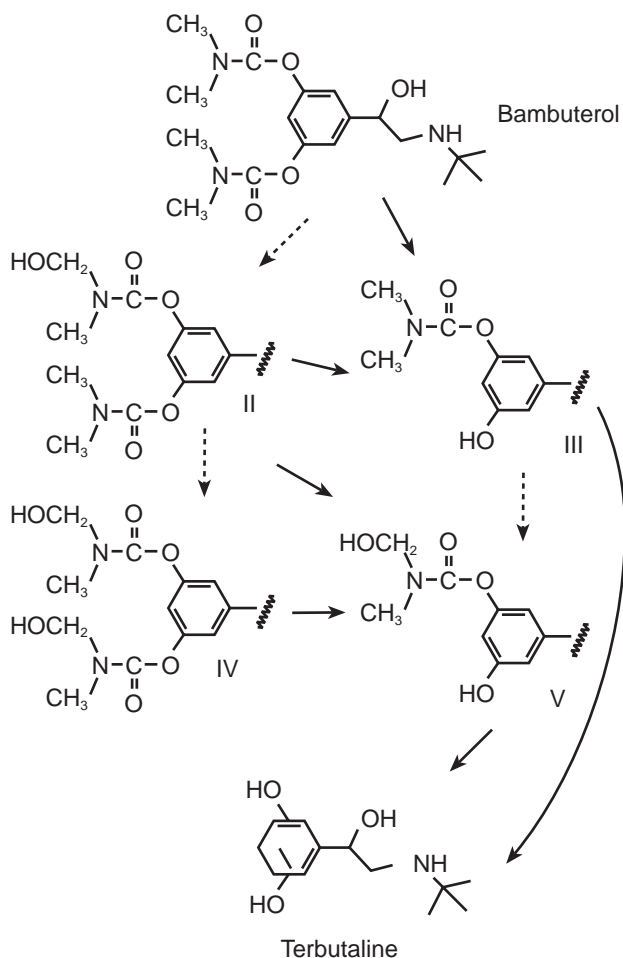


Figure 1 Oxidative (---->) and hydrolytic (→) metabolic pathways of bambuterol to terbutaline.

In Denmark, a register has been built up about subjects with abnormal genotypes of pChE [3]. This offered a possibility to recruit a group of individuals for a study of the fate of bambuterol and the formation of terbutaline in subjects with deficient pChE but with normal oxidative (CYP2D6) capability.

Methods

The study was of an open design. It was run at Herlev Hospital, Herlev, Denmark, in accordance with the Declaration of Helsinki, approved by the Local Ethics Committee, and registered by the Danish National Board of Health and Welfare. The subjects were fully informed verbally and in writing about the nature and objective of the study. Verbal consent to participation was obtained prior to the start of the study.

Subjects

Four healthy subjects, two men and two women (sisters), participated. The subjects were homozygous for the atypical gene for pChE according to the determination of genotype made by the Danish Cholinesterase Research Unit and selected from the card index of this unit.

Demographic data and biochemical findings indicative of pChE genotype of the subjects are given in Table 1. The genotype determination is a result of family studies (pedigree investigations) in combination with biochemical phenotyping using a battery of selective inhibitors [3, 9]. In this study, the family investigation assured that the four subjects were homozygous for the atypical gene (AA), otherwise this genotype could not have been distinguished from that with one silent and one atypical gene (AS).

Hydrolytic capability

The subjects' basal pChE activities measured by the hydrolysis of benzoylcholine [10] were 311–461 (mean 389) U l^{-1} in blood samples taken 17 days before start of treatment (Table 1). Normal range is 690–1560 U l^{-1} [3].

Oxidative capability

As a part of the exclusion criteria, and before entering the study, the subjects were tested for their ability to metabolize debrisoquine [11]. The metabolic ratio (MR) between debrisoquine and its metabolite 4-hydroxy-debrisoquine depends on the activity of the enzyme CYP2D6. All subjects were found to be fast metabolizers of debrisoquine (MR < 12.6, Table 1) and, on average, 79% of the measured 6 h urinary excretion was the hydroxylated metabolite.

Treatment plan

The subjects were given repeated doses of bambuterol hydrochloride as rapidly dissolving tablets (Bambec[®], Astra, Sweden) over a period of 10 days (days 1–10). The dose was 20 mg (50 μmol) taken in the morning after breakfast, then no food was allowed for at least 2 h.

Blood samples were obtained before treatment and up to 36 h after the last dose (days 10–11). The blood samples (10 ml) were collected into sodium-heparinized Venoject[®] sampling tubes into which 100 μl of a 1 mM solution of an esterase inhibitor (D2456, a bambuterol analogue [12]) had been added. The tubes were immediately centrifuged, and the plasma was stored at -20°C .

Urine was collected in tared polyethylene bottles. Two initial collection periods were 6 h, then 12 h collections ensued until 108 h after the last dosing. Two 10 ml portions were stored at -20°C until analysed.

Laboratory assessments

Bambuterol was measured by gas chromatography plus mass spectrometry [12]. At concentrations of 1 nmol l^{-1} in plasma and 8 nmol l^{-1} in urine (spiked samples), the coefficient of variation (CV) was about 4% for plasma and 5.4% for urine at the time of validation of the methods (within-day variation). The lower limit of quantification was set to 0.5 nmol l^{-1} in plasma and to 4 nmol l^{-1} in urine. Terbutaline was measured by high performance liquid chromatography with electrochemical detection. The published procedure [13] was slightly modified for analysis of the urine samples. At the time of validation of the methods, the CV (spiked samples) at 8 nmol l^{-1} in plasma was 2.1%

Table 1 Demographic data and biochemical characteristics of the subjects before treatment with bambuterol.

Subject	Sex	Age (years)	Weight (kg)	Height (cm)	MR*	pChE† (U l ⁻¹)	% inhibition by:				
							Dibucaine	Fluoride	Urea	Scoline	Ro 2-0683
1	F	40	62	169	0.05	440	22	21	90	10	11
2	F	45	55	159	0.36	461	23	19	96	10	15
3	M	50	89	175	0.56	344	23	18	93	8	17
4	M	42	70	180	0.17	311	23	20	100	11	21
Mean		44	69	171	0.20‡	389					
Reference values§: (percentiles 2½–97½)				genotype UU		690–1560	79–87	55–66	41–53	88–94	80–100
				genotype AA		190–732	14–27	16–30	86–100	4–22	9–27

*Metabolic ratio of debrisoquine.

†Plasma cholinesterase activity ($\mu\text{mol benzoylcholine converted min}^{-1}/\text{l}^{-1}$ [3]).

‡Geometrical mean.

§Given in [3]; values for Ro 2-0683 in [9].

within a day and 3.6% between days. In urine at concentrations of 120 and 320 nmol l⁻¹, CV was 7.2 and 4.6%, respectively (between-day variation). The lower limit of quantification was set to 4 nmol l⁻¹ in plasma and to 80 nmol l⁻¹ in urine.

Plasma cholinesterase activity during and after treatment was measured at steady state on days 10–11 (0, 1, 3, 12, and 24 h after dose) and 3 weeks after the last dose of bambuterol.

Plasma and 6 h collections of urine for the determination of creatinine clearance were sampled before treatment and on day 10. The samples were taken from collections also intended for the determination of debrisoquine (before treatment) or bambuterol and terbutaline (day 10). One plasma sample was taken in the middle of the urine collection interval.

Pharmacokinetic parameters

AUC = area under the curve of plasma concentration *vs* time during the dosing interval, calculated by the linear trapezoidal rule.

C_{ss} = average plasma concentration, calculated as AUC divided by the length of the dosing interval (24 h).

C_{max} = peak plasma concentration

T_{max} = time of peak plasma concentration

C_{min} = trough plasma concentration

Ae = amount excreted in urine during the dosing interval.

Renal clearance was calculated as Ae divided by AUC.

dAe/dt = urinary excretion rate, calculated as the amount excreted in a collection period divided by the length of the collection period. The calculated rate was time-located at the midpoint of the period.

ω = terminal disappearance rate constant, calculated from the slope found after linear regression of $\ln[dAe/dt]$ *vs* time data during the last, monoexponential phase of the curve.

Terminal half-life = $\ln 2/\omega$

Creatinine clearance was calculated by dividing dAe/dt of creatinine by plasma creatinine.

Statistical test

A two-tailed *t*-test for difference in means was applied to compare the terminal half-lives of bambuterol and terbutaline

found in this study and the ones found in a previous study in healthy subjects with normal pChE activity [8]. The test was performed on the assumption of an additive model. The level of significance was defined as $P=0.05$.

Results

Pharmacokinetic parameters are given in Table 2.

Plasma data

No subject had detectable plasma concentrations of bambuterol or terbutaline before entering the study. Measured plasma concentrations in the four subjects during a dosing interval after 10 days treatment are shown in Figure 2.

Bambuterol was measurable in plasma during the whole steady-state dosing interval, but at 0 h in subject 1 and at 24 h in subject 3 values were slightly below the limit of quantification of 0.5 nmol l⁻¹. Rough estimates of values were 0.4 and 0.5 nmol l⁻¹, respectively, but, because of the uncertainty of the estimates, they were not used for calculations of C_{max}/C_{min} ratios. Such ratios were calculated only for subjects 2 and 4 (Table 2).

Mean AUC for bambuterol over the 24 h dosing interval was 64.4 (44.5–88.7) nmol l⁻¹h, corresponding to an average plasma concentration of 2.7 nmol l⁻¹. The lowest and the highest values were in subjects 2 and 4, respectively. Peak plasma concentration of bambuterol was reached 1–3 h (median 2.0 h) after the last dose. The mean peak concentration was 10.5 (5.1–17.9) nmol l⁻¹.

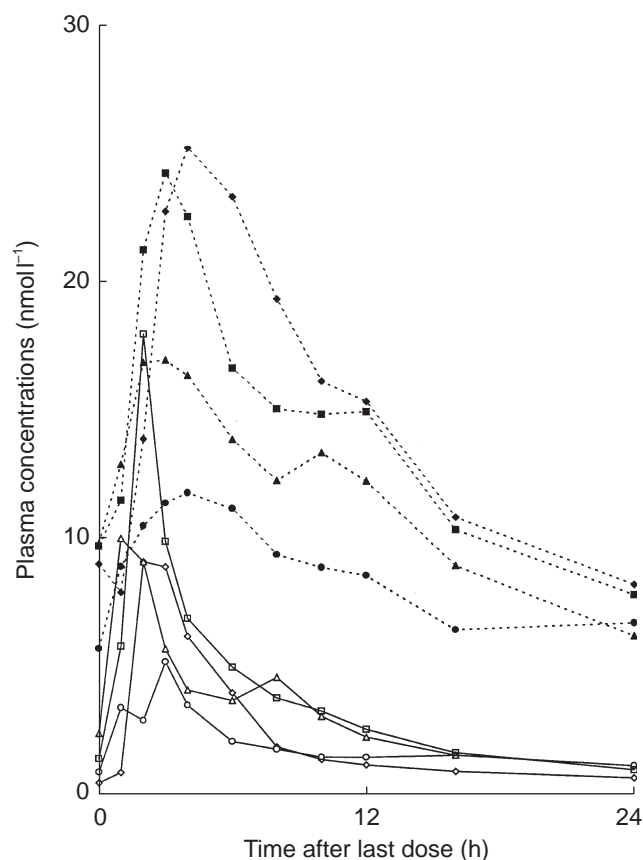
Terbutaline was measurable in plasma up to at least 24 h. Mean AUC for terbutaline over the dosing interval was 286 (200–348) nmol l⁻¹. C_{ss} (mean 11.9 nmol l⁻¹) was time-located 12–13 h (mean 12.8 h) after dose intake. The individual C_{ss} -values had the same rank order as the peak concentrations. Peak plasma concentration of terbutaline occurred 3–4 h (median 3.5 h) after the last dose. Mean peak concentration was 19.5 (11.7–25.2) nmol l⁻¹. As with bambuterol, the lowest peak was found in subject 2. The trough concentrations of terbutaline (mean 6.8 nmol l⁻¹) differed less than the peaks between subjects. The geometrical mean peak/trough concentration ratio was 2.8.

Table 2 Characteristics of the plasma concentration–time curves, renal clearances, and terminal half-lives of bambuterol and terbutaline.

Subject sex	C_{max} (nmol l ⁻¹)	T_{max} (h)	C_{min} (nmol l ⁻¹)	C_{max}/C_{min}	AUC (h × nmol l ⁻¹)	C_{ss} (nmol l ⁻¹)	Renal clearance (ml min ⁻¹)	Terminal half-life (h)	Creatinine clearance (ml min ⁻¹)*
<i>Bambuterol</i>									
1F	9.0	2	(0.4)	—	52.9	2.2	146	9.4	91
2F	5.1	3	0.8	6.6	44.5	1.9	127	12.6	88
3M	9.9	1	(0.5)	—	71.6	3.0	165	4.8	141
4M	17.9	2	0.9	19.0	88.7	3.7	143	8.0	120
Mean	10.5	2.0 (median)			64.4	2.7	145	8.7	110
<i>Terbutaline</i>									
1F	25.2	4	7.8	3.2	348	14.5	95	16.5	
2F	11.7	4	5.6	2.1	200	8.3	95	21.8	
3M	16.9	3	6.2	2.7	269	11.2	132	16.5	
4M	24.2	3	7.8	3.1	326	13.6	122	19.8	
Mean	19.5	3.5 (median)	6.8	2.8†	286	11.9	111	18.6	

*During treatment.

†Geometrical mean.

**Figure 2** Steady-state plasma concentrations of bambuterol and terbutaline in four subjects with atypical plasma cholinesterase who ingested 20 mg of bambuterol hydrochloride once daily for 10 days.

Urinary data

No subject had detectable urine concentrations of bambuterol or terbutaline before entry into the study. The mean urinary excretion of bambuterol over the dosing interval was 568 (338–763) nmol, corresponding to 1.14% of the administered dose. Within the interval, the major part was excreted during

the initial 12 h (mean 80%). Urinary excretion data during the dosing interval reflected the plasma concentration of bambuterol with the lowest excretion in subject 2 (0.68% of the dose) and the highest excretion in subject 4 (1.53%). Mean renal clearance of bambuterol over the entire dosing interval was 145 ml min⁻¹, which was higher than creatinine clearance in all participants. Creatinine clearance was of the same order before and during bambuterol treatment.

The mean amount of terbutaline excreted during the dosing interval was 1903 (1134–2376) nmol, corresponding to 3.81% of the dose. The individual values ranked like those found with bambuterol. About 60% of the excretion within the interval took place during the first 12 h after dosing. Mean renal clearance of terbutaline over the entire dosing interval was 111 ml min⁻¹, similar to creatinine clearance in all participants.

Terminal half-lives

The mean terminal half-life of bambuterol was 8.7 (4.8–12.6) h, shorter than the 14.2 h previously found [8] in subjects with normal pChE activity ($P=0.044$). The mean terminal half-life of terbutaline was 18.6 (16.5–21.8) h, similar to the value found in the normal subjects (mean 19.7 h).

Plasma cholinesterase activity

The inhibition by the treatment was low. The greatest inhibition was found in the samples taken 3 h after the last administration of bambuterol: at the most 26% (subject 4).

Adverse events

Adverse events were recorded on study days 1–14. Participants except subject 2 experienced mild or moderate adverse events, consistent with those commonly seen with β_2 -adrenoceptor agonists: tremor, restlessness, headache and palpitations.

Discussion

Surprisingly, the four subjects with atypical cholinesterase were all capable of producing terbutaline from bambuterol as efficiently as subjects with normal hydrolytic and oxidative capacity [8]. We had expected to find a higher AUC of bambuterol and a diminished and slower production of terbutaline as a result of these subjects' low basal activity of plasma cholinesterase and the poor affinity of the atypical enzyme for bambuterol. Instead, the pharmacokinetic parameters found in the present study (Table 2) were very similar to those found in a study of bambuterol in subjects with normal pChE activity given the same dose [8]. The subjects with atypical cholinesterase had a shorter terminal half-life and slightly higher plasma concentrations of bambuterol. They had a somewhat higher peak concentration of terbutaline with an increased fluctuation (C_{max}/C_{min}) within the dosing interval. However, the AUC values of terbutaline were the same as those found in subjects with normal pChE activity.

This contradicts partly the hypothesis (based on *in vitro* studies) [1] that both the oxidative and the hydrolytic pathways are required for optimum bioconversion of bambuterol to terbutaline.

Formation of terbutaline from bambuterol cannot occur without hydrolysis (Figure 1). Plasma cholinesterase is considered to be the blood enzyme predominantly catalysing the hydrolysis. However, the quantitative importance of hydrolysis of bambuterol in plasma relative to hydrolysis in tissues like intestine and liver has hitherto not been determined. Liver microsomes from various species have been tested for their capacity to hydrolyse bambuterol [1]. Microsomes from human livers had intermediate activity and those from rat liver no activity, while liver microsomes from guinea pigs had the highest activity. The enzymes catalysing these processes may be non-specific carboxyesterases known to be very active in liver microsomes or nonspecific cholinesterases probably present in the microsomal preparations [1]. In the absence of plasma cholinesterase, these enzymes may contribute to a greater extent to the hydrolytic conversion of bambuterol to its active metabolite. Also, the hydroxylated metabolites (II, IV, and V in Figure 1) have been shown to be converted partially to *N*-demethylated compounds which hydrolyse spontaneously with half-lives of a few hours at body temperature and physiological pH [14].

Debrisoquine was chosen as a marker of oxidative metabolism in the present study, because bambuterol has a molecular structure that fulfils requirements for being a substrate for CYP2D6 [15, 16]. Indeed, preliminary experiments showed disappearance of bambuterol and/or rise of hydroxylated metabolites when bambuterol was incubated with cDNA-expressed CYP2D6 (Gentest Corp, Woburn, MA, USA; G Hallström, G Jönsson, personal communication). The prestudy debrisoquine test revealed that our four subjects were markedly extensive metabolizers [17]. As about 90% of the Danish population are characterized as fast metabolizers of debrisoquine/sparteine, this finding might be incidental [18]. However, it cannot be excluded that a high oxidative metabolism in the present subjects might have compensated for their low plasma cholinesterase activity

by promoting formation of intermediary metabolites less stable towards hydrolysis. The oxidative pathway of bambuterol metabolism may then be quantitatively more important *in vivo* than previously believed. Further analysis of intermediate metabolites would be helpful to determine the relation between oxidative and hydrolytic metabolism and to identify the enzymes involved.

Bambuterol terminal half-life after oral administration is a measure of uptake rather than elimination [8]. Therefore, the shorter terminal half-life of bambuterol and the higher peak concentrations of bambuterol and terbutaline found in the four atypical subjects as compared with genetically normal subjects indicate a faster absorption or bioavailability rate of the prodrug. It is not known how much bambuterol is biotransformed by pChE or by enzymes other than cholinesterases during passage through the intestinal mucosa. The normal pChE enzyme may act as a limiting factor regarding the fraction of the absorbed dose that reaches the blood and subsequently the liver intact. In the four subjects studied, the lower activity of pChE and its low affinity for bambuterol might have allowed a larger intact fraction to pass the intestinal mucosa into the portal blood, thereby also promoting a fast oxidative metabolism in the liver.

In conclusion, after repeated oral administration of 20 mg bambuterol hydrochloride, four subjects, homozygous for the atypical gene for plasma cholinesterase, had average plasma concentrations of bambuterol and its active metabolite terbutaline which were grossly similar to those seen in subjects with normal plasma cholinesterase activity. This might be explained by an altered uptake and metabolism in the absence of plasma cholinesterase, or the importance of this enzyme for the formation of terbutaline from bambuterol *in vivo* may have been overestimated.

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