

5-BENZYLBARBITURIC ACID DERIVATIVES, POTENT AND SPECIFIC INHIBITORS OF URIDINE PHOSPHORYLASE

FARDOS N. M. NAGUIB,*† DIANE L. LEVESQUE,‡ ENG-CHI WANG,§
RAYMOND P. PANZICA‡§ and MAHMOUD H. EL KOUNI*

*Department of Pharmacology and Comprehensive Cancer Center, University of Alabama at Birmingham, Birmingham, AL 35294, U.S.A.; ‡Departments of Medicinal Chemistry and Chemistry, University of Rhode Island, Kingston, RI 02881, U.S.A.; and §School of Chemistry and Pharmacy, Kaohsiung Medical College, Kaohsiung City 80708, Taiwan, Republic of China

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Abstract—5-Benzylbarbituric acid derivatives were synthesized as a series of new, specific, and potent inhibitors of uridine phosphorylase. Among these, 5-(*m*-benzyloxy)benzyl-1-[(2-hydroxyethoxy)methyl] barbituric acid (5-benzyloxybenzylbarbituric acid acyclonucleoside, BBBA) was found to be the most potent with K_i values of 1.1 ± 0.2 and 2.6 ± 0.3 nM with uridine phosphorylase from human and mouse livers, respectively. BBBA exhibited competitive inhibition with uridine phosphorylase from both human and mouse livers. The 5-benzylbarbituric acid derivatives are specific inhibitors of uridine phosphorylase, as they had no effect on thymidine phosphorylase (EC 2.4.2.4), thymidine kinase (EC 2.7.1.21), uridine-cytidine kinase (EC 2.7.1.48), orotate phosphoribosyltransferase (EC 2.4.2.10), orotidine 5'-monophosphate decarboxylase (EC 4.1.2.23), and dihydrouracil dehydrogenase (EC 1.3.1.2). These compounds are more potent, easier to synthesize, and have better water solubility than their uracil counterparts as inhibitors of uridine phosphorylase. Furthermore, the 5-benzylbarbituric acids were found to be better inhibitors of human uridine phosphorylase than the murine enzyme, whereas the reverse holds true for the 5-benzyluracil derivatives. The 5-benzylbarbituric acid derivatives have potential usefulness in the therapy of cancer and AIDS, as well as other pathological and physiological disorders.

Two distinct pyrimidine nucleoside phosphorylases occur in mammalian cells [1, 2], uridine phosphorylase (UrdPase, EC 2.4.2.3) and thymidine phosphorylase (dThdPase, EC 2.4.2.4), which catalyze the general reaction:

Pyrimidine (deoxy)riboside + $P_i \rightleftharpoons$

Pyrimidine base + (deoxy)ribose-1-P

UrdPase from most mammalian sources primarily

cleaves pyrimidine (except cytosine) ribosides but is relatively non-specific, since it also cleaves pyrimidine 2'- and 5'-deoxyribosides [1–14]. dThdPase, on the other hand, seems to be specific for pyrimidine 2'- and 5'-deoxyribosides [3, 4, 9–12, 15–20].

UrdPase plays a crucial role in the chemotherapy of cancer and AIDS. The importance of UrdPase in cancer chemotherapy stems from the fact that UrdPase is the enzyme responsible for the activation and deactivation of several chemotherapeutic analogues, most notably the 5-fluoropyrimidines: 5-fluorouracil (FUra), 5-fluorouridine (FUrd), 5-fluoro-2'-deoxyuridine (FdUrd), and 5-fluoro-5'-deoxyuridine (5'-dFUrd) [3, 7, 9–12, 14, 16, 21–23], as many human tumours are apparently devoid of dThdPase activity [1, 3, 11, 15, 24–31]. Moreover, UrdPase, but not dThdPase, exhibits a circadian rhythm in mice [32, 33], which is opposite to that observed for the anticancer efficacy of FdUrd [34]. In addition, host-toxicities of some of these anticancer (e.g. FUra) [35–38], as well as anti-HIV (e.g. 3'-azido-3'-deoxythymidine, AZT) [39, 40] drugs are antagonized by uridine, the availability and concentration of which are controlled by UrdPase [32, 41–45]. The detrimental role played by UrdPase in reducing the effectiveness of various pyrimidine analogues in the chemotherapy of cancer and AIDS has generated a strong interest in developing inhibitors for this enzyme. Such inhibitors would enhance the chemotherapeutic efficacy of these drugs by preventing their degradation and/or host-toxicity.

† Corresponding author: Dr. Fardos N. M. Naguib, Department of Pharmacology, The University of Alabama at Birmingham, Birmingham, AL 35294. Tel. (205) 934-1131; FAX (205) 934-8240.

|| Abbreviations: AZT, 3'-azido-3'-deoxythymidine; BAU, benzylacetyluridine or 5-benzyl-1-[(2-hydroxyethoxy)methyl]uracil; BB, 5-benzylbarbituric acid; BBAU, 5-benzyloxybenzylacetyluridine or 5-(*m*-benzyloxy)benzyl-1-[(2-hydroxyethoxy)methyl]uracil; BBB, 5-benzyloxybenzylbarbituric acid or 5-(*m*-benzyloxy)benzylbarbituric acid; BBBA, 5-benzyloxybenzylbarbituric acid acyclonucleoside; or 1-[(2-hydroxyethoxy)methyl]-5-(*m*-benzyloxy)benzylbarbituric acid; BBBA acetate, 5-benzyloxybenzylbarbituric acid acyclonucleoside acetate or 1-[(2-acetoxyethoxy)methyl]-5-(*m*-benzyloxy)benzylbarbituric acid; AM-BBAU, aminomethyl-BBAU or 5-(*m*-benzyloxy)benzyl-1-[(1-aminomethyl-2-hydroxyethoxy)methyl]uracil; BBU, 5-benzyloxybenzyluracil; BU, 5-benzyluracil; DTT, dithiothreitol; 5'-dFUrd, 5-fluoro-5'-deoxyuridine; dThdPase, thymidine phosphorylase; FdUrd, 5-fluoro-2'-deoxyuridine; FUra, 5-fluorouracil, FUrd, 5-fluorouridine; P_i , orthophosphate; and UrdPase, uridine phosphorylase.

5-Benzylacetyluridines were developed as specific inhibitors of UrdPase [4, 10, 13, 14, 46–50]. They were shown to potentiate the efficacy of FdUrd *in vitro* and *in vivo* [14, 46, 47], to increase the level and duration of uridine in plasma [41–45] and heart-perfusate [51], to enhance the salvage of uridine by various tissues [42–45], and to reduce host-toxicity of FUra [42, 44, 45], FdUrd [52], and AZT [53, 54]. Yet, these inhibitors have less than favorable chemical and pharmacological properties. For example, the potency and/or poor water solubility of the benzylacetyluridines may limit their clinical usefulness. Furthermore, the more potent derivatives, i.e. AM-BBAU (aminomethyl analogue of 5-benzoyloxybenzylacetyluridine), were toxic (unpublished observation). In this paper we report the synthesis and testing of certain 5-benzylbarbituric acid derivatives as specific and potent inhibitors of UrdPase. Compared with their respective 5-benzyluracil counterparts, they are far more potent, have better water solubility and are easier to synthesize. A preliminary report has been presented [55].

MATERIALS AND METHODS

Chemicals

[2-¹⁴C]Uridine (56 Ci/mol), [2-¹⁴C]thymidine (56 Ci/mol), and [6-¹⁴C]uracil (56 Ci/mol) were obtained from Moravsek Biochemicals, Inc., Brea, CA; [6-¹⁴C]orotate (46.9 Ci/mol) was from NEN Research Products, the DuPont Co., Boston, MA; silica gel G/UV₂₅₄ polygram, polyethyleneimine cellulose 300 PEI/UV₂₅₄ and cellulose CEL 300 UV polygram thin-layer chromatography plates were from Brinkmann Instruments, Westbury, NY. The protein assay kit was obtained from Bio-Rad Laboratories, Richmond, CA. 5-Benzyluracil (BU), 5-benzoyloxybenzyluracil (BBU), 5-benzylacetyluridine (BAU), and 5-benzoyloxybenzylacetyluridine (BBAU) were synthesized and provided by Dr. S. H. Chu, Brown University, Providence, RI. All other chemicals were obtained from the Sigma Chemical Co., St. Louis, MO.

Chemical synthesis

Heterocycles. The 5-benzylbarbituric acid (BB) analogues were prepared from barbituric acid and the appropriate aldehyde as outlined in Scheme 1. The 5-arylidene intermediates and the target 5-arylalkylbarbiturates were obtained in good yields (ca. 90%). Their synthesis followed known methodology [56, 57]. Reduction of the 5-arylidenes was carried out in MeOH with NaBH₄ [58]. The known *m*-bromo- [59], *m*-chloro- [59], and *m*-nitro- [57] 5-benzylbarbituric acids (XVII–XIX) were prepared for this study as described above.

Acyclonucleosides: 1-[(2-Hydroxyethoxy)methyl]-5-(*m*-benzyloxy)benzylbarbituric acid or 5-benzoyloxybenzylbarbituric acid acyclonucleoside (BBBA, XXVI). BBBA was prepared by two different pathways as depicted in Scheme 2. The first pathway involved alkylation of persilylated barbituric acid (XX) with (2-acetoxyethoxy)methyl bromide [60] (XXI) in dry acetonitrile (CH₃CN) to provide 1-[(2-acetoxyethoxy)methyl]barbituric acid

(XXII) in 54% yield after purification by silica gel column chromatography. Deprotection of XXII with NaOMe furnished XXIII in near-quantitative yield. Either XXII or XXIII could be used in the next step leading to the substituted arylidene, but we selected XXII for ease of work-up due to better organic solubility. Reacting XXII with IV afforded the arylidene XXIV (71%) which was then reduced to BBBA acetate (XXV, 82%) followed by deprotection to provide BBBA (XXVI, 80%). Alternatively, BBBA (XXVI) could be prepared from the persilylated derivative (XXVII) of XVI. Treatment of XXVII with XXI in anhydrous 1,2-dichloroethane in the presence of a catalytic amount of AlCl₃ furnished BBBA acetate (50%) which after deprotection provided BBBA. Satisfactory elemental analyses and physicochemical data were obtained for all compounds. Complete experimental details will be published elsewhere.

Biological testing

Buffers

The following buffers were used: Buffer A [100 mM Tris-Cl, pH 7.5; 2 mM dithiothreitol (DTT); 5 mM MgCl₂] and Buffer B [20 mM potassium phosphate, pH 8.0; 1 mM DTT, 1 mM EDTA]. Buffer A was used when the activities of the nucleoside kinases were to be determined. Buffer B was employed when the activities of the other enzymes under study were to be determined.

Animal tissues

Human liver specimens were from donors and obtained through the Alabama Regional Organ Bank. The gall bladder was removed and the liver was sliced, perfused with cold normal saline and stored at –70° until used. Mouse livers were obtained from female Swiss Albino (CD1) mice weighing 20–24 g (Charles River Laboratories, Wellington, MA). Mice were killed by cervical dislocation and the livers removed.

Preparation of extracts

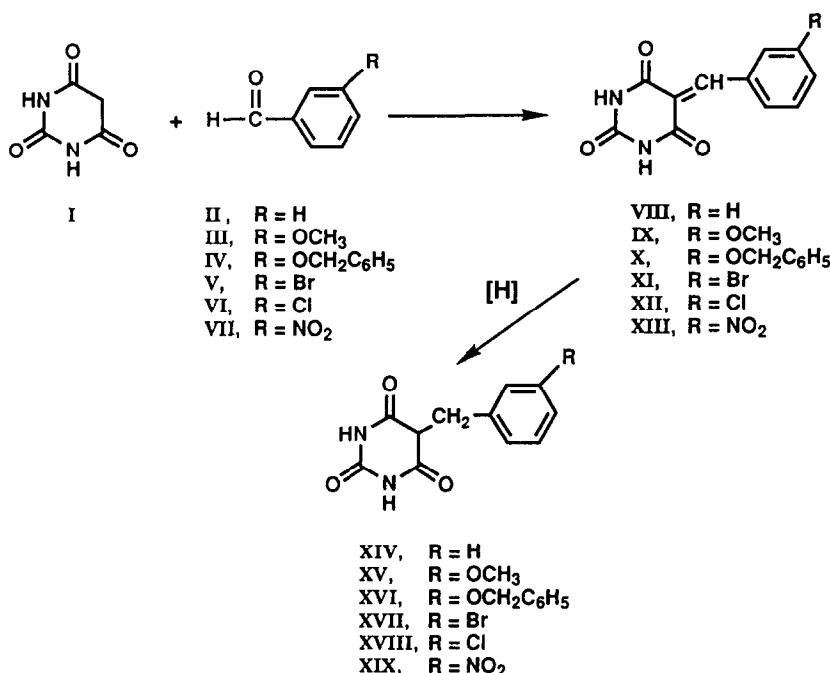
Organs were weighed, minced and homogenized in ice-cold (3:1, v/w) appropriate buffer using a Polytron homogenizer (Brinkman Instruments). The homogenates were centrifuged at 105,000 g for 1 hr at 4°. The supernatant fluids (cytosol) were collected and used as the enzyme source.

Enzyme assays

All assays were run at 37° under conditions where enzyme activity was linear with respect to time and enzyme concentration. For each inhibitor five concentrations were used ranging from 50 to 900 μM unless otherwise specified. Reactions were started by the addition of extract and stopped by boiling in a water bath for 2 min followed by freezing. Precipitated proteins were removed by centrifugation. Substrates were separated from products in the supernatant by TLC and the radioactivity in the spots was determined on a percentage basis using a Berthold LB-2821 Automatic TLC-Linear Analyzer.

Pyrimidine nucleoside phosphorylases. Nucleoside cleavage was measured isotopically by following the

Scheme 1



formation of nucleobases from their respective nucleosides as previously described [49]. The reaction mixture contained 20 mM potassium phosphate (pH 8), 1 mM EDTA, 1 mM DTT, 1 mM [2-¹⁴C]-uridine or [2-¹⁴C]thymidine (6 Ci/mol) and 25 μL cytosol in a final volume of 50 μL . The incubation was terminated after 30 min. Uridine and thymidine were separated from their respective nucleobases on silica gel TLC plates developed with chloroform:methanol:acetic acid (90:5:5, by vol.). The R_f values were: uridine, 0.07; uracil, 0.43; thymidine, 0.14; and thymine, 0.62.

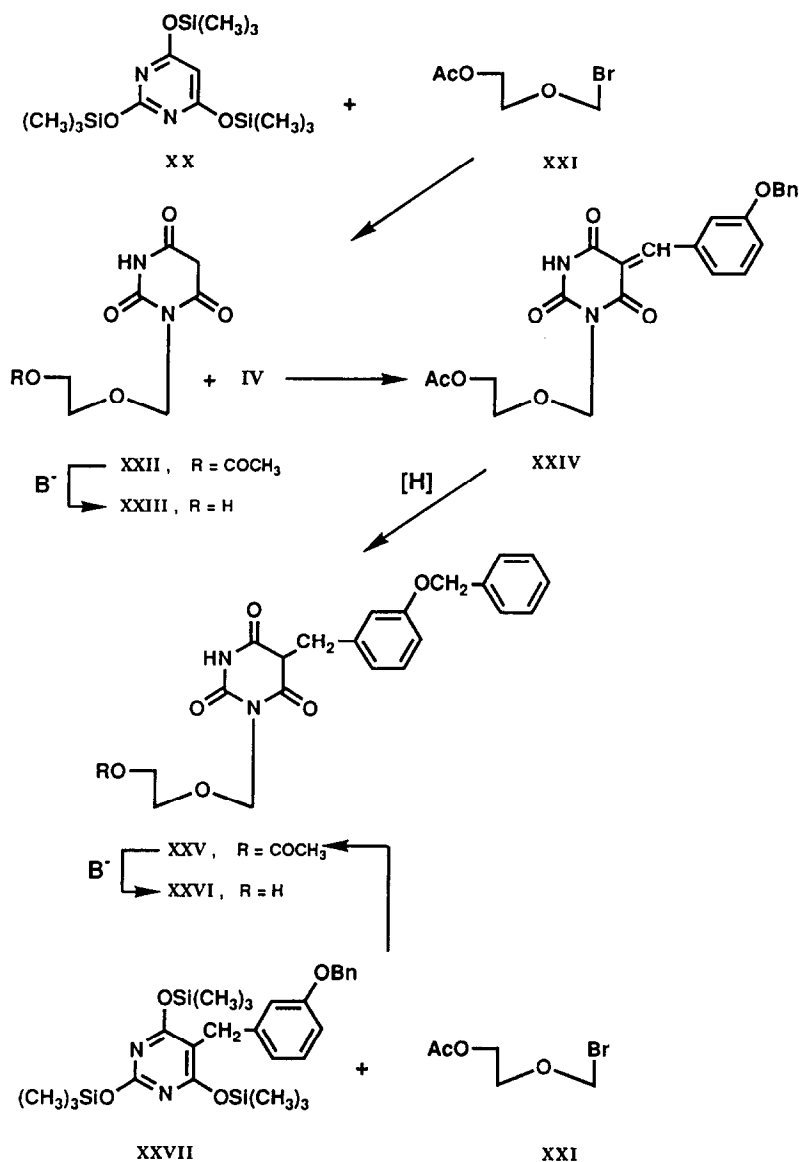
Dihydrouracil dehydrogenase (EC 1.3.1.2). Dihydrouracil dehydrogenase activity was measured by following the formation of dihydrouracil, carbamyl- β -alanine and β -alanine from [6-¹⁴C]uracil, as previously described [61]. The reaction mixture contained 20 mM potassium phosphate (pH 8), 1.0 mM EDTA, 2 mM DTT, 5 mM MgCl₂, 25 μM [6-¹⁴C]uracil (56 Ci/mol), 100 μM NADPH and 25 μL cytosol in a final volume of 50 μL . The incubation was terminated after 15 min. Uracil, dihydrouracil, carbamyl- β -alanine and β -alanine were separated on cellulose TLC plates developed in the top phase of a mixture of *n*-butanol:water:ammonia (90:45:15, by vol.). The R_f values were: dihydrouracil, 0.46; uracil, 0.23; and carbamyl- β -alanine, 0.09, respectively. Dihydrouracil dehydrogenase activity was determined as the sum of the products dihydrouracil, carbamyl- β -alanine and β -alanine.

Orotate phosphoribosyltransferase (EC 2.4.2.10) and orotidine 5'-monophosphate decarboxylase

(EC 4.1.2.23). Enzyme activity was measured by following OMP, orotidine, UMP, uridine and uracil formation from [6-¹⁴C]orotate [62]. The standard assay mixture contained 20 mM potassium phosphate buffer (pH 8.0), 2.5 mM 5-phosphoribosyl- α -1-pyrophosphate, 1 mM EDTA, 5 mM MgCl₂, 2 mM DTT, 0.05 mM (56.2 Ci/mol) [6-¹⁴C]orotate, and 25 μL cytosol extract, in a final volume of 50 μL . The incubation was terminated after 30 min. A 5- μL aliquot of the supernatant fluid was spotted on prewashed PEI-cellulose plates. The plates were developed first in distilled water to a front of 10 cm. They were then dried and redeveloped in 0.2 M LiCl. The R_f values were: OMP, 0.16; UMP, 0.51; orotate, 0.62; orotidine, 0.77; and uridine and uracil, 0.95. Orotate phosphoribosyltransferase activity was measured as the sum of OMP, orotidine, UMP, and uridine and uracil. Orotidine 5'-monophosphate decarboxylase activity was measured as the sum of UMP, uridine and uracil.

Pyrimidine nucleoside kinases. Thymidine kinase (EC 2.7.1.21) and uridine-cytidine kinase (EC 2.7.1.48) activities were determined isotopically as the sum of nucleotides formed from [2-¹⁴C]-thymidine and [2-¹⁴C]uridine, respectively [62]. The assay mixture contained 50 mM Tris-Cl (pH 7.4), 125 μM radiolabeled nucleoside (0.4 Ci/mol), 2.5 mM ATP, 5 mM MgCl₂, 25 mM NaF, and 30 μL cytosol extract in a final volume of 60 μL . After termination of the reaction and centrifugation, a 10- μL aliquot of the supernatant was spotted on silica gel TLC plates. The plates were developed with

Scheme 2



chloroform:methanol:acetic acid (56:24:4, by vol.). The R_f values were: thymidine, 0.77; thymidine nucleotides 0.15; uridine, 0.78; and uridine nucleotides, 0.15.

Kinetics studies

Determination and significance of apparent K_i values. Using uridine (1 mM) and five different concentrations of the inhibitor ranging from 50 to 900 μ M, apparent K_i values were estimated from Dixon plots ($1/v$ vs [I]) [63] of the data by a computer program with least squares fitting. The program was written by Dr. Sungman Cha and modified to fit IBM BASIC by Dr. F. N. M. Naguib. Apparent K_i

values are related to K_i values by the following equation:

$$\text{apparent } K_i = \frac{K_{is}(1 + [S]/K_m)}{1 + ([S]/K_m)(K_{is}/K_{ii})}$$

where K_{is} and K_{ii} are inhibition constants that would have been estimated from replots of slope and intercept, respectively, of a Lineweaver-Burk plot vs [I]. If a compound is a competitive inhibitor with respect to uridine, $K_{ii} = \infty$ and $K_{is} = K_i$. Therefore,

$$\text{apparent } K_i = K_i(1 + [S]/K_m).$$

Thus, for UrdPases from mouse and human livers

Table 1. Apparent K_i values for various 5-benzylbarbituric acid analogues in comparison with those of 5-benzyluracils and 5-benzylcyclohydridines for uridine phosphorylase from mouse and human livers

Inhibitor	Apparent K_i * (μ M)	
	Mouse	Human
Uracil	2,060 \pm 186	863 \pm 42
Acyclouridine	113 \pm 9.8	68.7 \pm 6.9
BU	288 \pm 47	275 \pm 68
BBU	66.4 \pm 11.3	96.4 \pm 4.6
BAU	3.12 \pm 0.21	3.97 \pm 0.30
BBAU	1.25 \pm 0.10	1.95 \pm 0.23
Barbituric acid (I)	19,810 \pm 10,680	
Barbituric acid acyclonucleoside (XXIII)	880 \pm 350	
BB, 5-Benzylbarbituric acid (XIV)	158 \pm 6.6	
5-(<i>m</i> -Nitrobenzyl)barbituric acid (XIX)	51.5 \pm 5.7	
5-(<i>m</i> -Methoxybenzyl)barbituric acid (XV)	26.5 \pm 0.9	
5-(<i>m</i> -Bromobenzyl)barbituric acid (XVII)	19.2 \pm 1.8	
5-(<i>m</i> -Chlorobenzyl)barbituric acid (XVIII)	12.4 \pm 1.2	2.14 \pm 0.26
BBB, 5-(Benzyloxybenzyl)barbituric acid (XVI)	11.7 \pm 0.5	2.77 \pm 0.53
BBBA, BBB acyclonucleoside (XXVI)	0.022 \pm 0.00	0.010 \pm 0.008
BBBA acetate (XXV)	7.0 \pm 0.3	
5-(<i>m</i> -Bromobenzylidene)barbituric acid (XI)	2,400 \pm 690	
5-(<i>m</i> -Chlorobenzylidene)barbituric acid (XII)	883 \pm 111	
5-(<i>m</i> -Nitrobenzylidene)barbituric acid (XIII)	321 \pm 45	

* Values are means \pm standard error of estimation from at least three determinations measured at 20 mM inorganic phosphate, 1.0 mM uridine and different concentrations of inhibitors.

which have K_m values of 143 and 242 μ M uridine, respectively [49], the apparent K_i of a competitive inhibitor, measured at a uridine concentration of 1 mM, is approximately 8- and 5-fold higher than their respective K_i values. It should be noted, however, that we have not characterized all of the compounds used in this study with regard to the type of inhibition (competitive, non-competitive or uncompetitive) or whether they are substrates for the enzyme.

Determination of K_i values. K_i values were estimated from the slopes and intercepts of the double-reciprocal plots by the methods of Wilkinson [64] and Cleland [65], using uridine concentrations ranging from 30 to 700 μ M.

Protein determination

Protein concentrations were determined spectrophotometrically by the method of Bradford [66] using bovine γ -globulin as a standard.

RESULTS

Inhibition constants and mechanisms of the 5-benzylbarbituric acid derivatives as inhibitors of UrdPase

Table 1 shows the apparent K_i values of various 5-benzylbarbituric acid derivatives, compared with those of the 5-benzyluracils (BU and BBU) and the 5-benzylacyclouridines (BAU and BBAU). Addition of a benzyl group to the 5-position of barbituric acid (5-benzylbarbituric acid, BB, XIV) increased the binding of barbituric acid to mouse liver UrdPase by 125-fold. Addition of a nitro, methoxy, bromo,

chloro, or benzyloxy group at the *meta* position of the benzyl group of BB further improved the binding of barbituric acid to UrdPase, by 385-, 745-, 1000-, 1600- and 1700-fold, respectively, with 5-benzyloxybenzylbarbituric acid (BBB, XVI) being the most potent inhibitor.

The binding of 5-benzyl- (BB, XIV) and 5-benzyloxybenzyl- (BBB, XVI) barbituric acids to mouse liver UrdPase was better than that obtained by their uracil counterparts (Table 1). BB had an apparent K_i value (158 μ M) 1.8-fold better than that of BU (288 μ M). Similarly, the apparent K_i value of BBB (11.7 μ M) was 6-fold superior to that of BBU (66.4 μ M). These results encouraged us to examine the effect of attaching the (2-hydroxymethoxy)methyl chain, i.e. the "acyclo" tail, to BBB (XVI) applying the same rationale that was used to enhance the binding of uracil, BU and BBU to UrdPase [3, 4, 10].

Addition of the (2-hydroxyethoxy)methyl "acyclo" tail enhanced the binding of barbituric acid (I) and BBB (XVI) to UrdPase from mouse liver by over 22- and 500-fold, respectively. The acyclo derivative of BBB, 5-benzyloxybenzylbarbituric acid acyclonucleoside (BBBA, XXVI), had an apparent K_i value of 0.022 μ M (Table 1). This apparent K_i was 57-fold better than that of BBAU (1.25 μ M) and 140-fold better than that of BAU (3.12 μ M) with UrdPase from mouse liver.

BBB and BBBA were also superior inhibitors of human UrdPase to their uracil counterparts, BBU and BBAU (Table 1). BBB inhibited human UrdPase 35-fold better than BBU. The acyclonucleoside, BBBA (apparent K_i value = 10 nM), was a 195- and 397-fold better inhibitor than BBAU (apparent K_i =

Table 2. Inhibitory potencies of 5-benzylbenzylbarbituric acid acyclonucleoside (BBBA) and 5-benzylacetyluridines (BAU and BBAU) for uridine phosphorylase from mouse and human livers

Inhibitor	K_i^* (nM)	
	Mouse	Human
BAU	420 \pm 40	1200 \pm 200
BBAU	170 \pm 0	220 \pm 20
BBBA (XXVI)	2.6 \pm 0.3	1.1 \pm 0.2

* Values are means \pm standard error of estimation measured at 20 mM inorganic phosphate, 30–700 μ M uridine and inhibitor concentrations ranging from 50 to 900 nM. The number of determinations for these values is found in Fig. 1.

1950 nM) and BAU (apparent K_i = 3970 nM), respectively. Furthermore, the inhibitory effects of BBB and BBBA were greater on human liver UrdPase than on mouse liver UrdPase. The apparent K_i of BBB (2.77 μ M) with the human UrdPase was 4-fold better than that obtained with mouse liver UrdPase (11.7 μ M). Similarly, BBBA (XXVI) had a 2-fold better apparent K_i (10 nM) with human UrdPase than with murine UrdPase (22 nM). This trend is the opposite of that found for the 5-benzyluracil derivatives, BBU, BAU and BBAU, which are better inhibitors of murine UrdPase than the human enzyme (Tables 1 and 2, and [13, 49]).

BBBA, the most potent inhibitor identified, was subjected to further kinetic studies to determine the mechanism of inhibition and the K_i value, and was also compared with BAU and BBAU. All three inhibitors, BBBA, BBAU and BAU, showed competitive inhibition with UrdPase from mouse and human livers. Figure 1 shows the Lineweaver-Burk plot of BBBA with human liver UrdPase.

BBBA had K_i values of 2.6 and 1.1 nM for UrdPase from mouse and human livers, respectively (Table 2). These K_i values were 65- and 200-fold better than those obtained with BBAU and over 160- and 1000-fold better than those obtained with BAU using mouse and human UrdPases, respectively. These results confirm that BBBA is a superior inhibitor to BAU and BBAU. They also demonstrate that, in contrast to BAU and BBAU, BBBA is a better inhibitor of the human UrdPase than the murine enzyme.

Inhibitor specificity

To check the specificity of the 5-benzylbarbituric acid analogues, these compounds were tested against various enzymes that could utilize the substrate (uridine or thymidine) or products (uracil or thymine) of UrdPase. Under our experimental conditions, the 5-benzylbarbituric acid analogues were neither inhibitors nor substrates for thymidine phosphorylase, uridine-cytidine kinase, thymidine kinase, dihydrouracil dehydrogenase, orotate phosphoribosyltransferase or orotidine 5'-monophosphate decarboxylase, as there was no significant effect on their activity in the presence of these

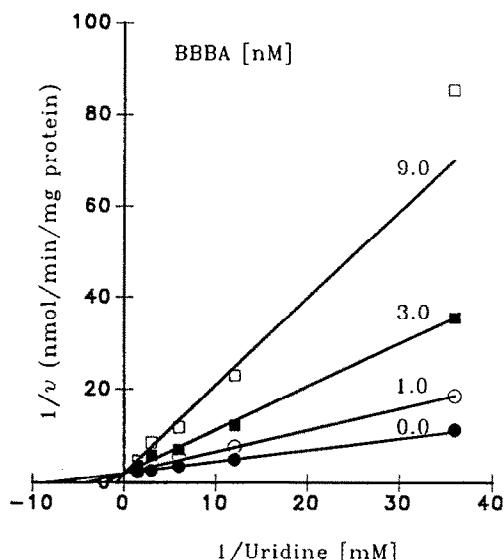


Fig. 1. Inhibition of human liver UrdPase by BBBA. Plots of $1/\text{velocity}$ versus $1/[\text{uridine}]$ at various fixed concentrations of BBBA. Each point represents at least three determinations.

compounds. These experiments indicate that the 5-benzylbarbituric acids are specific inhibitors of UrdPase.

Water solubility of 5-benzylbarbituric acid analogues

Table 3 shows the solubility of various 5-benzylbarbituric acid analogues when compared with those of BAU and BBAU. These compounds were at least 3-fold more soluble in water than either BAU or BBAU, with BBBA being 10-fold more soluble.

Lack of toxicity of BBBA

The toxicity of the most potent inhibitor among the various 5-benzylbarbituric acid analogues, BBBA, was tested in mice (CD1, Charles River Laboratories, Wellington, MA). Mice were injected i.p. with BBBA (50 mg/kg/day for 5 consecutive days) and monitored for 2 weeks. The mice did not show any mortality, and the mean body weight did not differ significantly from that of the controls (data not shown). Similar results were observed when BBBA (50 mg/kg/day) was injected i.v. into the tail vein of the mice.

DISCUSSION

In the present study we prepared from barbituric acid a series of new, highly specific and potent inhibitors of UrdPase. Among these, 5-benzyl-oxymethylbarbituric acid acyclonucleoside (BBBA, XXVI) has been identified as the most potent inhibitor of UrdPase with K_i values of 1.1 ± 0.2 and 2.6 ± 0.3 nM with UrdPase from human and mouse livers, respectively.

Although, barbituric acid (I) is a very poor

Table 3. Water solubility of 5-benzylbarbituric acid analogues and 5-benzylacynucleosides

Inhibitor	Solubility* (mM)
BAU	1
BBAU	<1
Barbituric acid (I)	>3
BB (5-Benzylbarbituric acid, XIV)	>3
5-(<i>m</i> -Methoxy)BB (XV)	>3
BBB (5-Benzylloxybenzylbarbituric acid, XVI)	>3
BBBA (BBB acynucleoside, XXVI)	>10

* In water at room temperature.

inhibitor of UrdPase (Table 1 and [10]), addition of a benzyl or a *meta*-substituted benzyl group to the C5 of the pyrimidine ring yielded extremely powerful inhibitors of UrdPase. Additional attachment of a (2-hydroxyethoxy)methyl "acyclo" tail to N1 further increased the potency of these inhibitors (Table 1). The data presented here together with results from previous studies [3, 4, 10, 13, 49, 50] allow us to postulate a mechanism(s) for the stronger binding of the 5-benzylbarbituric acid derivatives to the active site of UrdPase relative to their unsubstituted counterparts, barbituric acid (I) and its acynucleoside (XXIII). The fact that the most potent inhibitor among the N1-unsubstituted 5-benzylbarbituric acid derivatives was BBB (XVI) is consistent with the notion that increased hydrophobicity at the 5-position of the pyrimidine ring enhances binding of pyrimidines to UrdPase by interacting with a hydrophobic pocket on the enzyme adjacent to the 5-position of the pyrimidine ring [3, 4, 10, 13, 49, 50]. The poor binding of the 5-benzylidene derivatives of barbituric acids (XI, XII, XIII) as compared with their 5-benzyl counterparts (XVII, XVIII, XIX) (Table 1) could be explained by the relative rigidity of the 5-benzylidene group. The presence of the double bond between the C5 of barbituric acid and the benzyl methene would restrict the movement of the phenyl ring around the double bond and hinder its proper orientation to fit adequately into the hydrophobic pocket of the enzyme. The lack of proper orientation of substituents at C5 of the pyrimidine ring is probably also the main reason underlying the reported poor binding of the "classical" 5-disubstituted barbituric acid analogues (e.g. phenobarbital and pentobarbital) [10, 67]. The presence of two hydrophobic groups (e.g. phenyl and ethyl) at C5 would cause a steric hindrance at the hydrophobic pocket of the enzyme and reduce binding.

In addition to hydrophobicity, ionization of 5-benzylbarbituric acid derivatives may also play a role in enhancing the binding of these compounds to UrdPase. Niedzwicki *et al.* [10] have shown that addition of an electron-withdrawing group (e.g. nitro, bromo, chloro) at the 5-position of uracil enhances the binding of these analogues to UrdPase. Since such electron withdrawing groups enhance the acidity of these analogues, it was postulated that the

dissociated anionic forms bind more strongly to UrdPase than the uncharged species [10]. Under our assay conditions (pH 8), barbituric acid and its 5-substituted derivatives are most likely present as monoanions [10]. Addition of a benzyl group at the 5-position increases the acidity of 5-benzylbarbituric acid ($pK_a = 3.03$) over barbituric acid ($pK_a = 4.01$) by approximately 10-fold [59]. This acidity is further increased by addition of a nitro, bromo or chloro group at the *meta*-position of the benzyl group of 5-benzylbarbituric acid [59].

However, the decrease in pK_a values and increased hydrophobicity alone cannot explain the better binding of 5-benzylbarbituric acid (XIV) and its derivatives (XVI, XXVI) to UrdPase when compared with their uracil counterparts, particularly in view of the fact that the 5-unsubstituted barbituric acids (I, XXIII) bind weaker than their uracil analogues (Table 1). We postulate that the addition of a benzyl group at C5 of barbituric acid, may favor a tautomeric form of 5-benzylbarbituric acid derivatives with better binding capacity than their uracil analogues. In contrast to uracils, and, for that matter, 5-benzylidenobarbituric acid derivatives, as well as the "classical" 5-disubstituted barbiturates, where the monoanions are formed by the loss of a proton from N1 or N3 [10], evidence presently available suggests that the anions of 5-benzyl derivatives of barbituric acid (e.g. BBB, XVI), and presumably their acynucleosides (e.g. BBBA, XXVI), are formed by the loss of a proton from the C5 position of the pyrimidine ring with subsequent enolization of either the 4- or 6-position oxo-group [10, 68]. The charged oxygen probably enhances the binding of a preferred tautomer of the 5-benzyl derivatives of barbituric acid to UrdPase. Furthermore, the resulting negative charge localized on the exocyclic oxygen at C4 or C6 of the ionized 5-benzylbarbituric acid derivatives or their acynucleosides may also be involved in orienting and/or stabilizing the C5 benzyl group, and certain *meta*-substituents attached to the benzyl moiety, in a conformation better suited to fit into the adjacent hydrophobic pocket on the enzyme. Clearly, a better understanding of the binding of the 5-benzyl derivatives of barbituric acid to UrdPase must wait for more conclusive studies on the tautomeric equilibrium, ionization, and structural conformation of these compounds.

The enhanced binding of BBBA over BBB is probably due to the involvement of the hydroxyl group of the (2-hydroxyethoxy)methyl "acyclo" tail in hydrogen bonding to the second binding site on UrdPase. Similar multisubstrate binding was postulated for several acyclouridines [3, 4, 10, 49]. Therefore, the potency of BBBA can be increased further by additional chemical modification of the "acyclo" tail. This would include the addition of a terminal hydroxymethyl of an aminomethyl group to the "acyclo" tail, i.e. replacement of the (2-hydroxyethoxy)methyl "acyclo" tail with a (1,3-dihydroxy-2-propoxy)methyl or a (1-aminomethyl-2-hydroxyethoxy)methyl side chain. Such modifications increase the potency of BAU and BBAU by approximately 2.5- and 8.5-fold, respectively [49]. We are currently in the process of synthesizing and testing such derivatives.

BBBA has an excellent potential in various aspects of chemotherapy. The usefulness of UrdPase inhibitors has already been established in the field of experimental chemotherapy of cancer and AIDS. We have previously developed the 5-benzylacyclouridines as specific inhibitors of UrdPase [4, 10, 14, 46–50]. These inhibitors were shown to enhance the efficacy of FdUrd against human tumors *in vitro* and *in vivo* [14, 46, 47]. Furthermore, the benzylacyclouridines were shown to elevate the concentration and prolong the half-life of uridine in the plasma [41–45], as well as increase the salvage of uridine by various tissues [42, 44, 45]. Therefore, these inhibitors were used to protect against or rescue from host-toxicity of anticancer (i.e. FUra) [42, 44, 45] and anti-HIV (i.e. AZT) [52, 53] drugs, the toxicities of which were shown to be antagonized by administration of exogenous uridine [35–40]. The new acyclonucleoside BBBA is likely to share the various chemotherapeutic characteristics of the older inhibitors. BBBA, however, has certain advantages over the 5-benzylacyclouridines: (1) more potent; (2) cheaper and easier to make in large quantities; and (3) much more soluble in water. The last property is important in practical chemotherapy since the benzylacyclouridines are insoluble in water at the concentrations and volumes required to be administered to patients or experimental animals. Furthermore, the benzylbarbituric acid derivatives are, in general, better inhibitors of human UrdPase than the murine enzyme, whereas the reverse is true for the 5-benzyluracil derivatives. This is the first time an inhibitor of UrdPase is shown to be a better inhibitor of the human enzyme than its murine counterpart (Tables 1 and 2 [13, 49]).

Therefore, BBBA can be useful as a "protective" and/or "rescuing" agent against drug toxicity that can be rescued or prevented by uridine or similar nucleosides. In this respect, BBBA can substitute, totally or partially, for the "uridine rescue regimens" in treating cancer, viral, and other diseases. BBBA alone or in combination with low concentrations of uridine can maintain an adequate uridine concentration for a long enough time to rescue selectively the normal or uninfected cells from drug toxicity, without having to suffer from the toxic side-effects (e.g. phlebitis, pyrogenic reactions, changes in body temperature and diarrhea) observed with

the high concentration of uridine required to achieve the rescuing or protective effects in the absence of the inhibitor (43, 53, 69–72]. The use of BBBA in manipulating the uridine pool is not limited to the treatment of cancer or AIDS but can also be extended to treat other pathological and physiological disorders, where administration of uridine has been shown to be useful. Such disorders are quite numerous and include CNS disorders (e.g. cerebrovascular disorders and convulsions) [73–84], sleep promotion [85], muscle performance [86, 87], liver diseases [88–90], and cardiac damage [91–96]. The potential use of BBBA and other UrdPase inhibitors in treating such disorders has not yet been recognized in contemporary experimental or applied chemotherapy.

In conclusion, we have synthesized the 5-benzylbarbituric acids as specific and superior inhibitors of UrdPase to those previously known. These compounds could have a wide application in the therapy of cancer and AIDS, as well as other pathological and physiological disorders.

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