



Synthesis and *in vitro* evaluation of potential sustained release prodrugs via targeting ASBT

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ABSTRACT

The objective was to synthesize prodrugs of niacin and ketoprofen that target the human apical sodium-dependent bile acid transporter (ASBT) and potentially allow for prolonged drug release. Each drug was conjugated to the naturally occurring bile acid chenodeoxycholic acid (CDCA) using lysine as a linker. Their inhibitory binding and transport properties were evaluated in stably transfected ASBT-MDCK monolayers, and the kinetic parameters K_t , K_t , $normJ_{max}$, and P_p were characterized. Enzymatic stability of the conjugates was evaluated in Caco-2 and liver homogenate. Both conjugates were potent inhibitors of ASBT. For the niacin prodrug, substrate kinetic parameter K_t was 8.22 μM and $normJ_{max}$ was 0.0917. In 4 h, 69.4% and 26.9% of niacin was released from 1 μM and 5 μM of the conjugate in Caco-2 homogenate, respectively. For the ketoprofen prodrug, K_t was 50.8 μM and $normJ_{max}$ was 1.58. In 4 h, 5.94% and 3.73% of ketoprofen was released from 1 μM and 5 μM of the conjugate in Caco-2 homogenate, and 24.5% and 12.2% of ketoprofen was released in liver homogenate, respectively. *In vitro* results showed that these bile acid conjugates are potential prolonged release prodrugs with binding affinity for ASBT.

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1. Introduction

The human apical sodium-dependent bile acid transporter (ASBT, SLC10A2) is expressed at high levels in the terminal ileum where it mediates bile acid recovery (Dawson et al., 2003). ASBT is critical for intestinal reabsorption of bile acid in the enterohepatic recirculation. ASBT's high affinity and capacity for native bile acids affords it to be a potential prodrug target for intestinal drug absorption (Balakrishnan and Polli, 2006; Tolle-Sander et al., 2004), as well as a contributing factor to prolong drug release via enterohepatic recirculation. Previously, bile acid conjugates or bile acid derived compounds showed potential to provide sustained systemic concentrations of the drug after oral administration (Gallop and Cundy, 2002a,b). Prolonged drug release from the drug conjugated to a bile acid may be achieved via enterohepatic recirculation, where parent drug is slowly released into the systemic circulation from the bile acid conjugate.

Niacin (nicotinic acid) has been used in the management of hyperlipidemia for over 50 years. Clinical use of niacin has been limited by a relatively high incidence of cutaneous flushing. Niacin undergoes extensive first-pass metabolism in the liver via the conjugation and the amidation pathways (Menon et al., 2007; Piepho,

2000; Stern et al., 1992). The conjugation metabolic pathway has low affinity and high capacity and results in the formation of glycine conjugates of niacin which has been associated with flushing. The cutaneous flushing is mediated by a G-protein-coupled receptor (GPR109A) effect involves release of prostaglandin (PGE2, PGD2) (Benyo et al., 2005; Pike, 2005). The amidation pathway has high affinity and low capacity. When the amidation pathway is saturated, niacin is mainly metabolized by the conjugative pathway, resulting in flushing. The pharmacokinetic and pharmacodynamic profiles of different niacin formulations reflect differences in their absorption rates and metabolic depositions. Slower niacin absorption causes lower flushing as most of the drug undergoes amidation (Pieper, 2002, 2003). Extended-release niacin has demonstrated reduced rates of flushing with a t_{max} of 4–5 h, compared to the immediate-release niacin of 30–60 min (Knopp et al., 1998; Morgan et al., 1998).

Ketoprofen is a potent non-steroidal anti-inflammatory drug (NSAID). It is associated with higher gastrointestinal (GI) bleeding/perforation compared to other NSAIDs (Gonzalez et al., 2010). It has a short half-life of 2 h, requiring frequent doses to maintain therapeutic efficacy (Kantor, 1986).

The objective of this study is to design respective prodrugs of niacin and ketoprofen by conjugating them with bile acid and targeting ASBT for absorption. Prolonged availability of the parent drug may be expected if prodrug is slowly hydrolyzed during enterohepatic circulation. This approach could reduce flushing from niacin and extend the pharmacological effect of ketoprofen. Moreover, the approach could minimize ketoprofen GI side effects

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by preventing the parent drug direct contact with the GI tract. In this study, niacin and ketoprofen were conjugated with the naturally occurring bile acid chenodeoxycholate (CDCA) via lysine as a linker. CDCA was chosen due to its high ASBT binding affinity (Balakrishnan et al., 2006b). *In vitro* results indicated that both conjugates were potential prodrugs with affinity for ASBT.

2. Materials and methods

2.1. Materials

[³H]-Taurocholic acid was purchased from PerkinElmer (Waltham, MA). Taurocholic acid, nicotinoyl chloride hydrochloride, and ketoprofen were purchased from Sigma (St. Louis, MO). Protected lysine analogs were from Novabiochem (Gibbstown, NJ). Fetal bovine serum (FBS), trypsin, and Dulbecco's modified Eagle's medium (DMEM) were purchased from Invitrogen Corporation (Carlsbad, CA). CDCA was obtained from TCI America (Portland, OR). Pooled IGS Sprague–Dawley male rat liver S9 fraction was purchased from Xenotech (Lenexa, KS).

2.2. Synthetic procedures

Among the native bile acids, CDCA was selected on the basis of its favorable transport across ASBT-MDCK monolayers (Balakrishnan et al., 2006b). Pyridine-3-carboxylic acid and ketoprofen were conjugated to CDCA using lysine as the linker. Synthetic approaches of both prodrugs are shown in Scheme 1.

2.2.1. Synthesis of CDCA–lysine (4)

First, 5 g of CDCA **2** were coupled to 0.5 equiv. of H-lys(Z)-OBzl HCl **1** using HBTU (1.1 equiv.) and HOBT (0.5 equiv.) as the coupling reagent in 10 mL of anhydrous DMF, along with 2.2 equiv. of triethylamine (TEA), at 60 °C for overnight. The reaction was terminated by adding 30 mL of water. The reaction mixture was extracted with ethyl acetate (EtOAc) (3×). The combined organic layer was washed with 1N HCl (3×), 1N NaOH (3×), and brine (1×). The organic extract was dried with anhydrous sodium sulfate, filtered, and evaporated under vacuum to yield orange oil. The crude product was further purified by silica gel column chromatography using a mobile phase of EtOAc and hexane (80:20). Fractions were collected. Mass spectrometry (MS) showed an appropriate peak [M+H]⁺ 745.3. Solvent

of fractions in TLC and MS that contained the product were collected and evaporated to give a white fluffy solid compound **3**.

The neutral compound **3** was subjected to catalytic hydrogenation for the removal of the benzyl and CBZ protection groups using 10% Pd/charcoal in ethanol at 50 psi overnight. Suspension was filtered through Celite® and solvent was evaporated under vacuum to give white solid intermediate **4**. MS showed an appropriate peak [M+H]⁺ 521.4.

2.2.2. Synthesis of CDCA–lysine–niacin (6)

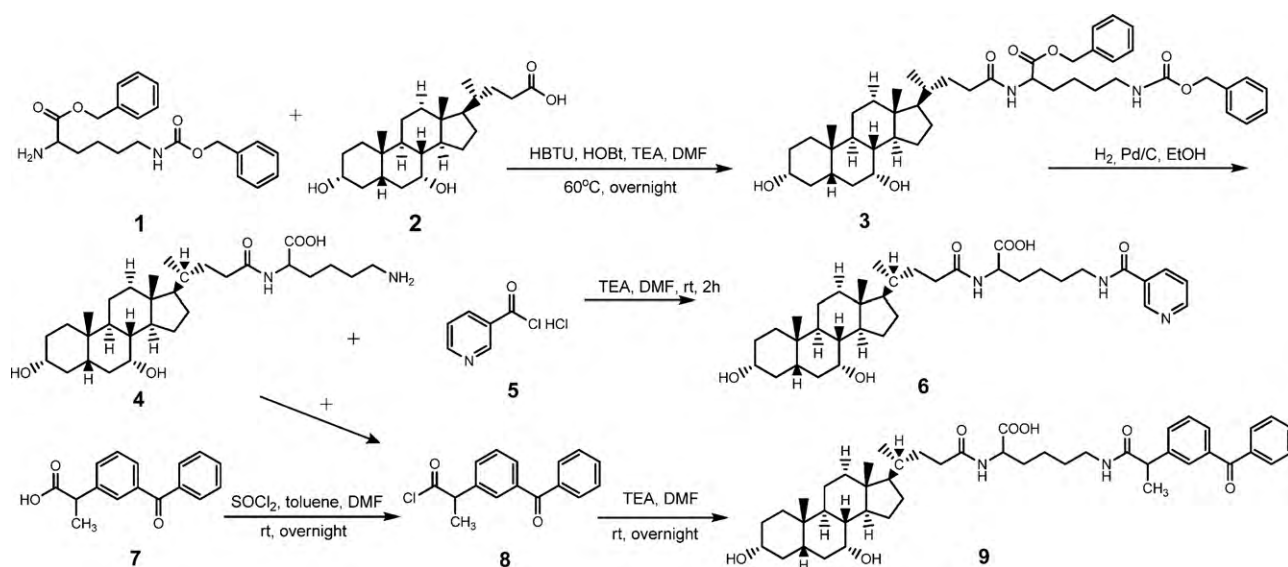
Compound **4** (0.5 g) was coupled to 2.0 equiv. of nicotinoyl chloride hydrochloride **5** in 5 mL of anhydrous DMF along with 3.0 equiv. of TEA. The reaction was conducted at 4 °C for 4 h. The reaction was terminated by adding 30 mL of water. The reaction mixture was extracted with EtOAc (3×). The combined organic layer was washed with 1N HCl (3×) and water (3×). Then organic layer was extracted with 1N NaOH (3×). The combined water layer was washed with EtOAc (3×). HCl solution (37%) was added drop by drop to the aqueous layer until the pH was about 2. Then the solution was extracted with EtOAc (3×) again and the organic layer was washed with brine (1×). The organic extract was dried with anhydrous sodium sulfate, filtered, and evaporated under vacuum to yield the final compound **6**. MS showed an appropriate peak [M+H]⁺ 626.3.

2.2.3. Synthesis of CDCA–lysine–ketoprofen (9)

Ketoprofen **7** (5 g) was activated using 20% thionyl chloride in 10 mL of toluene with a few drop of DMF as the catalyst at RT for 30 min. The reaction was terminated by evaporating the solvent under vacuum to yield a white solid **8**. The resulting compound **8** was coupled to 0.5 equiv. of **4** in anhydrous DMF along with 3.0 equiv. of TEA at RT overnight. The reaction was terminated by adding 30 mL of water. The reaction mixture was extracted and purified as described above (see synthesis of CDCA–lysine–niacin). The organic extract was dried with anhydrous sodium sulfate, filtered, and evaporated under vacuum to yield the final compound **9**. MS showed an appropriate peak [M+H]⁺ 757.5.

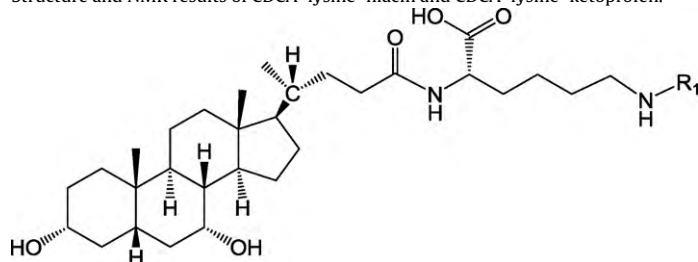
2.3. Characterization of the prodrugs

Identity and purity were characterized by TLC, LC–MS, ¹H NMR (proton, COSY), and LC–MS/MS. The mass spectra gave a molecular



Scheme 1. General synthetic approach to obtain CDCA–lysine–niacin (**6**) and CDCA–lysine–ketoprofen (**9**).

Table 1
Structure and NMR results of CDCA–lysine–niacin and CDCA–lysine–ketoprofen.



Conjugate	R ₁	¹ H NMR (DMSO- <i>d</i> ₆) ^a
CDCA–lysine–niacin		11.95 (1H, s, γ), 8.99 (1H, s, α), 8.69 (1H, d, α), 8.64 (1H, t, β), 8.17 (1H, d, α), 7.92 (1H, d, β), 7.48 (1H, t, α), 4.30 (1H, m, θ),
CDCA–lysine–ketoprofen		11.65 (1H, s, γ), 7.98 (1H, d, β) 8.04 (4, t, β) 7.78–7.58 (9H, α), 4.13(1H, m, γ) 3.68 (1H, m, α), 1.37 (3H, s, α)

^a Only characteristic peaks are shown. CDCA and methylene peaks are not shown. Values are in ppm relative to TMS. α indicates assignment to proton peaks in R₁; β indicates assignment to protons in amide; γ indicates assignment to protons in carboxylic acid region; θ indicates the proton in α carbon.

ion in agreement with the calculated mass of the prodrug. No CDCA impurity was detected in any final conjugate (Gonzalez and Polli, 2008). ¹H NMR results are compiled in Table 1. The chemical shifts are reported in parts per million (ppm) relative to tetramethylsilane (TMS) ($\delta = 0.0$ ppm). ¹H NMR spectra were recorded on a Varian Inova 500 MHz (Varian Inc., Palo Alto, CA).

2.4. Cell culture

Stably transfected ASBT-MDCK were cultured as described (Balakrishnan et al., 2005). Briefly, cells were grown at 37 °C, 90% relative humidity, and 5% CO₂ atmosphere and fed every 2 days by the DMEM supplemented with 10% FBS, 50 units/mL penicillin, and 50 μg/mL streptomycin. Geneticin was used at 1 mg/mL to maintain selection pressure. Cells were passaged every 4 days or after reaching 90% confluence.

Caco-2 cells were maintained in DMEM with 10% fetal bovine serum, 1% nonessential amino acid solution, 1% antibiotic solution (penicillin/streptomycin), and 1% glutamine solution and grown in flasks. Cells were incubated at 37 °C with 5% CO₂. The medium was changed every 2 days.

2.5. Enzymatic stability study

Prodrug was subjected to Caco-2 homogenate and liver S9 fraction. Intact prodrug and the liberated drug were quantified.

2.5.1. Caco-2 homogenate

Caco-2 cells grown in 225-cm² culture flask for 21–25 days were washed twice with ice-cold phosphate-buffered saline (PBS). The cells were scraped off in 12 mL ice-cold PBS and the scraped cells were transferred to a centrifuge tube. Tubes were stored at –80 °C overnight. The next day, tubes were quickly thaw at 37 °C (leave some ice), sonicated the cells twice at 1 s pulses with a sonicator probe (F60 Sonicator, Thermo Fisher Scientific Inc., Waltham, MA). The cell homogenate (protein content = 0.63 mg/mL) was stored at –80 °C (Tantishaiyakul et al., 2002).

2.5.2. Stability study

To begin the stability assay, each prodrug was added into a centrifuge tube containing 1000 μL Caco-2 cell homogenate or liver S9 fraction (20 mg/mL) to yield a prodrug concentration of 1 μM or 5 μM. Tubes were placed in a 37 °C incubator and shaken. Samples (100 μL) were removed at selected time points over 4 h and immediately added to a new centrifuge tube pre-filled with 100 μL ice-cold acetonitrile to precipitate cellular protein and stop enzymatic activity. Samples were centrifuged at 12,000 rpm for 30 min at 4 °C. The supernatant was carefully transferred to another centrifuge tube. The samples were stored at –80 °C until further analysis.

2.6. ASBT biological activity assessment

Each prodrug was subjected to ASBT inhibition and uptake studies.

2.6.1. Inhibition of taurocholate uptake by prodrugs

To characterize prodrug binding affinity to ASBT, cis-inhibition studies of taurocholate uptake by prodrugs were conducted as previously described (Zheng et al., 2009). Briefly, cells were exposed to donor solution containing 2.5 μM taurocholate (including [³H]-taurocholate) in the presence of a prodrug at eight concentrations (1–100 μM) for 10 min at 37 °C. At the end of the study, cell lysate was counted for taurocholate radioactivity using a liquid scintillation counter (Beckmann Instruments, Inc., Fullerton, CA).

2.6.2. Uptake of prodrug into ASBT-MDCK

To characterize prodrug substrate properties across ASBT, uptake studies of prodrugs were conducted as described (Zheng et al., 2009). Briefly, cells were exposed to donor solution containing eight different prodrug concentrations (1–500 μM) for 10 min at 37 °C. The studies were conducted in both HBSS buffer or modified HBSS buffer (i.e. sodium-free). Since ASBT is a sodium-dependent transporter, studies using sodium-free buffer enabled passive permeability measurement of the prodrug. Subsequently, to quantify uptake, cells were lysed as pervious described by adding 0.25 mL acetonitrile to each well (Rais et al., 2008). Following evaporation

Table 2
LC–MS/MS methods of analyzing CDCA–lysine–niacin and CDCA–lysine–ketoprofen and the parent drugs niacin and ketoprofen.

Compound	Exact mass ^a	Q1 mass (m/z) ^b	Q3 mass (m/z)	Retention time (min)	Gradient timetable (B%) ^c
CDCA–lysine–niacin	625.41	626.40	189.12	1.26	0–0.2 (45%), 0.2–3.0 (45–95%), 3.0–4.0 (95%), 4.0–4.2 (95–45%), and 4.2–7.0 min (45%)
Niacin	123.03	124.06	80.15	1.02	0–3.1 (5%), 3.1–3.2 (5–90%), 3.2–4.2 (90%), 4.2–4.5 (90%–5%), and 4.5–8.0 min (5%)
CDCA–lysine–ketoprofen	756.47	757.50	485.30	4.77	0–0.2 (40%), 0.2–3.0 (40–95%), 3.0–4.0 (95%), 4.0–4.2 (95–40%), and 4.2–7.0 min (40%)
Ketoprofen	254.09	255.10	198.36	2.36	0–0.2 (40%), 0.2–3.0 (40–95%), 3.0–4.0 (95%), 4.0–4.2 (95–40%), and 4.2–7.0 min (40%)

^a Calculated monoisotopic mass.

^b Q1 and Q3 mass are experimental mass [M+H]⁺.

^c The mobile phase was A, 0.1% formic acid in water; and B, 0.1% formic acid in acetonitrile. The gradient times were listed as the percent organic phase, B%.

at RT, 1 mL solution containing 50% water and 50% acetonitrile was added to each well for 10 min. Sample was stored in –80 °C until analyzed by LC–MS/MS.

2.7. Analytical methods

Samples from uptake studies were quantified by LC–MS/MS on a Thermo Finnigan Surveyor HPLC system, equipped with a Thermo Finnigan Surveyor autosampler and a Thermo TSQ quantum mass spectrometer (Thermo Fisher Scientific Inc., Waltham, MA). For both prodrugs and parent ketoprofen, the column was a Phenomenex Luna C8 (50 mm × 2.0 mm, 3 μm; Phenomenex; Torrance, CA). For parent niacin, the column was a Thermo Aquasil C18 (50 mm × 2.1 mm, Thermo Fisher Scientific, MA). Columns were heated to 45 °C. The mobile phase was A, 0.1% formic acid in water; and B, 0.1% formic acid in acetonitrile, with a flow rate of 0.4 mL/min. The injection volume was 10 μL. Detection was achieved under positive ion electrospray [M+H]⁺ tandem mass spectrometry, as positive mode provided the greatest sensitivity. The mobile phase and retention time for both conjugates and parent drugs are detailed in Table 2.

2.8. Kinetic analysis

Inhibition data was analyzed in terms of inhibition constant K_i as previously described (Zheng et al., 2009), where a modified Michaelis–Menten competitive inhibition model that considers aqueous boundary layer (ABL) resistance was used (Balakrishnan et al., 2007).

$$J = \frac{P_{ABL}(J_{max}/(K_t(1 + (I/K_i))) + S) + P_p}{P_{ABL} + (J_{max}/(K_t(1 + (I/K_i))) + S) + P_p} S \quad (1)$$

where J is taurocholate uptake, J_{max} and K_t are the Michaelis–Menten constants for ASBT-mediated transport, S is taurocholate concentration, P_p is the passive taurocholate permeability, I is the inhibitor concentration, K_i is the inhibition constant, and P_{ABL} is the aqueous boundary layer permeability (i.e. 150×10^{-6} cm/s) (Balakrishnan et al., 2007). K_t was 5.03 μM, obtained from a pooled data analysis approach. J_{max} was measured from taurocholate uptake studies using 200 μM of taurocholate; P_p was estimated from taurocholate uptake studies in the absence of sodium. K_i was calculated using WinNonlin Professional (Pharsight Corporation; Mountain View, CA). J_{max} and P_p of taurocholate were measured on each study occasion.

For uptake studies, uptake data into monolayers was analyzed in terms of substrate affinity K_t and was fitted to a previously

developed modified Michaelis–Menten model (Eqs. (2) and (3)) (Balakrishnan et al., 2007):

$$J = \frac{P_{ABL}((J_{max}/(K_t + S)) + P_p)}{P_{ABL} + ((J_{max}/(K_t + S)) + P_p)} S \quad (2)$$

$$J = \frac{P_{ABL}P_pS}{P_{ABL} + P_p} \quad (3)$$

where J is prodrug uptake. Eqs. (2) and (3) were applied simultaneously to data measured with sodium and without sodium to estimate K_t , J_{max} , and P_p . Data were analyzed using WinNonlin Professional.

3. Results

3.1. Prodrug synthesis

Bile acid conjugates CDCA–lysine–niacin and CDCA–lysine–ketoprofen were successfully synthesized. Their structures are depicted in Table 1. For each, the synthetic strategy was based on coupling the carboxyl group at the C-24 position of CDCA (Scheme 1) to the carboxyl group of ketoprofen or niacin using lysine as a linker. Prodrug properties were characterized using NMR and LC–MS, and the resulting data are shown in Tables 1 and 2, respectively. Both LC–MS and NMR data were consistent with the target structures. TLC showed a single spot.

3.2. ASBT inhibition and uptake

Inhibition of taurocholate uptake was carried out for each prodrug using stably transfected ASBT-MDCK monolayers. Fig. 1 illustrates the concentration-dependent inhibition profile by CDCA–lysine–niacin and niacin (Fig. 1A), and CDCA–lysine–ketoprofen and ketoprofen (Fig. 1B), respectively. Taurocholate uptake was reduced over 2-fold in the presence of 50 μM of each prodrug. Neither ketoprofen nor niacin was ASBT inhibitor. The K_i value indicated that each prodrug was a potent inhibitor and is listed in Table 3, with K_i of 12.9 μM for niacin prodrug and 15.6 μM for ketoprofen prodrug, respectively. Compared to the K_i range of native bile acids from 0.427 μM to 22.6 μM, with taurocholate $K_i = 4.96$ μM, both prodrugs were potent inhibitors and within the range of native bile acids (Balakrishnan et al., 2006b).

Given favorable binding affinity for ASBT, prodrugs were evaluated for ASBT-mediated uptake. Fig. 2 illustrates the concentration-dependent transport of CDCA–lysine–niacin and CDCA–lysine–ketoprofen into ASBT-MDCK monolayers. Table 3

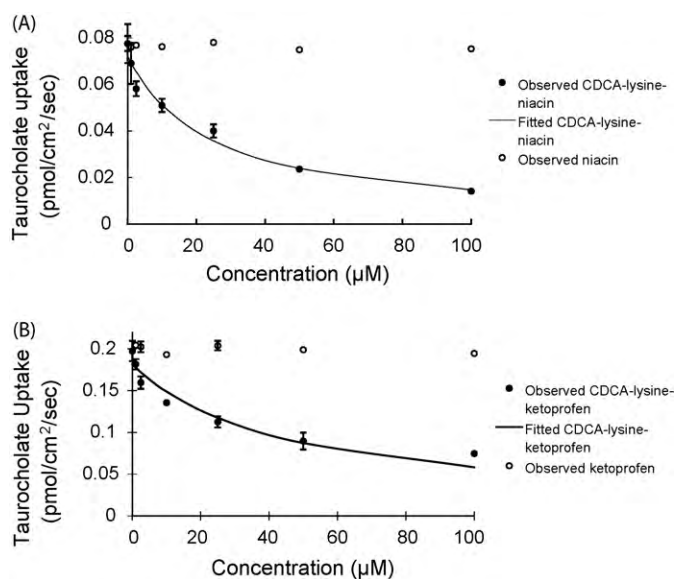


Fig. 1. Concentration-dependent inhibition of taurocholate uptake into ASBT-MDCK monolayers by (A) CDCA-lysine-niacin (●) and niacin (○); (B) CDCA-lysine-ketoprofen (●) and ketoprofen (○) for 10 min. Circles indicate observed data points, while the solid line indicates model fit. Taurocholate uptake into ASBT-MDCK cells was reduced significantly by both conjugates, where K_i was 12.9 μM for niacin prodrug and 15.6 μM for ketoprofen prodrug. Taurocholate uptake was not reduced by niacin or ketoprofen.

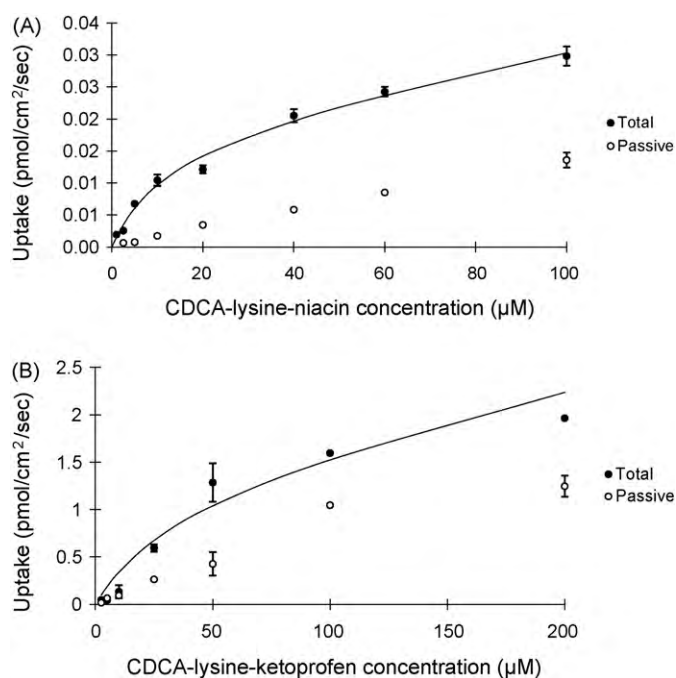


Fig. 2. Concentration-dependent uptake of CDCA-lysine-niacin (A) and CDCA-lysine-ketoprofen (B) into ASBT-MDCK monolayers for 10 min. The conjugate was measured at varying concentrations in the presence and absence of sodium to delineate ASBT-mediated and passive uptake components. Data indicate total uptake in the presence of sodium (●) and passive uptake in the absence of sodium (○), while the solid line indicates model fit of the total uptake.

Table 3
ASBT inhibition and uptake kinetics of CDCA-lysine-niacin and CDCA-lysine-ketoprofen.

Conjugate	K_i (μM)	K_t (μM)	$normJ_{max}^a$	$P_p \times 10^6$ (cm/s)
CDCA-lysine-niacin	12.9 (± 2.6)	8.22 (± 3.63)	0.0917 (± 0.0102)	0.367 (± 0.023)
CDCA-lysine-ketoprofen	15.6 (± 2.7)	50.8 (± 23.2)	1.87 (± 0.36)	6.38 (± 0.44)

Data were summarized as mean (\pm SEM) of three measurements.

^a To accommodate variation in ASBT expression levels across studies, J_{max} of each bile acid was normalized against taurocholate J_{max} from the same study yielding $normJ_{max}$.

Table 4

Niacin and ketoprofen released from CDCA-lysine-niacin and CDCA-lysine-ketoprofen, respectively in Caco-2 and liver S9 after 4 h.

	Caco-2	Liver S9
Percent niacin released in 1 μM CDCA-lysine-niacin	69.4%	– ^a
Percent niacin released in 5 μM CDCA-lysine-niacin	26.9%	–
Percent ketoprofen released from 1 μM CDCA-lysine-ketoprofen	5.94%	24.5%
Percent ketoprofen released from 5 μM CDCA-lysine-ketoprofen	3.73%	12.2%

^a Cannot be measured due to a high baseline niacin level in rat liver S9 fraction.

lists the kinetic uptake parameters for both compounds. Niacin prodrug had a high substrate binding affinity with a $K_t = 8.22 \mu\text{M}$, but the transport capacity is significantly reduced compared to taurocholate, such that $normJ_{max}$ of niacin prodrug was only 0.0917. $normJ_{max}$ was obtained by dividing prodrug J_{max} by taurocholate J_{max} . Therefore, ASBT's capacity to transport niacin prodrug is only about 9% of that of taurocholate.

Ketoprofen exhibited a more modest substrate binding affinity of $K_t = 50.8 \mu\text{M}$, which is 10-fold higher than the taurocholate $K_t = 5.03 \mu\text{M}$. However, the transport capacity of this prodrug was 1.87-fold of the capacity for taurocholate. The passive permeability of CDCA-lysine-ketoprofen was $6.38 \times 10^{-6} \text{ cm/s}$, which was 20-fold higher compared to CDCA-lysine-niacin of $0.367 \times 10^{-6} \text{ cm/s}$. This difference reflects the greater lipophilicity afforded by the ketoprofen moiety ($\text{Log } P = 3.31$), compared to the niacin moiety ($\text{Log } P = 0.25$). When $[S] = 1/2K_t$, passive permeability accounts for 10.3% for total niacin prodrug uptake, and 44.1% for ketoprofen prodrug.

3.3. Enzymatic stability of prodrugs and drug release

Each prodrug was designed with the employment of two amide linkages, which are stable in acid and base. Chemical stability of prodrugs was evident from successful synthesis, which entailed extracting prodrug using strong acidic and basic solutions; no breakdown to potential degradants was observed in MS. Therefore, general chemical stability of the prodrugs was concluded.

Prodrug hydrolysis can occur in many locations during enterohepatic circulation. In order to assess prodrug hydrolysis, enzymatic stability was conducted in Caco-2 homogenate and rat liver homogenate, using prodrugs concentrations of 1 μM and 5 μM . Time-dependent prodrug degradation and corresponding parent drug release were monitored. Table 4 summarizes the percentage of conversion of prodrug to the parent drug after 4 h.

3.3.1. Niacin prodrug

In Fig. 3, the niacin prodrug decreased with time in Caco-2 homogenate, while niacin increased with time. Under the same conditions, 5 μM prodrug displayed a slower hydrolysis profile compared to 1 μM , with 26.9% and 69.4% of niacin released in 4 h, respectively. The mass balance during niacin prodrug hydrolysis was over 100% for the last two sample times (i.e. 180 min and

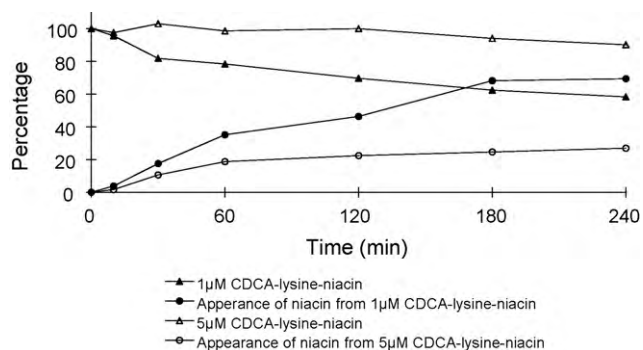


Fig. 3. Hydrolysis of CDCA-lysine-niacin in Caco-2 homogenate. Data indicate the time course of disappearance of 1 μM (\blacktriangle) and 5 μM (\triangle) of CDCA-lysine-niacin and appearance of niacin from 1 μM (\bullet) and 5 μM CDCA-lysine-niacin (\circ).

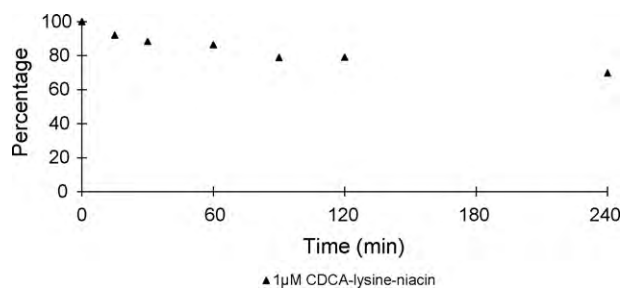


Fig. 4. Hydrolysis of CDCA-lysine-niacin in rat liver S9 fraction. Data indicate the time course of the disappearance of 1 μM of CDCA-lysine-niacin (\blacktriangle).

240 min). The error was possibly caused by evaporation via incubation at 37 °C over this time frame, which involved several sample withdraws.

LC-MS/MS showed a high baseline level of niacin (i.e. nicotinic acid) in rat liver S9 fraction. Therefore, the generation profile of niacin in liver homogenate was not able to be measured. Fig. 4 shows hydrolysis profile of 1 μM niacin prodrug in liver homogenate. About 30.3% of the prodrug was degraded after 4 h, which was slightly less than the 41.7% degradation in Caco-2 homogenate.

3.3.2. Ketoprofen prodrug

Figs. 5A and 6A show the stability profile of the ketoprofen prodrug in Caco-2 and liver homogenate, respectively. Like the niacin prodrug, under identical conditions, the hydrolysis rate of 5 μM prodrug was slower than that of 1 μM prodrug. Table 4 shows only 5.94% and 3.73% of ketoprofen was released from 1 μM to 5 μM of prodrug in Caco-2 homogenate after 4 h; while 24.5% and 12.2% was released in liver homogenate. The time-dependent appearance of ketoprofen is re-plotted using a smaller scale in Figs. 5B and 6B. The prodrug demonstrated high metabolic stability in Caco-2 homogenate. Similarly, conversion of prodrug to ketoprofen rate was slower in liver homogenate.

4. Discussion

4.1. Enterohepatic circulation of bile acid conjugates

The prodrugs were designed to be absorbed at the terminal ileum by the high capacity and high affinity transporter ASBT, followed by a slow release of the parent drug via enzymatic hydrolysis during enterohepatic circulation. Bile acid enterohepatic circulation is a very efficient system where 3–5 g bile acid pool cycles six times daily and results in a turnover of 12–18 g of bile acid each day in humans. Only less than 0.5 g bile acid is lost in feces each day,

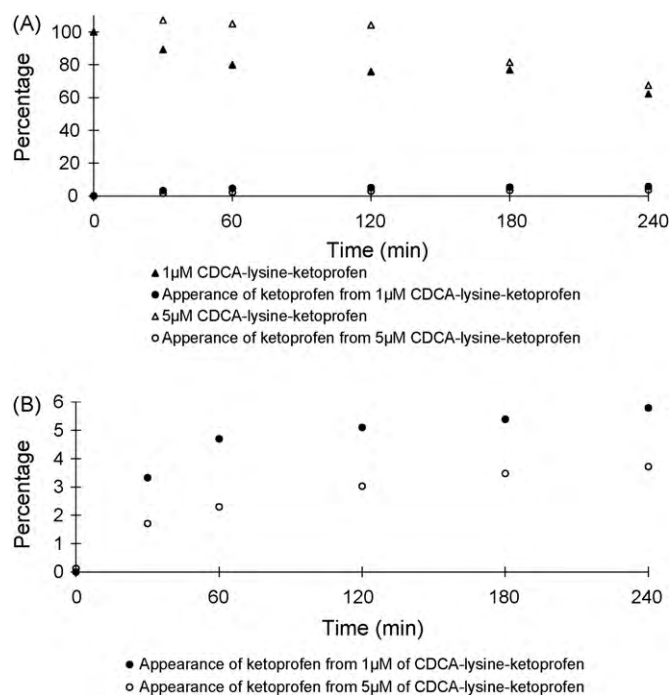


Fig. 5. Hydrolysis of CDCA-lysine-ketoprofen in Caco-2 homogenate. In panel A, data indicate the time course of the disappearance of 1 μM (\blacktriangle) and 5 μM (\triangle) of CDCA-lysine-ketoprofen and the appearance of ketoprofen from 1 μM (\bullet) and 5 μM (\circ) CDCA-lysine-ketoprofen. In panel B, ketoprofen appearance from 1 μM (\bullet) and 5 μM (\circ) prodrug is re-plotted on a smaller scale.

with a recovery efficiency over 97% (Dawson et al., 2009; Dawson and Oelkers, 1995; Wong et al., 1996).

In order to be efficiently enterohepatically recycled, high hepatic uptake and high secretion into bile are also critical which also

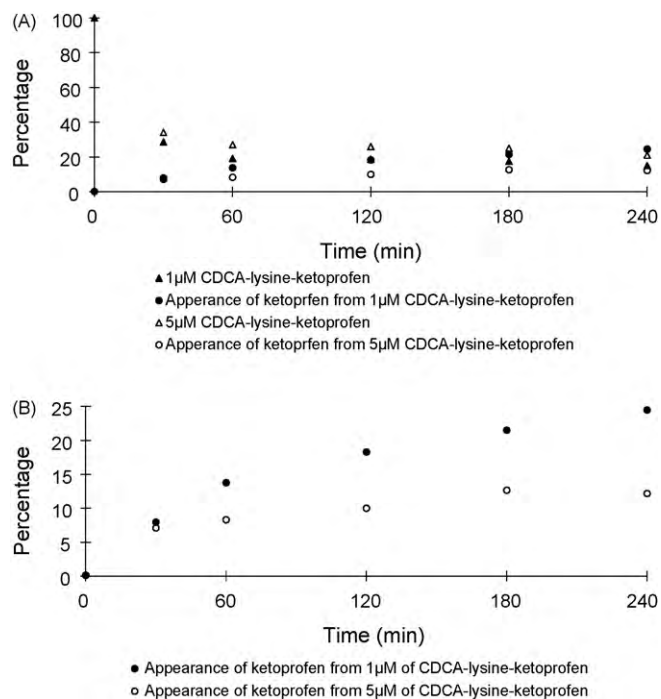


Fig. 6. Hydrolysis of potential prodrug CDCA-lysine-ketoprofen in rat liver homogenate. In panel A, data points indicate the time course of the disappearance of 1 μM (\blacktriangle) and 5 μM (\triangle) of CDCA-lysine-ketoprofen and the appearance of ketoprofen from 1 μM (\bullet) and 5 μM (\circ) CDCA-lysine-ketoprofen. In panel B, ketoprofen appearance from 1 μM (\bullet) and 5 μM (\circ) prodrug is re-plotted on a smaller scale.

involve transporters. Hepatic bile acid uptake is mainly mediated by sodium/taurocholate co-transporting polypeptide (NTCP), as well as members of the sodium independent organic anion-transporting polypeptides (OATPs) family (i.e. OATP-A, OATP-C, OATP-8) (St-Pierre et al., 2001). Bile salt exports pump (BSEP) and multidrug resistance protein (MRP2) transport bile acid into the bile canaliculus (Alrefai and Gill, 2007). MPR4 and MPR3 are able to remove bile acid through the sinusoidal surface (Alrefai and Gill, 2007). These bile acid transporters generally possess broader substrate specificity than ASBT. As a result, a bile acid conjugate that is well recognized by ASBT has high potential to be a substrate of liver bile acid transporters.

Although numerous transporter proteins are involved in the bile acid transport system, ASBT knockout mice exhibit 10–20-fold elevated fecal bile acid levels and a 80% decrease in bile acid pool size (Dawson et al., 2003). In humans, some inherited ASBT mutations cause primary bile acid malabsorption (Montagnani et al., 2001; Oelkers et al., 1997; Wong et al., 1995). These results indicate ASBT is critical for bile acid intestinal reabsorption. Bile acid–drug conjugates that utilize ASBT have potential to be absorbed from the intestine to the liver and excreted into the bile, to achieve a circulating reservoir of conjugated drugs. The reservoir may slowly release drug to provide sustained systemic drug concentrations.

4.2. Advantage to target ASBT

The conventional approaches for prolonged drug release are formulations or devices which provide a slow release/dissolution of drug within the intestine. Such approaches require the drug to be well absorbed in the large intestine where formulations will largely reside for 12–24 h after oral administration. Drugs that are amenable to conventional sustained release approaches must be highly permeable. Moreover, it is difficult to control drug release, and dose dumping can cause toxicity. Previously, prodrug approaches for the oral delivery of ketoprofen have employed polymer and dextran (Larsen et al., 1991; Wang et al., 2002). Prodrug approaches for niacin have entailed dextran and diacylglycerol esters (Cordi et al., 1995; Filip et al., 2003). Extended-release formulations of ketoprofen and niacin have been marketed.

The carrier used in the prodrug approach here was CDCA, with lysine as the linker. Lysine was selected since it is a natural amino acid and allows for CDCA and drug to be incorporated via amide linkages. It also provided a negative charge, which is critical for ASBT substrate translocation (Balakrishnan et al., 2006a). As a targeting moiety for ASBT as well as transporters involved in enterohepatic recirculation, CDCA is an endogenous compound, as is lysine, such that the prodrug has the potential advantage of low toxicity. In addition, as bile acids are distinctive physiological molecules and are poorly metabolized, a bile acid conjugate may show metabolic stability (Kramer and Wess, 1996).

Several other intestinal influx transporters have been proposed as prodrug targets, such as nucleoside transporters (ENTs), organic cation transporters (OCTs), organic anion-transporting polypeptides (OATPs), and peptide transporter (Pept1). Pept1 translocates valacyclovir and increase acyclovir bioavailability 3-fold (Burnette and de, 1994). K_i of valacyclovir is 4.08 mM in *Xenopus laevis* oocytes (Balimane et al., 1998) and 1.10 mM in CHO-PePT1 (Han et al., 1998). In comparison to this peptide prodrug, the bile acid prodrug approach that targets ASBT has the advantage of higher binding affinity (i.e. micromolar range).

4.3. Bile acid prodrugs

Several bile acid conjugates were previously designed to provide site-directed liver and gallbladder delivery (Kramer et al., 1994, 1992; Petzinger et al., 1995, 1999). For example, bile acid conjugates

of a HMG-CoA reductase inhibitor were active against HMG-CoA reductase in rat liver after an oral administration (Kramer et al., 1994). Also, a bile acid conjugate increased acyclovir bioavailability by 2-fold in rat (Tolle-Sander et al., 2004).

In the present study, both prodrugs were ASBT substrates, with high affinity and low capacity for the niacin prodrug and moderate affinity and high capacity for the ketoprofen prodrug. Transporter-mediated translocation is a complex process. It minimally requires substrate binding to the transporter, translocation of substrate via transporter conformation changes, dissociation of substrate, and transporter conformation restoration. Additional steps involve ASBT cotransport of sodium. Regarding the marked difference in transport capacity between two prodrugs, any step in the transporting process could be rate limiting, and consequently limit transporter capacity. However, since both prodrugs were potent inhibitors, substrate binding to the transporter was not the rate-limiting step, even for the niacin prodrug, for which ASBT exhibited a lower capacity.

An appropriate hydrolysis rate would be critical for prolonged release of drug from prodrug. If hydrolysis is too rapid, then prolonged release is not achieved. If hydrolysis is too slow, therapeutic concentrations of drug are not obtained. *In vitro* results suggest slow hydrolysis of niacin prodrug in Caco-2 and liver homogenates. Considering niacin's half-life of 20–45 min (Porter and Kaplan, 2009), the prodrug showed potential to target ASBT and yield a prolonged pharmacokinetic profile, which may cause levels of the flushing side effect. Compared to the niacin prodrug, the ketoprofen prodrug was a weaker substrate for ASBT and exhibited greater metabolic stability in Caco-2 homogenate. The conversion of prodrug to ketoprofen was achieved slowly in liver homogenate. Enzymatic stability results suggest that the prodrugs may be hydrolyzed gradually *in vivo* after absorption via ASBT.

For both prodrugs, further examination is warranted, in terms of these bile acid conjugates being substrates for other transporters that contribute to enterohepatic recirculation, as well as *in vivo* studies to assess potential for extended pharmacokinetic drug profile after prodrug administration.

5. Conclusions

Bile acid conjugates of each niacin and ketoprofen were synthesized. Both prodrugs exhibited binding affinity for ASBT. Niacin prodrug showed higher ASBT substrate binding affinity, but lower substrate capacity compared to taurocholate. Ketoprofen prodrug was a weaker substrate. Stability results suggest that each prodrug exhibits potential enzymatic stability for the conjugates to provide sustained drug release. Overall, *in vitro* results showed that these bile acid conjugates are potential prolonged release prodrugs with binding affinity for ASBT. Future *in vivo* pharmacokinetic assessment of the conjugates is warranted.

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