Protein Catabolic Rate May Underestimate Dietary Protein Intake in Hemodialysis Patients

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Introduction

Protein catabolic rate (PCR) or its normalized value, nPCR, is widely used to define nutritional status in dialysis patients [1]. This term PCR obviously is a misnomer and refers to the net PCR since the total PCR is as high as 300 g/day and does not necessarily bear a relationship with protein intake [2]. In the steady state, the net PCR has to be equal to the dietary protein intake as long as the method to calculate PCR includes all the body nitrogen output.

The usual method to calculate PCR is from nitrogen appearance rate [3] based on a formula developed from a detailed study of 5 dialysis patients with very low residual renal function who were judged to be in nitrogen balance [4]. In this study, total nitrogen output from stool, urine and dialysate and urea nitrogen appearance rate were measured independently and the following regression equation, which included both terms, was developed: urea nitrogen generation = 0.154PCR – 1.7. The authors recognized that in dialysate, nitrogen contained in urea, creatinine and uric acid accounted for only about 94% of total nitrogen measured by the Kjeldahl method, and that the remainder 6%, or about 1.4 g of nitrogen, could be in the form of amino acids. These losses of about 8.75 g of amino acids (1.4 × 6.25) are very similar to the losses of about 8 g of amino acids per treatment that were documented in the early 80s by a different group of investigators in hemodialysis patients using conventional cellulosic membranes and glucose-free dialysate [5]. Moreover, this regression equation agreed very closely with the plot derived from data in nondialyzed patients with stable chronic renal insufficiency gathered independently by another group of investigators [6]. This latter equation, urea generation rate = 0.149PCR – 1.2, differed from the above formula only by the magnitude of the intercept on the Y-axis; of interest, this difference of about 0.5 g of N per day is numerically almost identical with the losses of about 8 g of amino acids per dialysis (0.5 g/day × 7 days divided by 3, the number of dialysis per week = 1.16 g of N × 6.25 = 7.3 g of amino acid per treatment).

Recently, several authors have documented a significant increase in the losses of amino acids and protein into the dialysate as the result of repetitive processing of dialyzers [7–10]. Thus, with the current use of high-flux dialyzers and with at least 80% of the dialysis units in this country practicing dialyzer reuse, this formula may not measure accurately PCR since a large amount of nitrogen loss, either in the form of amino acids or protein, is not included in this calculation. The purpose of this presentation is to demonstrate that the current method used to calculate PCR in hemodialysis patients may indeed underestimate this parameter and hence dietary protein intake [3].
Methods

Review of all the literature reporting on the loss of protein and/or amino acids during hemodialysis.

Results

There were four reports which described the loss of protein and/or amino acids during hemodialysis and are detailed below [7–10]. Ikizler et al. [7] measured the losses of amino acids into the dialysate with three different membranes: low-flux cuprophane (Terumo T150), low-flux polymethylmethacrylate (Toray B2) and high-flux polysulfone (Fresenius F80) and found the following averages: 7.2 ± 2.6, 6.1 ± 1.5 and 8.0 ± 2.8 g of amino acids per dialysis session respectively. When polysulfone dialyzers were studied at the 6th reuse, an average loss of 12.2 ± 4.4 g of amino acids per dialysis occurred. Thus, these results suggest that currently available dialyzers, when first used, lose about the same amount of amino acids as those in the past; reuse to 6 times increases the losses by 50% and it is quite possible that further reuse will increase the losses of amino acids even further. The same group also measured albumin losses into the dialysate with increasing reuse number as follows: the average amount of albumin lost was 0.98 ± 0.9 g/treatment (range 0–18 g) below the 10th reuse, 1.94 ± 1.5 g (range 0.3–8.6 g) for 10–14 reuses, 4.3 ± 3 g (range 0.5–9.4 g) for 15–19 reuses, 9.3 ± 5.5 g (range 1.1–20.8 g) for 20–24 reuses and 11 ± 7.9 g (range 1.1–25.6 g) for over 24 reuses. Bleach and formaldehyde were used to process dialyzers in this study.

Kaplan et al. [8] measured the dialysate protein losses from polysulfone dialyzers (F80) undergoing repetitive processing with bleach and formaldehyde. Mean total protein losses per treatment were: 1.2 ± 0.3 g during the 1st use, 1.8 ± 0.3 g during the 5th use, 2.9 ± 0.4 g during the 10th use, 6.4 ± 0.5 g during the 12th to 13th use, 10.4 ± 1.1 g during the 16th through 20th use and 17.5 ± 1 g during the 23rd through 25th use. To further validate the results, dialysate samples were also evaluated for albumin concentration which went from unmeasurable for the 1st and 5th use to 14.4 ± 3.2 mg/dl after 23–25 reuses. In those samples where albumin and total protein were measured, the ratio of albumin to total protein was 53%. A random sample of dialyzers processed without bleach revealed only minimal protein losses after repetitive reuses.

Kaysen et al. [9] in a study of mechanisms of hypoalbuminemia in hemodialysis patients reported albumin losses of up to 5 g/treatment with more than 10 reuses, up to 10 g/treatment for more than 20 reuses and up to 13 g/treatment for more than 30 reuses. The dialyzers were processed with bleach and formalin in this study.

Kirschbaum [10] reported a significant decline in serum albumin after conversion from low-flux to high-flux dialyzers in a large group of hemodialysis patients. Dialyzers were reprocessed with bleach and formaldehyde. He found that among 12 dialysates, the differences between total nitrogen and urea nitrogen averaged 29 mg/l. Because the average dialysis session lasted 210 min and the dialysate flow was 500 ml/min, the average loss of nonurea N was 3 g/dialysis (29 mg/l × 105 l), which would correspond to a protein or amino acid equivalent of 19 g/dialysis! (3 × 6.25). The drop in serum albumin was most likely the result of negative nitrogen balance induced by excessive losses in the dialysate although the author states that he did not detect protein in the dialysate using a pyrogallol red reagent [10].

Discussion

The four studies discussed in the above section suggest that a significant amount of nitrogen, either in the form of amino acids or protein, is actually lost during hemodialysis using high-flux dialyzers that have been processed with bleach and formaldehyde. In other words, there are enough reasons to suspect that we may indeed be underestimating the total nitrogen output with our current calculation of PCR during hemodialysis. If we take a patient who is being dialyzed for the 23rd time with the same polysulfone dialyzer, loss of protein may be 10.4 g (using Kaplan’s data) and extra losses of amino acids are at least 4 g (using Ikizler’s data); these losses represent an extra 15 g/dialysis or 6.5 g/day (15 g × 3/7), which are not accounted for in the current calculation of PCR. If the losses of amino acids increase significantly after the 6th reuse, say to 16 g/dialysis, this extra loss of nitrogen may become very significant, as much as 19 g of protein equivalents per dialysis or 8.1 g/day in the previous example. Thus, a hemodialysis patient who uses dialyzers processed with bleach and formaldehyde can be losing as much protein per day as the counterpart CAPD patient [11]. This fact has not been widely appreciated in the literature and has been completely ignored in the current methods used to calculate PCR.

As mentioned above, these significant losses of extra nitrogen into the dialysate are not being detected by our current estimate of PCR based on the nitrogen appear-
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In summary, we still believe that PCR is a very useful tool to approximately determine the daily protein intake of dialysis patients. We should keep in mind, however, that this parameter, as is currently calculated, may under-estimate true dietary protein intake.

References


10 Kirschbaum B: The decline in serum albumin after conversion to high flux, high effective dialysis. Artif Organs 1994;18:729–735.


