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Ultraviolet-Absorbing Organic Anions in Uremic Serum Separated by Capillary Zone Electrophoresis, and Quantification of Hippuric Acid

A. C. Schoots, T. P. E. M. Verheggen, P. M. J. M. De Vries, and F. M. Everaerts

Organic anions accumulated in blood serum of patients with chronic renal failure were separated by a novel technique: closed-system capillary zone electrophoresis (CZE) in a pH 6 carrier-electrolyte system. Hippuric acid (HA), \( \beta \)-hydroxyhippuric acid, and uric acid were identified by their co-elution with standards prepared in ultrafiltered normal serum and by comparison with the corresponding ultraviolet-detected peaks positively identified in the HPLC analyses. Analysis time for the entire profile is 8 min. Repeatabilities (CVs) of CZE migration times and peak areas of the three acids in serum samples were about 0.7% and 6%, respectively. We quantified HA in 10 ultrafiltered uremic serum samples and compared results with those by a previously described HPLC procedure. The very good agreement further supports the identification of hippuric acid. Accuracy and precision of the CZE method were similar to those for the HPLC gradient-elution method, but analysis time for HA (8 min) is much less than by HPLC (90 min). Our technique is very suitable for selective, rapid analysis for (ultraviolet-absorbing) anionic constituents in ultrafiltered uremic serum, without any sample pretreatment.

Additional Keyphrases: renal function · tubular secretion · anion gap · uremic toxins · HPLC compared · solvent effect

Substances that accumulate in the body fluids of patients with advanced impairment of renal function include aromatic or indolic compounds (e.g., phenols, phenolic acids, hippurates, indolic acids, amines), heterocyclic nitrogen-containing compounds (e.g., guanidines, creatinine, uric acid), trace elements (e.g., aluminum, nickel), low-molecular-mass proteins (e.g., \( \beta \)-microglobulin, lysozyme, retinol-binding protein), parathyrin, potassium, and so-called middle molecules (1–4).

Artificial methods for blood purification such as hemodialysis and continuous ambulatory peritoneal dialysis partly suppress or prevent the clinical symptoms observed in patients with advanced renal failure. Therefore, dialyzable toxins apparently must be directly or indirectly responsible for the multiple disorders observed in uremic patients. Until now, there has been no conclusive evidence for the toxicity of a single substance or a specific group of accumulating solutes. For the past 15 years, much attention has been paid to the "middle molecule" hypothesis (5), but this hypothesis is not convincingly supported by experimental evidence, partly because the "middle molecules" are ill-defined chemically (3, 6).

Various authors have pointed out the potential importance of organic anions that normally are excreted into urine by a highly efficient (proximal) tubular secretory mechanism in the kidney (7–9). These solutes inhibit cell-membrane ion transport in various tissues (8, 10). Recently, a multivariate statistical study showed that high concentrations of aromatic acids coincide with low nerve-conduction velocities in patients on maintenance dialysis (11). Hippuric acid (HA), the aromatic acid regularly present at the highest concentration in uremic serum, may account for part of the positive anion-gap (12) observed in uremic patients and has been shown to inhibit drug-protein binding in (dialyzed) renal patients (13, 14). Hippurates are glycone conjugates formed in the liver from benzoic acids. Benzoic acid originates from the diet (fruits and vegetables, food preservatives), from mitochondrial \( \beta \)-oxidation of phenyl-fatty acids with an odd number of carbons, and from oxidative breakdown of phenylalanine through bacterial action in the intestines (10, 15, 16). Furthermore, environmental contact with xylenes or toluenes results in increased HA concentrations in serum. Commercial heparin, frequently used in hemodialysis practice, may contain benzyl alcohol, the metabolism of which results in the generation of HA. Because HA may be a marker of accumulation of tubularly secreted organic acids (both endogenous and exogenous) and their glycone conjugates in the serum, it is important to develop reliable, rapid methods for detection and quantification of HA and other aromatic acids.

Organic acids in uremic serum can be analyzed, as a group, by use of temperature-programmed gas chromatography (17, 18) or "high-performance" liquid chromatography (HPLC) with gradient-elution (9, 19–21), both of which are relatively time-consuming procedures. Specific analysis for HA has also been done colorimetrically (22) and by HPLC (23), but these methods require sample pretreatment, such as extraction with organic solvents. Other methods used for selective isolation of urinary organic acids in general are anion-exchange and adsorption chromatography (24). Earlier we reported a preliminary study on the pre-separation of those anions present in uremic serum by isotachophoresis preceding HPLC (25). Here we describe a method for the rapid assay of anionic constituents in uremic serum by closed-system capillary zone electrophoresis (CZE), without sample pretreatment, and we compare results with those obtained with our HPLC method.

Materials and Methods

Serum and standard samples. Pre-dialysis blood, sampled from patients undergoing regular hemodialysis treatment, was allowed to clot before centrifugation, then stored at \(-70^\circ C\) until assay. Before analysis, the samples were deproteinized by centrifugal ultrafiltration (ambient temperature, 1500 \( \times \) g) through micropartition ultrafiltration units (Grace & Co., Amicon Div., Danvers, MA 01923). Thus, the HA concentrations measured represent the portion of HA not bound to serum proteins [38% of HA in

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uremic serum is protein-bound (21)]. We prepared standard solutions for calibration in CZE and HPLC by adding HA to ultrafiltered normal (nonuremic) reference serum to give final concentrations in the injection calibration solutions of 0, 12.5, 25, 50, and 125 μmol/L for CZE analysis and 0, 25, 50, 100, and 250 μmol/L for HPLC analysis. Before injection, the ultrafiltered normal serum in these solutions was diluted 10-fold for CZE, fivefold for HPLC. These same dilution factors were used with the ultrafiltered uremic sera.

**Capillary electrophoresis.** For zone-electrophoresis experiments we used a home-made apparatus, described in detail elsewhere (26). The sampling unit has a fixed-volume (0.6 μL) sampling compartment. For the separation compartment we used non-coated Teflon capillaries (200 μm i.d.). The electrode compartments were separated from the separation compartment by semipermeable membranes. We used a slightly modified Model UV-M (Pharmacia, Uppsala, Sweden) fixed-wavelength ultraviolet-absorbance detector, operated at 254 nm. During analysis, the electrical current was kept constant at 50 or 35 μA and the voltage at about 10 kV. An Alpha-Series power supply (Brandenburg, Thornton Heath, U.K.) was used. Two carrier electrolyte systems were used (Table 1).

**HPLC analyses** were performed as described earlier (9, 21) but with a different separation column. We used a 4.6 mm (i.d.) × 25 cm Ultrasound Octadecyl (C18-modified silica) column, packed with 5-μm particles, in conjunction with a 2 mm (i.d.) × 3 cm Ultrasound Octadecyl guard column, packed with 10-μm particles (both from Beckman Instruments Inc., Fullerton, CA). The 45-min solvent gradient was linear from 100% aqueous ammonium formate buffer (0.05 mol/L, pH 4) to 60% methanol/40% buffer. The flow rate was 1 mL/min. The separation column and the solvent were kept at 25 °C by means of a thermostated bath and column water jacket. The chromatograph consisted of a Model 421 controller, two Model 100A double-piston pumps, a Model 160 fixed-wavelength detector (wavelength 254 nm, range 0.02 A full scale), and a Model 500 autosampler (all from Beckman Instruments). The injection volume was 25 μL.

**Data acquisition and statistics.** Zone-electrophoresis elution profiles and HPLC chromatograms were acquired with a chromatography data system. We used a Model 761S data interface and Model 2600 chromatography software (both from Perkin-Elmer/Nelson, Cupertino, CA) and an IBM PC/XT computer. Sampling frequency was 10 Hz in CZE and 1 Hz in HPLC. For regression analysis we used the REG procedure from SAS statistical software (27).

**Materials.** 2-(N-Morpholino)ethanesulfonic acid and hip-

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**Table 1. Operational Systems Used in Capillary Zone Electrophoresis**

<table>
<thead>
<tr>
<th>System I</th>
<th>System II</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Carrier constituent</strong></td>
<td>Acetate</td>
</tr>
<tr>
<td><strong>Conc of carrier, mol/L</strong></td>
<td>0.01</td>
</tr>
<tr>
<td><strong>Counter constituent</strong></td>
<td>β-Alanine</td>
</tr>
<tr>
<td><strong>pH</strong></td>
<td>3.8</td>
</tr>
<tr>
<td><strong>Additive</strong></td>
<td>MHEC, 0.5 g/L</td>
</tr>
<tr>
<td><em><em>Capillary</em> length, cm</em>*</td>
<td>15</td>
</tr>
<tr>
<td><strong>Driving current, μA</strong></td>
<td>50</td>
</tr>
</tbody>
</table>

* Teflon, 0.2 mm (i.d.). MES, 2-(N-morpholino)ethanesulfonic acid. MHEC, methylhydroxyethylcellulose.
Tentative peak identities: 1, hippuric acid; 2, p-hydroxyhippuric acid; 3, uric acid. Detection was at 254 nm. Ultrafiltered serum was diluted 1:1 with water in A, 1:8 in B and C. "AU" is A, absorbance

Fig. 1. Capillary zone electropherograms of (A) uremic serum in pH 3.8 electrolyte (System I, Table 1), (B) uremic serum in pH 6 electrolyte (System II, Table 1), and (C) ultrafiltered normal serum in pH 6 carrier electrolyte (System II, Table 1)

Fig. 2. Corresponding CZE and HPLC profiles for uremic serum ultrafiltrates
CZE analysis was done in System II (Table 1). Peak identifications common to both CZE and HPLC profile (tentatively in CZE) same as in Fig. 1. Other neutral or cationic compounds in HPLC: a, creatinine; b, pseudouridine; c, hypoxanthine. Note differences of ordinate scale and injection volumes (0.6 μL in CZE, 25 μL in HPLC)

Fig. 3. Zone electropherograms (in System II; Table 1) of a 40-fold-diluted ultrafiltered uremic serum sample with different amounts of sodium chloride added: (top) 2.8 mmol/L (no Cl− added), (middle) 13.2 mmol/L, (bottom) 54.4 mmol/L
Tentative peak identifications same as in Fig. 1

Fig. 4. Corresponding CZE and HPLC profiles for uremic serum ultrafiltrates
CZE analysis was done in System II (Table 1). Peak identifications common to both CZE and HPLC profile (tentatively in CZE) same as in Fig. 1. Other neutral or cationic compounds in HPLC: a, creatinine; b, pseudouridine; c, hypoxanthine. Note differences of ordinate scale and injection volumes (0.6 μL in CZE, 25 μL in HPLC)

Tentative peak identifications: 1, hippuric acid; 2, p-hydroxyhippuric acid; 3, uric acid. Detection was at 254 nm. Ultrafiltered serum was diluted 1:1 with water in A, 1:8 in B and C. "AU" is A, absorbance

through co-elution of standards in CZE and, as reported earlier, by mass spectrometry and enzymatic peak shift in HPLC (4). Identification of the three prominent CZE peaks was supported by comparison with the prominent HPLC peaks, which were all detected at 254 nm. HA was eluted at t_m = 5 min in CZE and t_p = 20 min in HPLC. Total analysis time for the profiles is 8 min in CZE and 90 min in HPLC.

The length of the latter is due to the necessity of re-equilibrating the HPLC column after gradient development, before the next sample is injected. It should be noted, however, that in the HPLC profile, neutral, anionic, and cationic constituents of serum are measured simultaneously, whereas CZE selectively separates anionic substances under the present conditions—which is desirable, because earlier studies (8) have shown that especially the group of tubularly secreted organic anions may be of importance in the etiology of uremic manifestations.

The CZE profiles show a large, asymmetrical negative peak starting at migration time of about 1.5 min, which originates from the large amount of Cl− ions present in serum. Chloride is eluted as the first component because of its high absolute mobility of 80 × 10^-3 cm/V·s. Its concentration ordinarily is 110 mmol/L in serum water and varies only to a minor extent among most patients. This fact is of importance for the reproducibility of migration times. The presence of a major component in the injected sample may significantly influence migration times, as is illustrated in Figure 3. The electropherograms represent the analyses of diluted ultrafiltered sera to which different amounts of sodium chloride were added. The greater the amount of chloride added, the longer the migration times of the (other) sample constituents, the smaller their (diffusive) dispersion, and the closer their relative migration distances. Serum components eluting closely after the back edge of the chloride zone are very sharp, and peak-shape analysis, based on statistical moments, demonstrates theoretical plate numbers (34) of >1 000 000 for these compounds. Plate numbers measured for low-molecular-mass solutes, injected in the absence of a major component, are about 30 000 (26). We calculated that the plate number decreases rapidly with elution time for peaks eluting after the chloride zone. This delayed elution phenomenon—which is related to the sample composition upon injection and which is analogous to the "solvent effect" observed in gel filtration and in gas and liquid chromatography (35-38)—will be described elsewhere in more detail (Schoots et al., ms. submitted for publication). Peak heights of the minor components are higher when the amount of chloride added is greater (Figure 3). This "solvent effect" as expected, can be used to enhance detectability of specific (minor) sample constituents by adding a major component.
of appropriate nature and concentration.

The elution order for uric acid, \( p \)-hydroxyhippuric acid, and HA is reversed in CZE as compared with reversed-phase HPLC. In HPLC, under the given conditions (C18-modified silica, eluent pH 4) where the weak organic acids are only partly dissociated, the hydrophobic character of the solutes seems to determine the elution order: \( t_{\text{R, HA}} < t_{\text{R, PHA}} < t_{\text{R, UA}} \). In the CZE experiments the following order of increasing effective ionic mobility was found for the three organic acids, as analyzed in ultrafiltered serum: \( \mu_{\text{UA}} < \mu_{\text{PHA}} < \mu_{\text{HA}} \). The \( pK \) values (2.700, charge -1; 7.271, charge -2) and corresponding absolute mobilities (25.3 \( \times \) \( 10^{-5} \) and 55.1 \( \times \) \( 10^{-5} \) cm\(^2\)/V \( \cdot \) s, respectively) of HA have been reported by Hirokawa et al. (39). The first \( pK \) value, 2.7, is lower than the value reported (\( pK = 3.8 \)) in another publication (40). Effective mobility, as is expressed in migration time, depends on several factors such as effective molecular radius, \( pK \) value, and concentration or activity coefficient. Moreover, secondary equilibria such as complexation influence effective mobility and thus migration time. Therefore, the discrepancy between reported \( pK \) values cannot be judged from the present data for serum samples; measurements under standardized conditions are necessary (41).

Table 2 shows the repeatability (CVs) of migration times and peak areas in CZE of three components in uremic serum. At other concentrations these data are similar. Similar data have been reported earlier for the gradient HPLC analysis (20), the CVs being 0.6–1.0% and 3–6% for retention times and peak heights, respectively.

HA was assayed in various concentrations in the series of ultrafiltered uremic serum samples, and the CV for its migration time in CZE was 2.5% (\( n = 20 \), all samples run on the same day).

Quantification

HA was quantified in 10 samples of ultrafiltered serum from uremic patients who were undergoing hemodialysis. The ultrafiltered serum samples were diluted 10-fold before injection in the CZE procedure. The samples were all analyzed during one day, and calibration solutions were placed in between them. Peak area depends on migration velocity, because ultraviolet absorbance is a concentration-dependent detection principle. Therefore, in quantitative analysis based on peak area we used a constant electrical current, 35 \( \mu \)A, in all experiments. Quantification of HA by CZE was compared with quantification by HPLC, in which samples were diluted fivefold before injection. Injected amounts were 0.6 \( \mu \)L in CZE, 25 \( \mu \)L in HPLC—the latter being injected via an autosampler. For calibration, standard solutions were prepared by addition of HA to ultrafiltered normal serum (see Materials and Methods). Peak height is not linearly related to concentration in CZE under the present conditions, because solute peaks are intrinsically asymmetrical in CZE at these concentrations (i.e., \( >1 \) \( \mu \)mol/L) (30). The HA concentration range of the injected calibration solutions was 0 to 125 \( \mu \)mol/L. Although CZE peaks were asymmetrical at these concentrations, this does not influence peak area as long as sufficient resolution from

![Fig. 4. Method comparison of quantification of HA in CZE and HPLC (bottom) and calibration lines for quantification by CZE peak area (top) and by HPLC peak height (middle)](image-url)
other sample components is maintained. Therefore, HA concentrations up to 1250 μmol/L in ultrafiltered uremic serum can be measured because, in CZE, samples were diluted 10-fold before injection. Peak height has been found to be more reproducible than peak area in HPLC analysis (20). Therefore, peak area is used in CZE and peak height in HPLC. Calibration curves for HA in CZE and HPLC were obtained by injecting the calibration solutions in duplicate.

A linear model was fitted to the calibration data for HA, yielding the following relations:

\[ CZE: \text{Area}_{HA} = 437 + 174 \cdot (\text{concentration}_{HA}); \]
\[ r = 0.999 (P < 0.0001). \]

\[ \text{HPLC: Height}_{HA} = -481 + 60.83 \cdot (\text{concentration}_{HA}); \]
\[ r = 0.999 (P < 0.0001). \]

The \( y \)-axis intercepts were not significantly different from zero (by \( t \)-test) in either CZE and HPLC.

We compared the present CZE technique with HPLC (9, 21), with respect to quantification of HA. Figure 4 shows the outcome of this comparative study. The ultrafiltered serum samples were analyzed in duplicate both in CZE and in HPLC, and the data points represent mean values for these determinations. We found a good agreement for the quantification of HA. The regression line shown in Figure 4 follows the equation

\[ \text{Concn}_{HA,CZE} = -35.16 + 1.0860 \cdot (\text{Concn}_{HA,HPLC}); \]
\[ r = 0.981 (P < 0.0001). \]

The \( y \)-axis intercept was not significantly different from zero (\( t = -1.475, P = 0.18 \)), nor was the slope of the regression line (1.086) significantly different from unity (\( F = 1.29, P = 0.29 \)).

Conclusion

The present study demonstrates the reliability of a new, rapid, and selective method of analysis for ultraviolet-absorbing anionic constituents—especially HA—in uremic serum by CZE. The separation, based on differences in electrophoretic mobility of the anionic sample constituents, is therefore very suitable for the analysis of these types of compounds in uremic serum. HA, which is prominently present in uremic serum, can be quantified as accurately and precisely as with HPLC. Our method will facilitate the study of the usefulness of HA as a marker of accumulation of tuberculosis secreted organic acids and of its role in drug-protein binding in patients with chronic renal failure. The "solvent effect" we observed when there is a major component (chloride) in the injected sample may be useful for enhancing the detectability of minor components. Currently, commercial CZE equipment is available that enables organic acids in uremic serum to be resolved with even higher efficiency by use of separation capillaries with internal diameters of 75 μm or less and involving electrospray flow (42).

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