

H⁺ Extrusion by an Apical Vacuolar-Type H⁺-ATPase in Rat Renal Proximal Tubules

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Summary. The activity of Na⁺/H⁺-exchange and H⁺-ATPase was measured in the absence of CO₂/HCO₃ by microfluorometry at the single cell level in rat proximal tubules (superficial S₁/S₂ segments) loaded with BCECF [2',7'-bis(carboxyethyl)5-6-carboxyfluorescein-acetoxymethyl ester]. Intracellular pH (pH_i) was lowered by a NH₄Cl-prepulse technique. In the absence of Na⁺ in the superfusion solutions, pH_i recovered from the acid load by a mechanism inhibited by 0.1 μM bafilomycin A₁, a specific inhibitor of a vacuolar-type H⁺-ATPase. Readdition of Na⁺ in the presence of bafilomycin A₁ produced an immediate recovery of pH_i by a mechanism sensitive to the addition of 10 μM EIPA (ethylisopropylamiloride), a specific inhibitor of Na⁺/H⁺ exchange. The transport rate of the H⁺-ATPase is about 40% of Na⁺/H⁺-exchange activity at a similar pH_i (0.218 ± 0.028 vs. 0.507 ± 0.056 pH unit/min). Pre-exposure of the tubules to 30 mM fructose, 0.5 mM iodoacetate and 1 mM KCN (to deplete intracellular ATP) prevented a pH_i recovery in Na⁺-free media; readdition of Na⁺ led to an immediate pH_i recovery. Tubules pre-exposed to Cl⁻-free media for 2 hr also reduced the rate of Na⁺-independent pH_i recovery. In free-flow electrophoretic separations of brush border membranes and basolateral membranes, a bafilomycin A₁-sensitive ATPase activity was found to be associated with the brush border membrane fraction; half maximal inhibition is at 6 × 10⁻¹⁰ M bafilomycin A₁. It is concluded that in superficial rat proximal tubules a H⁺-ATPase and Na⁺/H⁺ exchange are colocalized in the apical membrane; H⁺ transport through the H⁺-ATPase is affected by intracellular ATP and Cl⁻ content. The transport capacity of apical H⁺-ATPase is sufficient to explain Na⁺-independent proximal tubular bicarbonate reabsorption.

Key Words bicarbonate reabsorption · Na⁺/H⁺-exchange · bafilomycin A₁ · ethylisopropylamiloride (EIPA)

Introduction

Secretion of H⁺ is the initial event leading to proximal tubular HCO₃⁻-reabsorption (for review *see* [1]). H⁺ secretion is to a large extent mediated by brush

border membrane Na⁺/H⁺ exchange. However, in the absence of Na⁺/H⁺-exchange activity, either in the absence of luminal Na⁺ and/or in the presence of inhibitors of Na⁺/H⁺ exchange, luminal acid secretion (bicarbonate reabsorption) continues, albeit at slower rates (for review *see* Ref. 1; e.g. *see* [7, 17]).

At a cellular level, this luminal acidification has been visualized as an alkalinization of intracellular pH (pH_i) which is only partially inhibited by Na⁺ removal or addition of amiloride (for review *see* [1]; e.g. [14, 17]). At a subcellular level, studies on isolated brush border membrane have provided evidence for two proton extrusion systems: Na⁺/H⁺ exchange and H⁺-ATPase (for review *see* [1, 6]; for H⁺-ATPase *see* [13, 20, 23, 26]). In studies on brush border membrane N-ethylmaleimide-sensitive H⁺-ATPase, it has been difficult to correlate the measured ATP-hydrolysis with functional H⁺ extrusion, mainly because of the right-side-out orientation of this membrane preparation and inaccessibility of ATP to the intravesicular space. A further complication in the interpretation of these vesicle studies is a possible contamination of brush border membranes by other cellular organelles containing this enzyme-/transport-activity [12, 13, 18–20, 23, 26] (for review *see* [9]). However, recent studies with monoclonal antibodies raised against bovine kidney medulla H⁺-ATPase have also shown immunolabeling of brush border membranes in rat proximal tubules; obviously, these experiments could not document proton-pump activity [5, 10]. Some authors have suggested that the H⁺-ATPase is inactive while inserted in the brush border membrane [6].

The goal of this paper was to provide some functional evidence for ATP-dependent H⁺-extrusion (H⁺-ATPase) in the plasma membrane of intact proximal tubular epithelial cells. By the analysis of alterations in cytosolic pH (pH_i) we could characterize some of the functional properties of this H⁺-ATPase and could also obtain an estimate of its pump activity relative to Na⁺/H⁺-exchange activity. Finally, membrane fractionation experiments

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permitted us to attribute H⁺-ATPase activity to the brush border membrane fraction.

Materials and Methods

PREPARATION OF PROXIMAL TUBULES AND BCECF/AM LOADING

A crude preparation of proximal tubules was obtained from 100-g male Wistar rats by cutting a 0.5–1.0 mm superficial slice and chopping it into small pieces (less than 1 mm) with a razor blade; tissue pieces were kept at 4°C in minimum essential medium (MEM; GIBCO), supplemented with 20 mM glucose and 1 mM adenosine but without phenol red. Prior to experiments on intracellular pH, tissue was incubated at 4°C for 30 min in this buffer containing 10 μM BCECF/AM [2'7'-bis(carboxyethyl)5-6-carboxyfluoresceinacetoxymethyl ester]. Tissue specimens were then immobilized by absorption on glass coverslips; these coverslips had been precoated with collagen by adding a solution of 2 mg/ml collagen in acetic acid and drying in air. Coverslips were then placed into a microscope chamber [15, 16] and superfused with the above medium containing 10 μM BCECF/AM. The chamber was subsequently centrifuged at 300 × g for 1 min at 4°C. This procedure immobilized small tubule fragments onto the collagen-coated coverslip so that the chamber could be perfused without tubule movement. Prior to perfusion with different experimental solutions, loading with BCECF/AM was completed by further incubation for 30 min at room temperature without perfusion. S₁ and S₂ segments of proximal tubules were identified by attachment of glomerular structures and/or the physical appearance of the large cells in convoluted tubules. Cells from S₁ and S₂ segments were studied with no quantitative differences in results.

FLUORESCENCE MEASUREMENTS

On the stage of an inverted microscope (Zeiss IM 35) proximal tubules were visualized so that the measuring area of a photomultiplier tube was focused on a single cell. The cells selected for study were either in single isolated tubules or were in tubules protruding from a group of several tubules. The microfluorometer was adjusted to monitor BCECF by excitation ratioing 390–440/475–490 nm and collecting emitted light between 515 and 565 nm as described previously [15, 16]; background correction was performed by subtracting the autofluorescence of comparable tubule cells without dye. Solutions used throughout the pH_i measurement/perfusion experiments were nominally CO₂/HCO₃⁻ free. The standard NaCl solution contained (in mM): 130 NaCl, 4 KCl, 1 MgSO₄, 1.7 CaCl₂, 1.0 NaH₂PO₄, 20 glucose, 20 HEPES adjusted with Tris to a pH of 7.4. Tetramethylammonium-chloride (TMACl) solution contained TMA⁺ instead of Na⁺, and NH₄Cl solution contained 20 mM NH₄⁺ and 110 mM TMA instead of Na⁺. In experiments without chloride, all chloride salts (Na⁺, K⁺, Ca²⁺, TMA⁺, NH₄⁺) were replaced by gluconate salts. Tetramethylammonium gluconate and ammonium gluconate were prepared by titration of corresponding hydroxides with D-gluconic acid lactones.

ISOLATION OF PLASMA MEMBRANE VESICLES BY FREE-FLOW ELECTROPHORESIS

Crude plasma membrane isolation and free-flow electrophoresis was performed essentially as described by Heidrich et al. [11] with a slightly modified chamber buffer (in mM: 100 mannitol, 10

triethanolamine, 10 acetic acid adjusted with NaOH to a pH of 7.4) at 180 V/cm and 95 mA at 4°C using a Bender Hobein VAP22 apparatus. After separation, 90 fractions were collected and in each fraction Na⁺-K⁺-ATPase and leucine aminopeptidase (LAP) activity were assayed to measure the distribution of basolateral (Na⁺-K⁺-ATPase) and brush border membranes (LAP), respectively [2, 11].

MEASUREMENTS OF BAFILOMYCIN A₁ (BA₁)-SENSITIVE ATPASE

BA₁-sensitive ATPase was measured by a P_i-liberation assay similar to that described by Dean et al. [8]. Briefly, vesicles (25 μg protein) were incubated for 15 min at 37°C in 200 μl of the following incubation medium (in mM): 250 sucrose, 100 KCl, 5 MgSO₄, 1 ouabain, 1 levamisole, 5 disodium ATP, 10 HEPES adjusted with Tris to a pH of 7.0; oligomycin was added at a concentration of 5 μg/ml and incubation was performed in the presence or absence of bafilomycin A₁ (usually 0.1 μM). After this incubation, 1 ml was added of a solution containing 2.9% ascorbic acid (wt/vol), 0.48% ammonium molybdate (wt/vol), 2.86% sodium dodecyl sulfate (wt/vol) in 0.48 N HCl, and samples were left on ice for 10 min. This was followed by an incubation at 37°C for 15 min, after addition of 1.5 ml of a solution containing 2% Na-arsenite (wt/vol), 2% Na citrate (wt/vol), 2% (wt/vol) acetic acid in H₂O. Absorbance at 705 nm was used to measure P_i release. The bafilomycin A₁-sensitive ATPase activity was expressed as the difference in ATPase activity with and without inhibitor. Prior to ATPase measurements, vesicles were permeabilized with 0.5% (wt/vol) octyl-beta-D-glucopyranoside (octylglucoside treatment) at 30°C for 15 min.

CHEMICALS

Unless indicated, all chemicals were obtained from Fluka (Buchs, Switzerland), Merck (Darmstadt, Germany), or Sigma (St. Louis, MO) and were of the finest grade available. Ethylisopropyl amiloride (EIPA) was the gift from Dr. Thomas Friedrich (Frankfurt/M, Germany), and bafilomycin A₁ was the gift from Prof. Dr. Karlheinz Altendorf (Osnabrück, Germany). H₂DIDS and BCECF/AM were from Molecular Probes (Eugene, OR).

Results

The activity of plasma membrane H⁺-extrusion mechanisms of isolated rat proximal tubular epithelial cells were measured by single cell microfluorometry of intracellular pH (pH_i), using cells loaded with BCECF [1, 14, 15, 16]. Changes in pH_i were always recorded in bicarbonate-free media and were monitored exclusively from cells located at the end of proximal tubular fragments, in order to ensure that both luminal and contraluminal sides were exposed to perfusion solutions. For the present experiments it was crucial to minimize the time of isolation and to eliminate the exposure of tissues to enzymes frequently used in preparing isolated tubular segments, such as collagenase and hyaluronidase. If these enzymes were used to prepare a tubular sus-

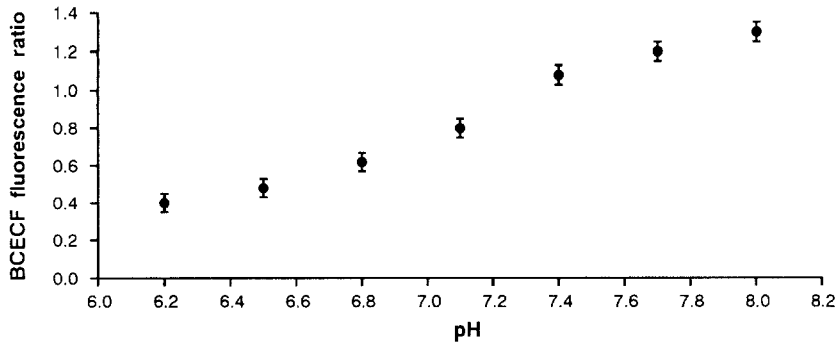


Fig. 1. Calibration of the BCECF fluorescence signal from the proximal tubules. Tubules were perfused with high K⁺ (110 mM) solution with addition of 5 μ M nigericin. Different media pHs were tested and the fluorescence ratio readings, from alternative excitation of BCECF at 390–440 nm and 475–490 nm, were plotted against extracellular pH (mean \pm SE; $n = 3$).

pension, Na⁺-independent H⁺-extrusion activity was either diminished or completely absent (*data not shown*).

To validate the use of BCECF in this preparation, the potassium/nigericin calibration method was used as originally described by Thomas et al. [25] to correlate the fluorescence ratios to pH_i [15, 16]. The curve that was determined (shown in Fig. 1) was similar to the calibration curves obtained previously in the same microfluorometer, using established renal cell lines grown on permeant supports [15, 16]. The mean steady-state pH_i of proximal tubular cells perfused with NaCl solution was 7.45 ± 0.04 ($n = 25$).

In most of the following experiments, H⁺-extrusion mechanisms were studied after cellular acidification created by the classical NH₄⁺-prepulse technique [3, 15, 16]. After perfusion with the standard NaCl solution (*see Materials and Methods*) the tissue was perfused with a Na⁺-free TMA medium containing 20 mM NH₄Cl (replacing isoosmotically TMA-Cl). The presence of NH₄Cl caused a cellular acidification (Fig. 2a) in contrast to most other previously examined cell types (e.g. in squid giant axons, OK cells and LLC-PK₁ cells) where addition of NH₄Cl produces transient alkalization (e.g. [3, 15, 16]). Since the acidification was not inhibited by 10 μ M EIPA (*data not shown*), it is unlikely to be due to reversal of Na⁺/H⁺ exchange. The data suggest the presence of a significant NH₄⁺-influx pathway, although it was not possible to identify the mechanism of NH₄⁺-ion uptake. The rapid acidification was not prevented by 1 mM furosemide (inhibitor of Na⁺/K⁺/2Cl⁻ symport), 10 mM tetraethylammonium, 10 mM BaCl₂ or 2 mM chloroquine (blockers of various K⁺ channels) (*data not shown*). After pH_i reached an apparent steady state, NH₄Cl was removed and replaced by the TMA-Cl solution (Fig. 2b). The resulting further acidification was followed by a strong alkalization and an overshoot in pH_i over the resting pH_i value (Fig. 2c). This recovery from an acid load was observed in the complete absence of Na⁺,

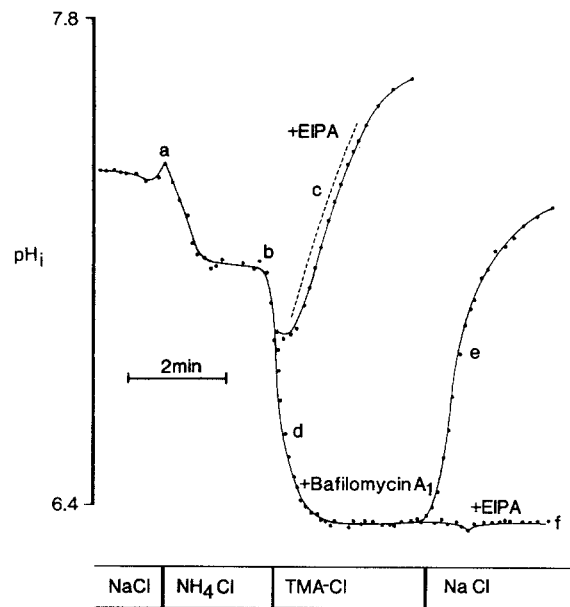


Fig. 2. Presence of bafilomycin A₁ (BA₁)-sensitive, Na⁺-independent pH_i recovery in rat proximal tubule cells. After perfusion with a NaCl solution, a NH₄Cl solution (*see Materials and Methods*) provoked an acidification (a). After removing NH₄Cl solution further rapid acidification occurred (b), followed by a pH_i recovery in a Na⁺-free medium (TMA-Cl) (c). This recovery (0.218 ± 0.028 pH_i unit/min) (c) was not inhibitable by 10 μ M EIPA (c, dotted line) or 100 μ M H₂DIDS (*not shown*). In the presence of 0.1 μ M BA₁ (d), acidification due to NH₄Cl removal was stronger and pH_i did not recover. Tubules treated with BA₁, after perfusion with NaCl medium, exhibited Na⁺/H⁺-exchange activity (0.507 ± 0.056 pH_i unit/min (e), blockable by 10 μ M EIPA (f).

was not dependent on extracellular chloride (*not shown*) and was not sensitive to 10 μ M EIPA (Fig. 2c; dotted line) or 100 μ M H₂DIDS (blocker of several anion exchange mechanisms; *data not shown*).

Bafilomycin A₁ (BA₁) has recently been described as a potent inhibitor of vacuolar (V) ATPases [4]. Addition of 0.1 μ M BA₁ completely blocked the Na⁺-independent recovery in pH_i and caused larger acidification after NH₄Cl removal (Fig. 2d). The in-

hibition of Na⁺-independent recovery by 0.1 μM BA₁ strongly suggests that Na⁺-independent pH_i recovery is related to a (V) ATPase. In the continuous presence of BA₁, subsequent addition of Na⁺ provoked an immediate return of pH_i to resting values (Fig. 2e). This Na⁺-dependent recovery in pH_i was prevented by 10 μM EIPA, documenting that it represents Na⁺/H⁺ exchange (Fig. 2f). Therefore, these experiments show that two separate mechanisms mediate proton extrusion from the cytosol of rat proximal tubular epithelial cells: a vacuolar type H⁺-ATPase and Na⁺/H⁺ exchange. If the pH_i recovery rates are calculated over the same ranges of pH_i (6.7 to 7.0), thus with similar intracellular buffering capacities, it is possible to compare rates of Na⁺/H⁺ exchange (0.507 ± 0.056 pH unit/min; n = 7) to those of H⁺-ATPase (0.218 ± 0.028 pH unit/min; n = 9). Thus, over a pH_i range close to physiological values, H⁺ extrusion by H⁺-ATPase is about 40% that by Na⁺/H⁺ exchange.

To show that the Na⁺-independent pH_i recovery rate was dependent on cellular ATP content, we included protocols known to lower ATP content in most cells. The combined application of 1 mM KCN (inhibitor of oxidative phosphorylation), 0.5 mM iodoacetate (IAA; inhibitor of glycolysis) and 30 mM fructose (hexokinase substrate prevented Na⁺-independent pH_i recovery (Fig. 3d), but not Na⁺-dependent recovery (Na⁺/H⁺ exchange) (Fig. 3e). V-ATPases [9], as well as kidney and macrophage endosomal ATPase are Cl⁻ dependent [12, 19, 21, 23, 24]. In tubules exposed for 2 hr to Cl⁻-free media (replacement by gluconate), H⁺-pump activity was reduced (Fig. 3c). Control experiments were performed using tubules from the same preparation maintained in the standard NaCl solution. This control group maintained normal H⁺-extrusion activity for 2 hr (*data not shown*). This suggests that removal of Cl⁻, and not tissue degradation, is responsible for the decrease in H⁺ extrusion in the experimental group. Thus, Na⁺-independent pH_i recovery seems to be related to a Cl⁻-dependent, ATP-dependent mechanism.

Measuring Na⁺-independent H⁺ extrusion resulted in alkalinization which was above normal steady state observed in NaCl medium (*see* Fig. 2). This alkalinization (pH = 7.93 ± 0.08; n = 15) could not be returned to the normal steady-state value unless NaCl medium was re-added; the rate of this recovery (alkali extrusion) was 0.251 ± 0.022 pH unit/min (n = 6). This phenomenon was not affected by replacements of Cl⁻ with gluconate (*data not shown*) and also was not inhibited by 10 μM EIPA (*data not shown*). However, as indicated in Fig. 4, this return to resting pH_i after readdition of Na⁺ was sensitive to 100 μM H₂-DIDS: readdition of NaCl

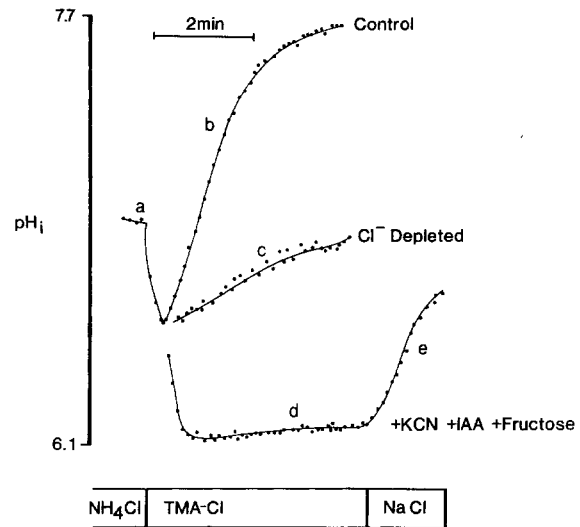


Fig. 3. Cl⁻ and ATP dependence of the Na⁺-independent pH_i recovery rate. NH₄Cl was withdrawn (a) to produce an acid load (*see also* Fig. 2) and a normal Na⁺-independent pH_i recovery rate was observed (b). Tubules pre-exposed to Na⁺-gluconate medium for 2 hr/37°C, in order to decrease the intracellular Cl⁻ content, gave much smaller pH_i recovery rate analyzed in the Cl⁻-free medium (c). In cells exposed to 30 mM fructose, 0.5 mM iodoacetate (IAA) and 1 mM KCN for 30 min at 4°C (to decrease the intracellular ATP content), Na⁺-independent pH_i-recovery rate was completely inhibited (d) (always in the presence of Cl⁻-containing media). The Na⁺/H⁺-exchange activity remained in the presence of KCN, IAA and fructose (e).

medium in the presence of 100 μM H₂DIDS did not lead to a fall in pH_i; removal of H₂-DIDS in the continuous presence of NaCl initiated an immediate pH_i-recovery mechanism which leveled off at the resting pH_i, observed in normal NaCl-perfusion conditions.

From the experiments described above, conclusions on the cellular location (basolateral *vs.* brush border) of the H⁺-extrusion systems cannot be given. To assess the plasma membrane location of the BA₁-inhibited H⁺-pump activity (*see* Fig. 2) we measured BA₁-sensitive ATPase activity in separated brush border (BBM) and basolateral membranes (BLM). We used a free-flow electrophoretic procedure for membrane separation; applying this method of isolation, contamination of BBM and BLM by endocytic vesicles, lysosomes, mitochondria and endoplasmic reticulum is minimal [11]. Figure 5A shows the separation of BBM (leucine-aminopeptidase activity, LAP) and BLM (Na⁺-K⁺-ATPase activity). In crude plasma membranes (PM), prior to application to the free-flow electrophoresis procedure, LAP was enriched by a factor of 4.7 ± 1.0 (n = 3) over activity in the homogenate and Na⁺-K⁺-ATPase enrichment was 7.0 ± 1.0 fold (n =

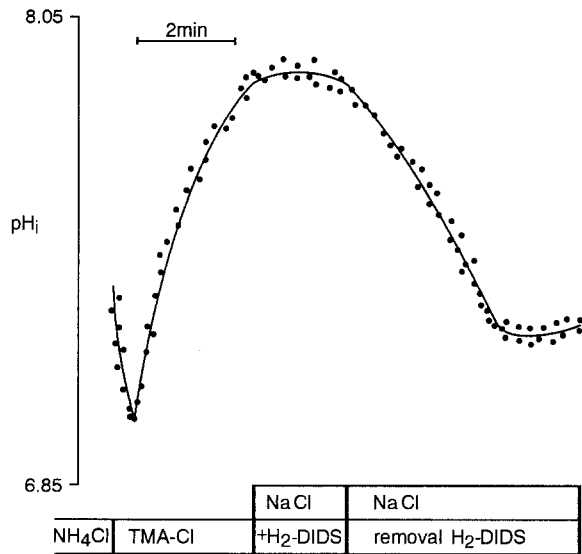


Fig. 4. Na⁺-dependent, H₂DIDS-inhibitable alkali extrusion. In CO₂/HCO₃⁻-free media, Na⁺-independent H⁺ extrusion produces intracellular alkalinization (mean pHi = 7.93 ± 0.08) which cannot be reversed in the absence of Na⁺ (not shown). By addition of Na⁺ to the extracellular fluid the pHi returns to resting pHi values observed in normal Na⁺-perfusion conditions. This Na⁺-dependent alkali extrusion was reversibly inhibited by 100 μM H₂DIDS; as indicated in the figure, the addition of Na⁺ in the presence of H₂DIDS does not lead to a recovery of pHi, removal of H₂DIDS is followed by alkali extrusion. The rate of this recovery was 0.251 ± 0.022 pH unit/min.

3). In pool 1, the BBM fraction, LAP activity was enriched 12.5 ± 3.2 fold (*n* = 3), whereas Na⁺-K⁺-ATPase activity was enriched 2.7 ± 0.1 fold of that in the homogenate. In pool 2, the BLM fraction, Na⁺-K⁺-ATPase activity was enriched 21.0 ± 5.1 (*n* = 3) and LAP activity 3.2 ± 0.4 (*n* = 3) fold. In Fig. 5, not all of the BA₁-sensitive ATPase activity from the crude plasma membrane fraction (PM) is copurified with LAP. This is likely to be due to either the presence of BA₁-sensitive ATPase in intracellular organelles which are eliminated during free-flow electrophoresis, or to degradation of the ATPase during electrophoresis. The vesicles obtained by the free-flow electrophoresis were permeabilized with 0.5% (wt/vol) octylglucoside in order to expose both sides of the membrane to the ATPase assay procedure. Figure 5B shows that there is a significant BA₁-sensitive ATPase activity, measured by a phosphate liberation assay, in crude plasma membranes (PM). Figure 5B also demonstrates that BA₁-sensitive ATPase activity is concentrated in the BBM fractions and almost absent in the BLM fractions. Together with experiments given in Fig. 2, the observation presented in Fig. 5 is strong evidence for the BA₁-sensitive H⁺ extrusion being associated with

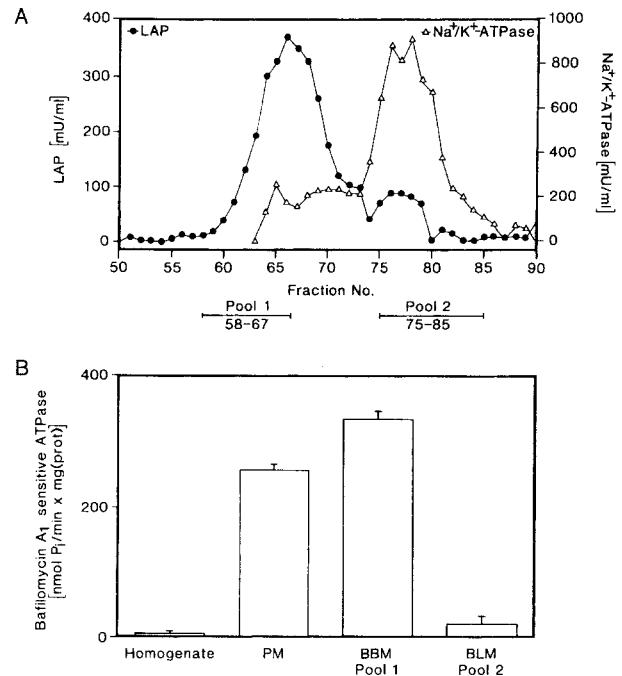


Fig. 5. Localization of the vacuolar type (V) H⁺-ATPase in brush border membranes. Crude plasma membrane vesicles were isolated and separated by free-flow electrophoresis (A) After separation, 90 fractions were collected and in each fraction Na⁺/K⁺-ATPase and leucine aminopeptidase (LAP) activity were assayed as the measure for the distribution of the basolateral (Na⁺-K⁺-ATPase) and the brush border membranes (LAP), respectively. (B) Fractions from free-flow electrophoresis were pooled as indicated in A and enzyme activities (H⁺-ATPase) measured in homogenates of kidney cortex slices and plasma membranes (PM) applied to free-flow electrophoresis and in basolateral (BLM) and brush-border membranes (BBM).

the BA₁-sensitive H⁺-ATPase activity located in the brush border membrane.

Since, at higher concentrations, BA₁ is known to inhibit other ATPases in addition to the V-type [4], we tested the sensitivity of the oligomycin-, ouabain- and levamisole-resistant ATPase activity to increasing concentrations of BA₁ (Fig. 6). 50% inhibition was observed at a BA₁ concentration of around 0.6 nM, a value far below that reported for inhibition of F₁F₀-ATPases (50% inhibition at μM concentration of BA₁; [4]). In comparison to N-ethylmaleimide (IC₅₀ in μM range) BA₁ is at least 10⁴ times more potent as an inhibitor of vacuolar H⁺-ATPases.

Discussion

In the present experiments it was important to minimize any lengthy manipulation of the tissue and to exclude enzymatic treatment, which might damage

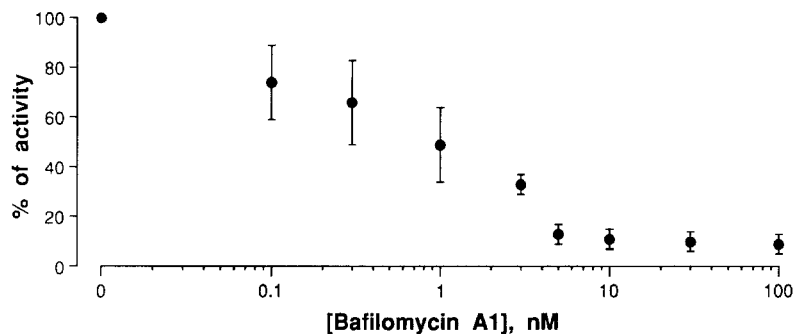


Fig. 6. Bafilomycin A₁ inhibition curve for oligomycin-, levamisole-, ouabain-resistant ATPase in brush border membranes (BBM). BBM obtained by the free-flow electrophoresis separation (*see* Fig. 5, *pool 1*) were treated with 0.5% (wt/vol) octylglucoside and exposed to various concentrations of bafilomycin A₁. ATPase assay was performed as described in Materials and Methods. Total oligomycin-, laevamisole-, ouabain-resistant ATPase activity in these experiments was 219 ± 50 nmol P_i/min · mg protein ($n = 3$). Bafilomycin A₁-insensitive ATPase (also oligomycin-, laevamisole-, ouabain-resistant) was 22.3 ± 11.7 nmol P_i/min · mg protein ($n = 3$). The values represent percent inhibition of the former activity in the presence of different concentrations of bafilomycin A₁. I₅₀ for bafilomycin A₁ was 0.61 ± 0.15 nM ($n = 3$).

some plasma membrane proteins. Only under such conditions do proximal tubules have the ability to extrude protons after acid load in the absence of Na⁺. The fact that the observed Na⁺-independent H⁺-extrusion activity (Fig. 2) and H⁺-ATPase in purified brush border membranes (Fig. 5) are inhibited by low concentration of bafilomycin A₁ strongly suggests the presence of a functional vacuolar H⁺-ATPase pump in the BBM. Indeed, studies with monoclonal antibodies have demonstrated the presence of the vacuolar proton pump in the proximal tubular brush border membrane [5]. Furthermore, Na⁺-independent H⁺ extrusion (bicarbonate reabsorption) has been shown earlier (for review *see* [1]; e.g. [7, 14, 17, 22, 28], although the mechanisms involved in this phenomenon have been difficult to define.

In the absence of Na⁺, H⁺ extrusion overshoots the steady-state pH_i, and addition of Na⁺ returns pH_i to the steady-state pH_i (observed under Na⁺-perfusion conditions), via H₂DIDS-sensitive, EIPA-insensitive, Cl⁻-independent mechanism (Fig. 4 and *data not shown*). Although the precise nature of this latter transport mechanism was not investigated in the present study, the possibility exists that it represents Na[OH⁻]HCO₃⁻ cotransport. This hypothesis presumes that sufficient cellular Na and base equivalents are available for net efflux. Such a mechanism could be operative in nominally bicarbonate-free solutions due to cellular CO₂ production and carbonic-anhydrase activity or due to efflux of hydroxyl ions. This DIDS-sensitive transport mechanism seems to have the potency to sense pH_i and to reduce its transport rate near resting pH_i. An alternate inter-

pretation is that at resting pH_i the rate of this alkali extrusion mechanism could match the rate of H⁺-extrusion mechanism. The detailed study of this phenomenon was not in the scope of the present paper but could be a basis for further investigations.

The data presented do not allow one to draw any conclusions concerning the activity of the Na⁺-independent H⁺-extrusion system at resting pH_i in the presence of full Na⁺. By the technique used (measurement of pH_i), inhibition of the pump by bafilomycin A₁ in the presence of Na⁺ cannot result in intracellular acidification as Na⁺/H⁺-exchange activity compensates and keeps pH_i at resting values. On the other hand, any pH_i alteration (acidification) or also a constant pH_i after amiloride inhibition of Na⁺/H⁺ exchange might be the result of (only partial) inhibition of Na⁺/H⁺ exchange which can (in part) be compensated by the H⁺ pump.

It is known that in acidifying epithelia H⁺-ATPase might be recruited from an intracellular compartment to the apical surface, e.g. due to acidification of intracellular space [14, 22, 27]. Two lines of evidence argue against the involvement of such a mechanism in the present study: (i) the presence of colchicine (100 μM after 30 min of preincubation) had no influence in the present study on the rate of Na⁺-independent bafilomycin-sensitive H⁺ extrusion (0.161 ± 0.051 pH unit/min in the absence of colchicine and 0.211 ± 0.054 in the presence of colchicine; $n = 3$; pH range for measuring of Na-independent pH_i recovery was between 6.7 and 7.0). This makes an exocytic insertion rather unlikely, in contrast to a previous study on rabbit S₃-proximal tubule seg-

ments [14]. (ii) Bafilomycin A₁-sensitive ATPase was also present in brush border membrane vesicles isolated from rat renal superficial cortex from the same animals not pre-exposed to an acidotic condition.

Finally, we believe that the activity measured in isolated brush border vesicles and the activity measured in H⁺ extrusion by microfluorometry is related to an identical molecular structure. The following arguments are in favor of such an interpretation: (i) H⁺-ATPase as well as H⁺ extrusion are inhibited by low bafilomycin A₁ concentrations; (ii) H⁺ extrusion is reduced by lowering intracellular [Cl⁻], and vacuolar ATPase enzyme activity in intracellular organelles is known to require Cl⁻ for optimum activity [19]; (iii) other investigators have documented a brush border location of vacuolar type H⁺-ATPase by immunohistochemical methods [5]; this enzyme activity is known to be sensitive to bafilomycin A₁ [4, 9].

Considering the physiological role of the proximal tubular brush border Na⁺-independent H⁺ pump, it is very likely that this mechanism plays a significant role in acid extrusion. It may also excrete protons at significant rates when intratubular pH is low, since Na⁺/H⁺ exchange is known to be inhibited by acidic pH in the extracellular medium. Thus, this H⁺ extrusion system might become more prominent with progressing/increased intratubular acidification. Furthermore, the Na⁺-independent H⁺-extrusion mechanism might also operate at more alkaline intracellular pH_i values than Na⁺/H⁺ exchange (*see* presented data). Thus, the relative contribution of Na⁺-independent H⁺ extrusion might also become more important with progressing intracellular alkalosis.

In conclusion, this study presents functional as well as biochemical data on brush border membrane location of a vacuolar type H⁺-ATPase in renal proximal tubules. These observations extend previous biochemical information on NEM-sensitive ATPase activity in isolated brush border membranes [13, 18–20, 23, 26], as well as on Na⁺-independent H⁺ extrusion [7, 14, 17, 22, 28] by attributing it to a bafilomycin A₁-sensitive, Cl⁻-dependent vacuolar type (V) ATPase. Furthermore, the present data suggest that the rate of this H⁺-extrusion activity in brush border membrane is sufficient to secrete the fractional amount of protons necessary for the Na⁺-independent mode of HCO₃⁻-reabsorption in the proximal tubules (ca. 30–40%).

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