

# Sodium Dependence of Nitrofurantoin Active Transport across Mammary Epithelia and Effects of Dipyridamole, Nucleosides, and Nucleobases

Phillip M. Gerk,<sup>1</sup> Linda Hanson,<sup>2</sup> Margaret C. Neville,<sup>2</sup> and Patrick J. McNamara<sup>1,3</sup>

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**Purpose.** The sodium dependence and effects of nucleoside and nucleobase transport inhibitors were determined to ascertain the role of sodium dependent nucleoside or nucleobase transporters in nitrofurantoin active transport across mammary epithelia.

**Methods.** Five lactating female rats received steady-state intravenous infusions of nitrofurantoin with and without the broad-based inhibitor dipyridamole. In the CIT3 murine model of lactation, <sup>14</sup>C-nitrofurantoin basolateral to apical permeability was examined in the presence of varying sodium concentrations, purine and pyrimidine nucleosides and nucleobases, and dipyridamole.

**Results.** Dipyridamole effectively inhibited <sup>14</sup>C-nitrofurantoin flux across CIT3 cells, with  $K_i = 0.78 \mu\text{M}$  (95% C.I. = 0.11 to 5.3  $\mu\text{M}$ ) and significantly decreased the milk-to-serum ratio of nitrofurantoin from  $29.2 \pm 5.0$  to  $11.0 \pm 6.3$  without changing systemic clearance. Nitrofurantoin active transport was significantly inhibited by complete sodium replacement. Adenosine and guanosine significantly inhibited nitrofurantoin permeability ( $54.5 \pm 2.6$  ( $\mu\text{l/hr/cm}^2$ ) and  $50.7 \pm 0.6$  ( $\mu\text{l/hr/cm}^2$ ), respectively, vs. control  $90.5 \pm 4.6$  ( $\mu\text{l/hr/cm}^2$ ) but uridine, thymidine, and the nucleobases had no effect.

**Conclusions.** Nitrofurantoin active transport was sodium dependent and inhibited by dipyridamole, adenosine, and guanosine, but known sodium dependent nucleoside or nucleobase transporters were not involved.

**Key words:** nitrofurantoin; dipyridamole; nucleosides; nucleobases; sodium-dependence.

## INTRODUCTION

In an effort to predict neonatal exposure due to the transfer of medications into breast milk, a passive diffusional model has been developed previously (1). Assuming only unbound, unionized drug crosses the mammary epithelium, this model predicts milk to serum ratios (M/S) for many medications entering milk by passive diffusion (2). However, nitro-

furantoin and cimetidine are actively transported into both human and rat milk (3–6). Furthermore, nitrofurantoin, an antibiotic used to treat urinary tract infections, reaches M/S values in humans and rats of 20 and 100 times those predicted by diffusion, respectively (4–6). To make better M/S predictions, it is necessary to understand how xenobiotics such as nitrofurantoin are actively transported by the mammary gland into milk.

We observed that hypoxanthine inhibited the specific transport of nitrofurantoin across a confluent murine mammary epithelial cell (CIT3) monolayer (unpublished data). This observation suggested a role for nucleobase and/or nucleoside transport systems. Facilitative (sodium independent) and concentrative (sodium dependent) nucleoside and nucleobase transporters have been found in various epithelial and endothelial tissues of several different species (7–9). Therefore, sodium replacement and several inhibitors of nucleoside/nucleobase transport were tested for their effects on nitrofurantoin transport in the CIT3 system.

There are several types of nucleoside transporters (7–10). The equilibrative nucleoside transporters are distinguished by their sensitivity to nitrobenzylmercaptapurine riboside (nitrobenzylthioinosine). They are both sodium independent, with a broad specificity for purines and pyrimidines, and are sensitive to 10  $\mu\text{M}$  dipyridamole. Next, there are sodium-dependent secondary active nucleoside transporters, which are functionally designated N1–N6 and differ in their substrate and inhibitor selectivities, as shown in Table I (8–11). Nucleoside transporters having N1 and N2 functionality have been cloned and named CNT2 and CNT1, respectively (11). N1 and N6 prefer purine nucleosides, N2 and N4 prefer pyrimidine nucleosides, N3 accepts both, and N5 transports uridine (Table I). In contrast to N5 and N6, N1–N4 are insensitive to NBMPR (Table I). If a xenobiotic is concentrated into milk, the concentrative transporters N1–N6 would be more likely involved than the equilibrative transporters.

In regards to nucleobase transport, two equilibrative types have been identified for hypoxanthine (10). In addition, at least one  $\text{Na}^+$  dependent nucleobase transport system has been identified, for which hypoxanthine is a substrate and guanine, uracil, and thymine are inhibitors, as shown in Table II (12–16). Furthermore, in two studies hypoxanthine transport was insensitive to 100  $\mu\text{M}$  guanosine, uridine, and thymidine (12,17). Unlike the active nucleoside transporters, none of the nucleobase transporters have been cloned and functionally expressed, so functional expression systems are non-existent. Recently, nucleobase transporter-like cDNAs have been reported in humans, mice, and pigs, but so far have not been functionally expressed (18). These cDNAs may show promise as nucleobase transporters in future studies.

To examine the role of nucleoside and nucleobase transporters *in vivo* as well as *in vitro*, dipyridamole was chosen. Dipyridamole is a broadbased inhibitor of several types of nucleoside and nucleobase transporters, as well as organic cation and p-glycoprotein transporters (8,10,19,20). Also, if a sodium dependent transporter is involved, then the transport activity is expected to be sodium dependent. To determine the role of sodium dependent nucleoside or nucleobase transporters, the following algorithm was used, based on Tables I and II. If a known sodium dependent nucleoside transporter

<sup>1</sup> University of Kentucky College of Pharmacy, Division of Pharmaceutical Science Lexington, Kentucky 40536-0082.

<sup>2</sup> University of Colorado Health Science Center, Department of Physiology, Denver, Colorado 80262.

<sup>3</sup> To whom correspondence should be addressed. (e-mail: pmcnamar@uky.edu)

**ABBREVIATIONS:** M/S, milk to serum ratio; N1–N6, sodium dependent nucleoside transporters; B–A, basolateral to apical; Pe, effective permeability; J, flux; A, area;  $C_d$ , donor concentration;  $J_{\text{max}}$ , maximal flux; NS, nonspecific diffusional flux; S, substrate concentration;  $K_m$ , substrate Michaelis-Menten constant;  $\text{IC}_{50}$ , half-maximally inhibitory concentration;  $K_i$ , inhibitor Michaelis-Menten constant.

**Table I.** Sodium-Dependent Nucleoside Transporters

Subtype	Name designation	N1 <i>cif</i>	N2 <i>cit</i>	N3 <i>cib</i>	N4 <i>cit</i>	N5 <i>cs</i>	N6 <i>csg</i>
<b>Substrates</b> (Km, $\mu$ M)	adenosine	1-14					
	guanosine	4-11		30			T
	uridine	9-40	7-13	18	4-5	T	
	thymidine	NT	5-8	13	27		
<b>Inhibitors</b> (IC50, $\mu$ M)	adenosine	8	12-21	I, 100	I, 100	I, 1000	I, 1000
	guanosine	15	640	8-30	I, 100	I, 1000	I, 1000
	uridine	I	8-22	8-10	I, 100	I, 1000	I, 1000
	thymidine	NI	13	13-16	I, 100		NI, 1000
	NBMPR	NI	NI	NI	NI	I	0.0007
	dipyridamole	NI	NI	NI	NI	I	I, 1000
	hypoxanthine						I, 1000
	guanine						I, 1000

These data are adapted from references (8-11) where several different studies are summarized.

T indicates agent is transported, NT indicates agent is not transported.

I indicates agent is inhibitory, or NI indicates not inhibitory, and the indicated concentration ( $\mu$ M).

Blanks indicate undetermined data.

is involved, then both adenosine and uridine should inhibit nitrofurantoin transport, as the functionally classified N1-N6 sodium dependent nucleoside transporters are inhibited by both (8-11). If a known sodium dependent nucleobase transporter is involved, then guanine and/or uracil should inhibit nitrofurantoin transport (12-15,21).

The purpose of the present research was to determine the role of the sodium dependent nucleoside and nucleobase transporters in the active transport of nitrofurantoin across mammary epithelia by determining the effects of sodium replacement and inhibitors of nucleoside and nucleobase transporters on the process.

## MATERIALS AND METHODS

### Chemicals

$^{14}$ C-nitrofurantoin (58 mCi/mmol) was obtained from Chemsyn Labas (Lexena, Kansas). Polyethylene glycol 400 (PEG 400) was obtained from Union Carbide Chemical Plastic Co., Danbury, Connecticut. Organic solvents were obtained from Fisher (Pittsburgh, Pennsylvania). Cell culture reagents were obtained from Gibco BRL (Rockville, Maryland) except for hydrocortisone and insulin (Sigma, St. Louis, Missouri), epidermal growth factor (Collaborative Biomedical Products, Bedford, Massachusetts), ovine prolactin (National Hormone and Pituitary Program, Torrance, Califor-

nia), fetal bovine serum (Gemini Bio-Products, Calabasas, California), sterile water for injection (Baxter, McGaw Park, Illinois), and compressed gases (Scott-Gross, Lexington, Kentucky). Agents used in transport studies (Fig. 1) and all other chemicals were obtained from Sigma.

### In Vivo Infusions

This research adhered to the "Principles of Laboratory Animal Care" (NIH publication #85-23, revised 1985) and protocol 97-0052M was approved by the University of Kentucky Institutional Animal Care and Use Committee. In a randomized crossover study, five 300 g Sprague-Dawley lactating female rats 12-15 days post-partum were cannulated under ketamine/xylazine anesthesia. After a day to recover from surgery, the rats were randomized to receive intravenous infusions of nitrofurantoin 0.4 mg/h either with or without dipyridamole 12 mg/h. The delivery vehicle was neat polyethylene glycol 400 (0.25 ml/h) and hourly intravenous injections of 1 ml normal saline were given to prevent dehydration. Blood samples were obtained at 3, 4, and 5 h after the start of the infusion and serum was harvested by centrifugation. Milk samples were obtained at 5 h under light ketamine/xylazine anesthesia. The rats were crossed over the following day to receive the other treatment. All samples were protected from light, aliquoted, and stored frozen ( $-20^{\circ}$ C) until analysis.

Serum and milk samples were extracted and analyzed by HPLC using a modified method of Pons *et. al.* (22). Briefly, furazolidone (25  $\mu$ l of 5  $\mu$ g/ml) was added as an internal standard to each 50  $\mu$ L aliquot of serum or milk. The sample was acidified and protein precipitated by adding 100  $\mu$ L of 1 N HCl and vortexing for 30 s. The sample was extracted into 1 ml methylene chloride, vortexed, and centrifuged. The organic layer was decanted, evaporated under nitrogen, and reconstituted with mobile phase. The sample was injected onto a Shimadzu (Kyoto, Japan) HPLC system with a C-18 column (Lichrosorb 5 RP18 125  $\times$  4.0 mm, Phenomenex, Torrance California) and eluted with 10% acetonitrile 90% 25 mM potassium phosphate buffer (pH 3.00) at 1.0 ml/min. UV absorbance was measured at 366 nm. Peak height ratios (ni-

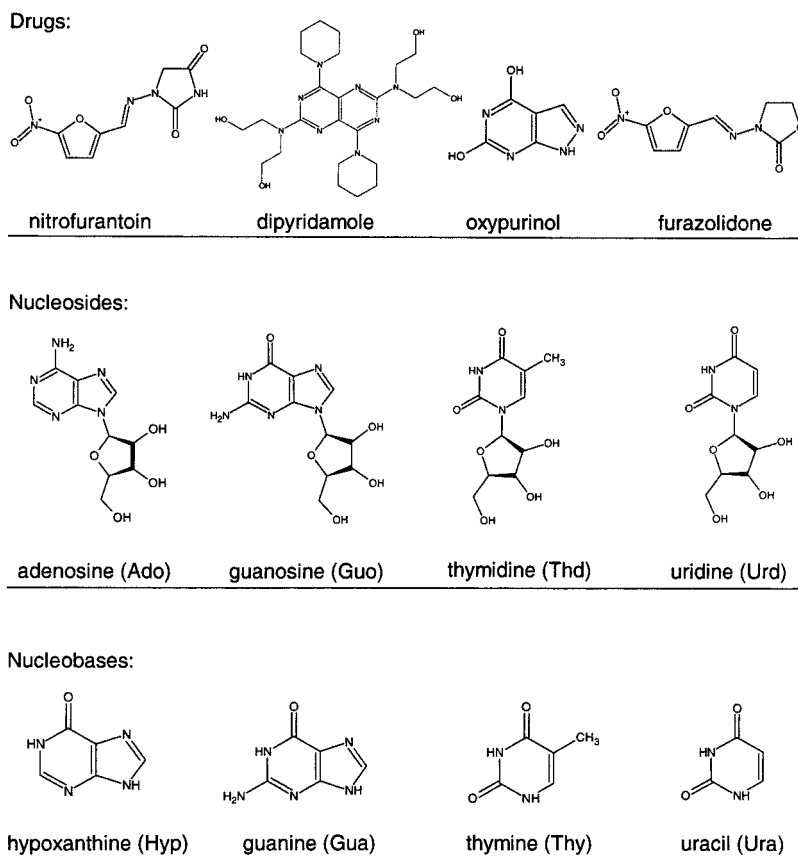
**Table II.** Sodium-Dependent Nucleobase Transport

	heart muscle	LLC-OK cells	renal PK1	renal BBMV	Choroid plexus	jejunum
<b>hypoxanthine</b>	8.1*	0.78*	0.79*	4.4*	31*	
<b>guanine</b>	6.4	0.6	5.8	1.8	I	I
<b>thymine</b>	5.5	4.2	3.5	7.6	I	21
<b>uracil</b>	7.2	3.9	2.9	30	64.6	47*
<b>dipyridamole</b>	30.2	0.9	4.6			

Note: This table adapted from reference (16).

Km (\*) or Ki values are given in  $\mu$ M.

I indicates inhibitory, NI indicates not inhibitory at the indicated concentration.



**Fig. 1.** Structures of agents used in transport studies. IUPAC-IUB abbreviations for nucleosides and nucleobases are in parentheses.

trofurantoin/furazolidone) were used for comparison with the standard curve.

### Cell Culture Studies

CIT3 cells were cultured according to the previously published protocol (23). Briefly, the cells (passages 15 to 21) were grown in 50:50 DMEM/Ham's F12 media supplemented with 2% fetal bovine serum, epidermal growth factor (5 ng/ml), insulin (10  $\mu$ g/ml), penicillin (100 U/ml), and streptomycin (100  $\mu$ g/ml). The cells were seeded on #3407 Snapwell polycarbonate 0.4  $\mu$ m, 1 cm<sup>2</sup> filter inserts (Corning-Costar, Corning, New York), then differentiated by removing epidermal growth factor and adding prolactin (3  $\mu$ g/ml) and hydrocortisone (3  $\mu$ g/ml).

All experiments were conducted in a buffer or medium free of antibiotics, serum, hormones, or proteins. Three types of solutions were used in the experiments. Early studies were performed in DMEM/F12 cell culture media (Gibco BRL) containing 14.3 mM NaHCO<sub>3</sub> and 10 nM HEPES at pH 7.4. Because the commercial medium contains hypoxanthine, thymidine, and other potentially interfering compounds, saline buffers were used for subsequent studies. Sodium replacement studies were conducted in Ringer's solutions containing 113 mM NaCl, 5 mM KHCO<sub>3</sub>, 2 mM CaCl<sub>2</sub>, 1 mM MgSO<sub>4</sub>, 20 mM glucose, 10 mM HEPES, and 23  $\mu$ M phenol red, pH adjusted to 7.4. NaCl was replaced with either lithium chloride or glucosamine hydrochloride. Finally, all other studies

were performed with an isotonic saline buffer containing 130 mM NaCl, 4.2 mM KHCO<sub>3</sub>, 1.05 mM CaCl<sub>2</sub>, 0.41 mM MgSO<sub>4</sub>, 0.14 mM MgCl<sub>2</sub>, 20 mM glucose, 10 mM HEPES, adjusted to pH = 7.4 bubbled to equilibrium with 95% oxygen 5% carbon dioxide. The calculated osmolarity was adjusted to 280-305 mOsm/L with mannitol. Humidified 95% oxygen 5% carbon dioxide was bubbled through both sides of each diffusion chamber at 20-25 ml/min, except air was bubbled in experiments where Ringer's solutions were used as above.

Before the experiments, <sup>14</sup>C-nitrofurantoin was purified by silica gel TLC (methylene chloride:acetone 3:1) and its radiochemical purity was determined by injecting an aliquot onto the HPLC system described above, sampling and counting the effluent to verify purity  $\geq$ 97%. The diffusion chambers were assembled with snapwells, filled with 5 ml on each side, and maintained at 37°C. Transepithelial resistances were measured with a Millicell meter (Millipore Corp., Bedford, Massachusetts) in each assembled diffusion chamber before, and at the end of the experiment. Resistances <300 ohm\*cm<sup>2</sup> were cause for exclusion of that chamber. <sup>14</sup>C-Nitrofurantoin (0.4  $\mu$ Ci) was spiked to the basolateral side. Samples (100  $\mu$ l) were obtained from the basolateral and apical chambers at 1, 20, 40, 60, 80, 100, and 120 min and were analyzed by mixing with 3a70B liquid scintillation cocktail (Research Products Inc., Mt. Prospect, Illinois) and counting on a Tri-Carb 2200CA (Packard Instrument, Meriden, Connecticut). Basolateral to apical (B-A) permeability was determined as in

Data Analysis below. Mass balance was complete, and less than 5% of the radiolabel was transferred to the opposite side in 120 min.

### CIT3 Study Designs

A saturation study on the B-A permeability of nitrofurantoin was performed in DMEM/F12 using total nitrofurantoin concentrations ranging from 1 to 1600  $\mu\text{M}$ . The results were fitted to the Michaelis-Menten equation by nonlinear regression (see Data Analysis).

The initial sodium dependence study was performed in Ringer's solutions isotonicly replacing NaCl and LiCl or glucosamine HCl in the absence or presence of 250  $\mu\text{M}$  cold nitrofurantoin and determining B-A permeability.

The first studies using adenosine, guanosine, hypoxanthine (500  $\mu\text{M}$  each), and the ATP synthase inhibitor oligomycin (10  $\mu\text{M}$ ) were performed in DMEM/F12, comparing the B-A permeability of nitrofurantoin in control (no inhibitor) vs. treated groups. A further characterization was done with adenosine, guanosine, thymidine, uridine, hypoxanthine, guanine, thymine, uracil, and oxypurinol (100  $\mu\text{M}$  each) in saline buffer. To determine the  $K_i$  value for dipyridamole, the B-A permeability of  $^{14}\text{C}$ -nitrofurantoin was determined at dipyridamole concentrations from 0.03  $\mu\text{M}$  to 100  $\mu\text{M}$  (see Data Analysis).

### Data Analysis

In the infusion studies, systemic clearance was determined by dividing the infusion rate by the steady-state serum concentrations. The observed milk to serum ratio (M/S) was calculated by dividing the concentration in milk by the steady state serum concentration.

In the CIT3 studies, effective permeability ( $P_e$ , ( $\mu\text{l/h}$ )/ $\text{cm}^2$ ) was determined by calculating flux ( $J$ , pmol/hr) and dividing by the area ( $1\text{ cm}^2$ ) and the donor concentration ( $C_d$ ,  $\mu\text{M}$ ) as given in Eq. (1) below. Sink conditions (no change in  $C_d$ ) were assumed. GraphPad Prism v. 2.01 (GraphPad Software, Inc., San Diego, California) was used for non-linear regression. The Michaelis-Menten equation [Eq. (2)] was fitted by nonlinear regression, where  $J$  is the observed flux,  $J_{\text{max}}$  is the maximal flux,  $S$  is the concentration of nitrofurantoin,  $K_m$  is the Michaelis-Menten constant, NS is a nonspecific diffusional flux.

$$J = A * P_e * C_d \quad (1)$$

$$J = \frac{J_{\text{max}} - \text{NS}}{1 + 10^{(\log S - \log K_m)}} + \text{NS} \quad (2)$$

Data from the dipyridamole inhibition experiment were fitted to Eq. (3) below to obtain  $\text{IC}_{50}$ , then equation 4 was solved to obtain  $K_i$  for dipyridamole. In Eqs. (3) and (4),  $J$  is the observed flux of nitrofurantoin in the presence of dipyridamole,  $J_{\text{max}}$  is the flux of nitrofurantoin in the absence of dipyridamole,  $I$  is the concentration of dipyridamole, NS is a nonspecific diffusional flux (determined above),  $S$  is the concentration of nitrofurantoin,  $\text{IC}_{50}$  is the concentration of the inhibitor which inhibits half of the active component of the observed flux, and  $K_m$  is the Michaelis-Menten constant for nitrofurantoin transport.

$$J = \frac{J_{\text{max}} - \text{NS}}{1 + 10^{(\log I - \log \text{IC}_{50})}} + \text{NS} \quad (3)$$

$$K_i = \frac{\text{IC}_{50}}{1 + \left(\frac{S}{K_m}\right)} \quad (4)$$

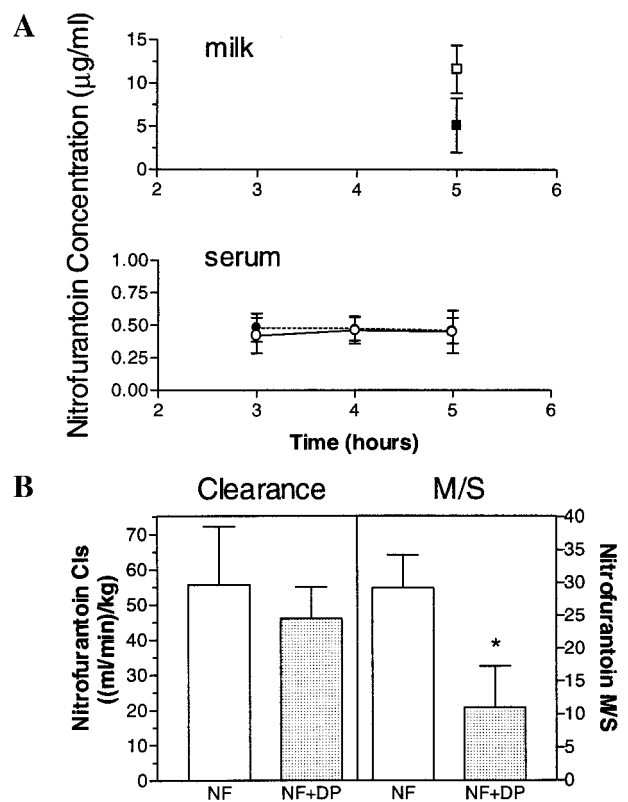
### Statistical Analysis

Means for M/S and systemic clearance from the rat infusion study were compared by paired  $t$ -tests, with  $\alpha = 0.05$ . Comparisons in the CIT3 determinations were made by one-way ANOVA with Dunnett's post-hoc test ( $\alpha = 0.05$ ) using GraphPad Prism v. 2.01.

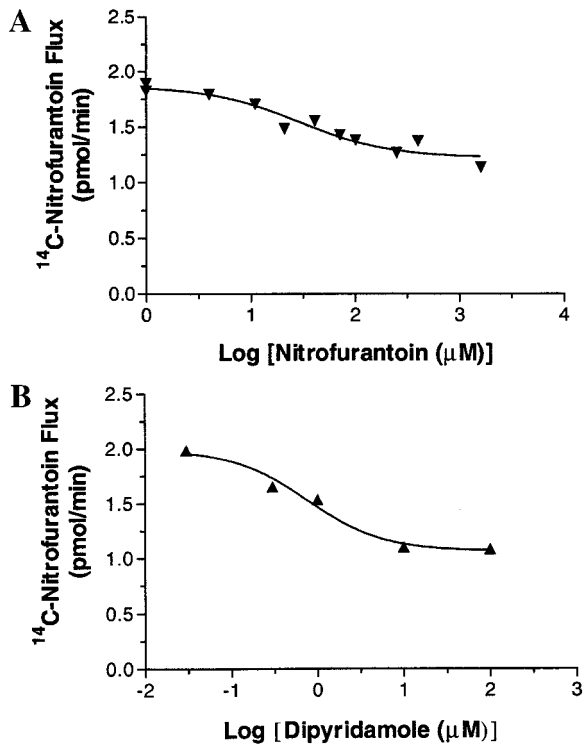
### RESULTS

Results from the infusion study in lactating rats are shown in Fig. 2. Figure 2a shows that nitrofurantoin serum concentrations reached a steady-state at 3 h that was maintained through 5 h. The systemic clearance of nitrofurantoin was not changed by co-administration of dipyridamole ( $55.7 \pm 16.6$  (ml/min)/kg vs.  $46.0 \pm 9.1$  (ml/min)/kg) (mean  $\pm$  S.D.) as shown in Fig. 2b. However, dipyridamole significantly decreased the M/S of nitrofurantoin from  $29.2 \pm 5.0$  to  $11.0 \pm 6.3$ , a 62% decrease in M/S (Fig. 2b). Both infusion regimens were well tolerated by the rats, and little or no hemolysis was observed.

Figure 3a shows the effect of increasing concentrations of cold nitrofurantoin on the flux of  $^{14}\text{C}$ -nitrofurantoin across



**Fig. 2.** Five lactating rats were infused with nitrofurantoin 0.4 mg/h alone (NF) or with dipyridamole 12 mg/h (NF+DP). Data are mean  $\pm$ SD. **(A)** Milk and serum steady-state concentrations of nitrofurantoin. Milk concentrations are indicated by squares, serum concentrations are indicated by circles; filled symbols indicate the presence of dipyridamole. **(B)** Effects of dipyridamole on the M/S and systemic clearance of nitrofurantoin 0.4 mg/h. Asterisk indicates  $P < 0.05$ .

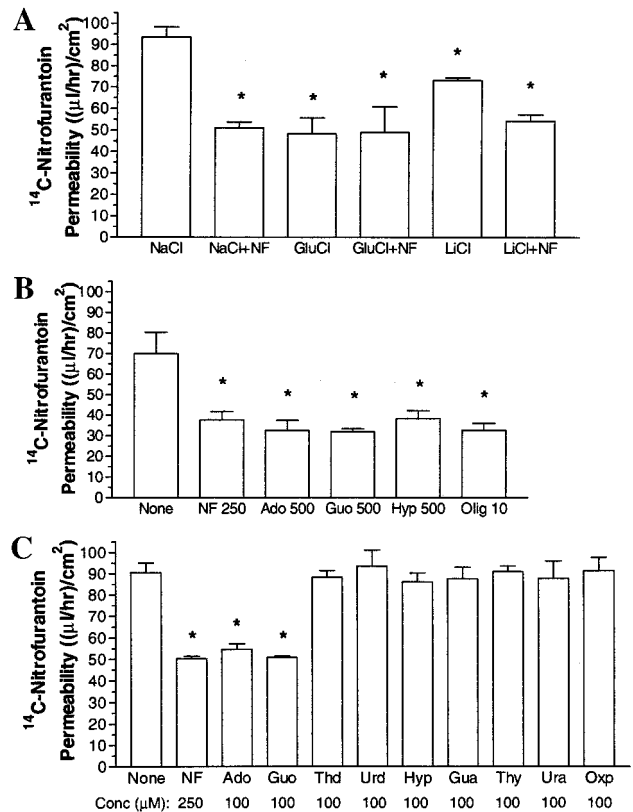


**Fig. 3.** Saturation and inhibition of <sup>14</sup>C-nitrofurantoin B-A flux. **(A)** Effect of cold nitrofurantoin. Regression line was fitted to equation 2. **(B)** Effect of dipyridamole on <sup>14</sup>C-nitrofurantoin B-A permeability. Regression line fitted to equation 4.

CIT3 cells grown on Snapwell filters. A portion of the <sup>14</sup>C-nitrofurantoin flux was inhibitable, and therefore apparently saturable, with  $K_m = 33.5 \mu\text{M}$  (95% CI 15.4 to 72.8  $\mu\text{M}$ ) as shown in Fig. 3. The observed total flux of <sup>14</sup>C-nitrofurantoin was 1.87 pmol/min (95% CI 1.75 to 1.99 pmol/min) and the non-saturable flux was 1.23 pmol/min (95% CI 1.11 to 1.34 pmol/min) and the saturable component was approximately 36%. Dipyridamole effectively inhibited <sup>14</sup>C-nitrofurantoin flux across CIT3 cells, with a  $K_i$  of 0.78  $\mu\text{M}$  (95% C.I. = 0.11 to 5.3  $\mu\text{M}$ ) as shown in Fig. 3b.

The effect on nitrofurantoin transport of complete replacement of sodium with lithium or glucosamine is shown in Fig. 4a and compared to the effect of a saturating concentration of cold nitrofurantoin (250  $\mu\text{M}$ ). These experiments were carried out in a modified Ringer's solution. Cold nitrofurantoin (250  $\mu\text{M}$ ) decreased the effective permeability of <sup>14</sup>C-nitrofurantoin from  $93.6 \pm 4.8 (\mu\text{l/h})/\text{cm}^2$  to  $51.0 \pm 2.7 (\mu\text{l/h})/\text{cm}^2$ . Replacing sodium with glucosamine HCl decreased nitrofurantoin permeability to a similar value of  $48.3 \pm 7.3 (\mu\text{l/h})/\text{cm}^2$ . LiCl decreased the permeability to  $73.3 \pm 1.4 (\mu\text{l/h})/\text{cm}^2$ ; additional inhibition was seen when cold nitrofurantoin was also added ( $54.3 \pm 3.1 (\mu\text{l/h})/\text{cm}^2$ ), suggesting that lithium can partly substitute for sodium.

In the next experiment, we examined the effects of adenosine, guanosine, hypoxanthine and oligomycin in DMEM/F12 medium are compared with 250  $\mu\text{M}$  cold nitrofurantoin (Fig. 4b). In the absence of any inhibitor, the effective permeability of <sup>14</sup>C-nitrofurantoin was  $70.0 \pm 10.4 (\mu\text{l/h})/\text{cm}^2$  but was decreased equivalently by 250  $\mu\text{M}$  cold nitrofurantoin (to  $37.6 \pm 4.1 (\mu\text{l/h})/\text{cm}^2$ ), adenosine (to  $32.4 \pm 4.9 (\mu\text{l/h})/\text{cm}^2$ ), guanosine (to  $31.8 \pm 1.4 (\mu\text{l/h})/\text{cm}^2$ ), hypoxanthine (to  $38.2 \pm$



**Fig. 4.** Sodium dependence of <sup>14</sup>C-nitrofurantoin (1  $\mu\text{M}$ ) transport and effects of nucleosides and nucleobases. **(A)** Effects of sodium replacement on <sup>14</sup>C-nitrofurantoin B-A permeability. NaCl: sodium chloride Ringer's solution; +NF: 250  $\mu\text{M}$  cold nitrofurantoin added to the basolateral side; LiCl: lithium chloride Ringer's solution; GluCl: glucosamine HCl Ringer's solution. Mean  $\pm$  SD of 3–5 determinations. Asterisks indicate  $P < 0.05$  vs. NaCl. **(B)** Effects of nitrofurantoin 250  $\mu\text{M}$  (NF), adenosine 500  $\mu\text{M}$  (Ado), guanosine 500  $\mu\text{M}$  (Guo), hypoxanthine 500  $\mu\text{M}$  (Hyp), and oligomycin 10  $\mu\text{M}$  (olig) on <sup>14</sup>C-nitrofurantoin B-A permeability. Numbers indicate concentrations ( $\mu\text{M}$ ) added to both sides, except NF added to basolateral side only. Mean  $\pm$  SD of 4–8 determinations. Asterisks indicate  $P < 0.05$  vs. control (none). **(C)** Effects of nucleosides and nucleobases on <sup>14</sup>C-nitrofurantoin B-A permeability. NF: 250  $\mu\text{M}$  cold nitrofurantoin added to both sides; 100  $\mu\text{M}$  of nucleosides or nucleobases on both sides indicated by the IUPAC-IUB abbreviations (Fig. 1); Oxp: 100  $\mu\text{M}$  oxypurinol. Mean  $\pm$  SD of 3–6 determinations. Asterisks indicate  $P < 0.05$  vs. control (none).

$3.9 (\mu\text{l/h})/\text{cm}^2$ ), and oligomycin (to  $32.5 \pm 3.5 (\mu\text{l/h})/\text{cm}^2$ ). A panel of nucleosides (adenosine, guanosine, thymidine, and uridine) and nucleobases (guanine, hypoxanthine, thymine, uracil, and oxypurinol) were studied in the saline (phosphate-free) buffer, shown in Fig. 4c. The control ( $90.5 \pm 4.6 (\mu\text{l/h})/\text{cm}^2$ ) decreased to  $50.3 \pm 1.1 (\mu\text{l/h})/\text{cm}^2$ ,  $54.5 \pm 2.6 (\mu\text{l/h})/\text{cm}^2$ , and  $50.7 \pm 0.6 (\mu\text{l/h})/\text{cm}^2$  in the presence of nitrofurantoin, adenosine, and guanosine respectively. The pyrimidine nucleosides uridine and thymidine and the nucleobases had no significant effects.

## DISCUSSION

Our studies of nitrofurantoin transport have shown an active transport component with characteristics similar but not identical to previously studied sodium dependent nucleo-

side or nucleobase transporters. In rats, we found a high M/S ratio for nitrofurantoin, which the broad-based inhibitor dipyridamole decreased by 62%. Kari *et al.* (4) reported a nitrofurantoin M/S approximately 75 times the predicted ratio for a passive process, implicating active transport into rat milk. In the rat, dipyridamole infusion (~10 mg/h) significantly inhibited incorporation of the nucleoside  $^3\text{H}$ -thymidine into the bone marrow and gastrointestinal epithelium (24). The infusion rate used in our study (12 mg/h) inhibited nitrofurantoin active transport into rat milk, but did not significantly inhibit the clearance of nitrofurantoin. The involvement of a nucleoside or nucleobase transporter would be consistent with this observation in lactating rats, but other transporters can not be ruled out.

Nitrofurantoin active transport across the mammary epithelium both *in vivo* and *in vitro* was sensitive to dipyridamole ( $K_i = 0.78 \mu\text{M}$ ), consistent with mediation by one of several possible transporters. Dipyridamole acts as a competitive or non-competitive inhibitor of several types of transporters, and is widely used as an inhibitor of nucleoside and nucleobase transporters (8,10). Regarding the concentrative nucleoside transporters, N1–N4 are insensitive to  $10 \mu\text{M}$  dipyridamole, but N5 and N6 are strongly inhibited by  $10 \text{ nM}$  and  $20 \mu\text{M}$  dipyridamole, respectively (10,25,26), although the type of inhibition has not been determined for either one. Sundaram *et al.* reported that dipyridamole competitively inhibits the human equilibrative nucleoside transporter hENT1 with an  $\text{IC}_{50}$  of  $198 \pm 24 \text{ nM}$  and a Hill coefficient of  $1.25 \pm 0.12$ , suggesting that one molecule binds to the binding site (27). Furthermore, chimeras they constructed of hENT1 with rENT1 (insensitive to dipyridamole) demonstrate that the inhibition is linked to specific domains of the protein (27).

Dipyridamole was shown to inhibit several other types of transporters. It was reported to inhibit organic cation transporters, although the type of inhibition was not reported (19). Bendayan *et al.* demonstrated that  $50 \mu\text{M}$  dipyridamole inhibited uptake of the organic cation *n*-methylnicotinamide, and that dipyridamole had an  $\text{IC}_{50}$  of  $7.7 \mu\text{M}$  against cimetidine uptake by LLC-PK1 cells (19). Dipyridamole inhibited p-glycoprotein (28); furthermore, Ayesb *et al.* showed that dipyridamole competed with verapamil for binding to p-glycoprotein, with a Hill coefficient of 2 (20). Additionally, it exerted a mixed competitive/non-competitive inhibition of bicarbonate-chloride exchange, with an  $\text{IC}_{50}$  of  $4 \mu\text{M}$  (29). Plagemann and Woffendin showed that dipyridamole inhibited 2-deoxy-D-glucose transport without changing the transition temperature of the transporter (30). Although dipyridamole is quite lipophilic ( $\log D = 2.2$  at pH 7.4; from ACD/LogP DB v4.0 using ACD/I-Lab service), this finding suggests that the inhibition of glucose transport by dipyridamole is not mediated by an alteration in membrane fluidity (30). The consensus amongst these studies is that although dipyridamole is a broad-based inhibitor of various transporters, the inhibition is due to binding of the drug to the various transport proteins rather than to changes in the properties of the membrane.

Toddywalla *et al.* reported a  $K_i$  of  $53 \pm 41 \mu\text{M}$  for nitrofurantoin transport across CIT3 cells grown on Snapwell filters, while performing the experiments in DMEM/F12 (23). In the present study we determined a  $K_m$  of similar magnitude,  $33.5 \mu\text{M}$  (95% CI  $15.4$  to  $72.8 \mu\text{M}$ ), for nitrofurantoin transport in saline buffer. Toddywalla *et al.* also reported a

B-A nitrofurantoin permeability of  $64 \pm 5 \times 10^{-3} \text{ cm/h}$  (or  $(\mu\text{l/h})/\text{cm}^2$ ) of which approximately 40% was inhibited by high nitrofurantoin concentrations (23). Similarly, in this study we observed a nitrofurantoin B-A permeability of  $70 (\mu\text{l/h})/\text{cm}^2$  in DMEM/F12, 47% of which was inhibitable. For experiments in Ringer's solution or the saline buffer, the B-A permeability ranged from  $74.5$  to  $97.4 (\mu\text{l/h})/\text{cm}^2$ , 35.7% to 51.5% of which was inhibitable. Because DMEM/F12 contains sodium and many potentially interacting substances (e.g., hypoxanthine, thymidine, phenol red, amino acids, and vitamins), other buffers were required for characterization studies. The Ringer's solution was chosen initially to conveniently replace sodium. However, the salt concentrations are different from the concentrations in the media in which the cells are cultured. The saline buffer (described above in Methods) was designed to provide similar concentrations of the major salt components, but without phosphates.

Sodium dependence was demonstrated in this study. Also, inhibition of active transport by the ATP synthase inhibitor oligomycin is consistent with an energy dependent process. Furthermore, lithium substitution only partially decreased nitrofurantoin B-A permeability, suggesting lithium may have substituted for sodium to some extent. By contrast, the nucleoside transporters generally show very high sodium specificity (10).

As mentioned previously, if the functionally classified N1–N6 sodium dependent nucleoside transporters were involved, then both adenosine and uridine should have inhibited nitrofurantoin transport (8–11). Because the  $K_m$ 's for these agents range from  $1$ – $45 \mu\text{M}$  (8,10), the  $100 \mu\text{M}$  concentrations used in this study would inhibit the nucleoside transporters by  $>50\%$ . However, this study showed that adenosine but not uridine inhibited nitrofurantoin transport. These results are not consistent with the known nucleoside transporters. Furthermore, guanosine but not thymidine inhibited transport, suggesting a preference for purine nucleosides.

If a known sodium dependent nucleobase transporter were involved, then guanine and/or uracil should inhibit nitrofurantoin transport (12–15,21). The  $K_i$ 's for guanine and uracil inhibition of hypoxanthine transport range  $0.6$ – $6.4 \mu\text{M}$  and  $2.9$ – $64.6 \mu\text{M}$ , respectively (12–15,17). Furthermore, the  $K_m$  for hypoxanthine transport range is  $0.78$ – $31.1 \mu\text{M}$  (14,15). At the concentrations used in this study, a nucleobase transporter similar to those previously reported would be inhibited by approximately 50% or more. Although hypoxanthine was inhibitory at  $500 \mu\text{M}$ , neither hypoxanthine nor the other nucleobases were inhibitory at  $100 \mu\text{M}$ . This finding is not predicted from the previously characterized sodium dependent nucleobase transport systems.

## CONCLUSIONS

Dipyridamole effectively inhibited the active transport of nitrofurantoin across mammary epithelia both *in vitro* with a high affinity and *in vivo*. Nitrofurantoin transport across the CIT3 model was sodium dependent. The effects of nucleosides and nucleobases on nitrofurantoin transport are not consistent with other studies on functionally classified nucleoside and nucleobase transporters. These results suggest that nitrofurantoin does not interact with the previously characterized nucleoside or nucleobase transporters. Further *in vivo*, *in vitro*, and molecular studies are required to elucidate the specific transport system for nitrofurantoin in mammary

epithelia. Understanding these mechanisms will allow better M/S predictions to be made, aiding clinicians to better assess the inadvertent exposure of suckling infants to various agents.

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