

Impaired endotoxin-induced interleukin-1 β secretion, not total production, of mononuclear cells from ESRD patients

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Impaired endotoxin-induced interleukin-1 β secretion, not total production, of mononuclear cells from ESRD patients. Lipopolysaccharide (LPS)-induced interleukin-1 beta (IL-1 β) and tumor necrosis factor alpha (TNF α) production and secretion from peripheral blood mononuclear cells (PBMC) were determined in a longitudinal study with repeated measurements in PBMC from patients with chronic uremia not on hemodialysis ($N = 8$), end-stage renal disease (ESRD) patients ($N = 8$), and healthy controls ($N = 7$). ESRD patients were studied while using low-flux Cuprophane dialyzers and again using high-flux AN 69 dialyzers. Total (cell-associated plus secreted) LPS-induced IL-1 β production was enhanced in uremic patients, but similar to controls in ESRD patients on Cuprophane. In contrast, LPS-induced IL-1 β secretion (secreted amounts in % of total production) was similar to controls in uremic patients, but significantly reduced in ESRD patients on Cuprophane ($P < 0.01$). During AN 69 hemodialysis, LPS-induced total IL-1 β production remained unchanged but IL-1 β secretion increased significantly ($P < 0.05$) compared to Cuprophane dialysis. Increased IL-1 β secretion coincided with a suppression in PGE₂ synthesis ($P < 0.02$). Similarly, blockade of endogenous PGE₂ by indomethacin increased LPS-induced IL-1 β secretion ($P < 0.01$) but did not enhance total IL-1 β production in PBMC from controls and patients on Cuprophane hemodialysis. Neither total production nor secretion of TNF α was different comparing the three study groups. We conclude that LPS-induced IL-1 β secretion, but not total production, is impaired in PBMC from ESRD patients on long-term Cuprophane hemodialysis. This functional change in the PBMC response is specific for IL-1 β , not due to uremia *per se* but hemodialysis-dependent and reversible. Hemodialysis with AN 69 suppresses endogenous PGE₂ synthesis in PBMC which is associated with increased LPS-induced IL-1 β secretion in the presence of unchanged total IL-1 β production. We speculate that PGE₂ could inactivate the IL-1 β converting enzyme which is essential for processing and secretion of mature IL-1 β .

The *in vitro* production of proinflammatory cytokines such as IL-1 and TNF from PBMC is one way to study differences in the response of these cells between chronic disease and health [1, 2]. As cytokines play an important role in cell-cell communication, the total cellular production of IL-1 and TNF as well as the release of these mediators from activated PBMC are important in the orchestration of the host response system [3, 4]. Although TNF α has a signal peptide similar to that of secretory proteins, TNF α can remain inside the cell [5, 6]. However, in sharp

contrast, the two interleukin-1 molecules (IL-1 α and IL-1 β) clearly lack signal peptides [4]. Despite the missing signal peptide, as much as 60% of IL-1 β has been found in the supernatant of lipopolysaccharide- (LPS) stimulated PBMC of healthy volunteers *in vitro* [7, 8].

Cytokine production of PBMC has been used to describe the effect of orally administered drugs (such as cyclooxygenase inhibitors) [9] or dietary supplementation [1, 10]. However, there is a remarkable intra-individual variability in LPS-induced IL-1 β production from PBMC studied on three different days within one week [1, 9]. Therefore, in order to judge whether a change in PBMC function due to disease or therapy is meaningful, it is a prerequisite to describe the intra-individual variability of *in vitro* PBMC cytokine production under stable conditions over a defined period of time.

Patients on chronic hemodialysis therapy have elevated cell-associated levels of cytokines such as IL-1 β and TNF α in PBMC determined by immediate lysis after separation or after incubation [2, 11–13]. Using single point measurements, we have demonstrated that PBMC from patients on hemodialysis with low-flux Cuprophane dialyzers produce similar amounts of total (secreted plus cell-associated) IL-1 β in response to LPS *in vitro* compared to PBMC from an age-matched control group [14]. In that study, LPS-stimulated PBMC secreted significantly less IL-1 β than PBMC from healthy controls (controls, $56.2 \pm 8.3\%$; Cuprophane hemodialysis, $31.2 \pm 6.3\%$, $P < 0.02$). The reduction in IL-1 β secretion seems to be stimulus dependent since *Staphylococcus epidermidis* induced the release of approximately 90% of total IL-1 β in both groups (controls, $88.1 \pm 4.1\%$; Cuprophane hemodialysis, $89.0 \pm 2.7\%$).

The mechanisms involved in the reduced ability to release IL-1 β in response to LPS remained unclear. Furthermore, it has not been determined whether the change in PBMC function is due to uremia *per se* or to repeated stimulation by the hemodialysis procedure. Also, the question whether the inflammatory nature of the dialyzer membrane-induced dysfunction [14, 15] was reversible remained unanswered.

In order to address these issues, we studied the variability of TNF α and IL-1 β production and release from LPS stimulated PBMC from healthy controls, uremic patients with chronic renal failure as well as from patients with end-stage renal disease (ESRD) on chronic hemodialysis. We used repeated measurements in PBMC per individual during a period of eight weeks. The

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same ESRD patients were studied in two consecutive eight-week periods in which they were dialyzed using low-flux Cuprophan versus high-flux AN 69 dialyzers. We measured PGE₂ concentrations in PBMC supernatants and studied the effect of indomethacin on LPS-induced IL-1 β and TNF α production and release from PBMC.

Methods

Patients and study groups

Two control groups were included in the study. For the longitudinal study over a time period of 8 weeks, the first group included seven healthy male volunteers with a mean age of 32.4 years (27 to 42). Blood was drawn from an antecubital vein once a week in the morning of the same weekday. In a second control group, 10 healthy volunteers, seven males, three females, donated blood once for PBMC separation in order to study the effect of indomethacin on LPS-induced IL-1 β and TNF α production and secretion. None of the volunteers took any medication (cortison, aspirin or other prostaglandin inhibitors) throughout the study period.

The first study group included eight uremic patients with chronic renal failure, two males, six females with a mean age of 53.9 years (31 to 78) who gave informed consent. The diagnosis of the kidney diseases were polycystic kidney disease in one, diabetic nephropathy in two, chronic glomerulonephritis in three and chronic interstitial nephropathy in two. These patients were studied on each occasion for which they were scheduled for a routine blood test in the out patient unit. This were three occasions during the eight-week study period with at least a two week interval between two samples. In this group, serum concentrations were 28.2 ± 4.0 mmol/liter for urea and 699 ± 168 μ mol/liter for creatinine. The creatinine clearance was 12.7 ± 4.9 ml/min.

The second group included eight male ESRD patients, mean age 52.9 years (34 to 64), who gave informed consent. These patients were on hemodialysis therapy with Cuprophan dialyzers for an average time of 61 months (16 to 132) when entering the study. The diagnosis leading to end-stage renal failure was polycystic kidney disease in four, chronic glomerulonephritis in three and chronic interstitial nephritis in one. Patients with hepatitis B or C, active hyperparathyroidism or requiring anti-inflammatory medication were not included.

During the first study period of eight weeks, hemodialysis was continued using Cuprophan[®] dialyzers (GF 120 H[®], Gambro, Lund, Sweden) for three hemodialysis sessions per week, with four to five hours each. Blood was drawn once per week from the arterio-venous fistula before the start of the dialysis session following the long inter-dialytic interval. Then all patients switched to AN 69[®] dialyzers (Filtral 12[®], Hospal, Nürnberg, Germany). Following a washout period of eight weeks in which no blood was drawn, the patients entered the second study period of eight weeks where blood was taken again once a week before hemodialysis after the long inter-dialytic interval. The effectiveness of the hemodialysis as indicated by the treatment time, blood flow in the dialysis machine, post-pre-treatment differences in serum urea and the weight loss over treatment, was the same in both study periods (Table 1). In addition, the endotoxin contamination of the dialysate (Limulus test: Coatest[®] Endotoxin, KabiVitrum GmbH, München, Germany), which is recognized to

Table 1.

Parameter	Cuprophan hemodialysis	AN 69 hemodialysis
Δ Body weight kg	3.00 ± 0.18	3.05 ± 0.24
RR pre-HD mm Hg	$149 \pm 2.1/88.9 \pm 2.7$	$154.5 \pm 3.9/90.6 \pm 2.3$
RR post-HD mm Hg	$140.6 \pm 4.8/86.3 \pm 3.5$	$140.6 \pm 3.9/85.8 \pm 4.4$
Urea pre-HD mmol/liter	25.36 ± 0.8	26.34 ± 1.1
Urea post-HD mmol/liter	8.94 ± 0.43	8.57 ± 0.54
Δ Urea mmol/liter	16.43 ± 0.71	17.77 ± 0.84
Dialysate endotoxin ^a pg/ml	19.5 ± 1.5	20.3 ± 0.6

Numbers represent the means \pm SEM of $N = 8$ averaged values per patient.

^a Dialysate samples were taken from the dialyzer outlet after 3 hours of hemodialysis

be a potent inducer of PBMC cytokine production [16, 17], was less than 21 pg/ml and did not differ significantly during the two study periods (Table 1).

To test the effect of indomethacin on LPS-induced IL-1 β production and secretion, PBMC were separated from 10 ESRD patients on Cuprophan hemodialysis (6 males, 4 females). These patients are different from those in the study group mentioned above.

Preparation of human blood mononuclear cells (PBMC)

For the longitudinal study blood was drawn under aseptic conditions into pyrogen-free tubes containing EDTA (1.5 mg/ml of blood) (Sarstedt, Nümbrecht, Germany) to which the proteinase inhibitor aprotinin was added (0.67 TIU/ml of blood) (Sigma, St. Louis, MO, USA). The blood was kept on ice and subsequently processed within 60 minutes. To study the effect of indomethacin, blood was drawn into pyrogen-free syringes containing 10 IU/ml heparin.

From each blood sample PBMC were separated by Ficoll-Hypaque (Seromed, Biochrom KG, Berlin, Germany) centrifugation. The PBMC were re-suspended to a concentration of 5×10^6 cells/ml in ultrafiltered [18] tissue culture medium (MEM, Wtittaker M.A. Bioproducts Inc., Walkersville, MD, USA) supplemented with 2 mM L-glutamine, 10 mM HEPES, 100 U/ml penicillin, 100 μ g/ml streptomycin (all from Wtittaker M.A.) and 2% heat-inactivated human AB serum. One milliliter aliquots (final concentration 2.5×10^6 PBMC/ml) were incubated in 24-well flat bottom polystyrene plates (Nunc, A/S Nunc, Roskilde, Denmark). In the longitudinal study, PBMC were stimulated with 100 ng/ml LPS from *E. coli* (phenol-extracted LPS: O55:B5, Sigma), a dose which induces maximal cytokine production in normal PBMC [8]. In additional experiments, PBMC were stimulated with 1 to 100 ng/ml LPS in the presence and absence of 0.5 μ g/ml indomethacin (Sigma).

During 18 hours of PBMC incubation, temperature was maintained at 37°C in an atmosphere containing 5% CO₂. After this period, cell culture supernatants were removed, transferred into 1.5 ml Eppendorf tubes and centrifuged for five minutes at 4°C to remove remaining cells. Following this centrifugation step, supernatants containing the extracellular amounts of cytokines were transferred into new 1.5 ml eppendorf tubes and frozen at -70°C. The cell pellets (in approximately 20 μ l of remaining medium) in the eppendorf tubes were re-suspended in 1 ml of fresh MEM, added to the adhering PBMC in the corresponding wells of the 24 well plates and kept frozen at -70°C. On the day of the cytokine

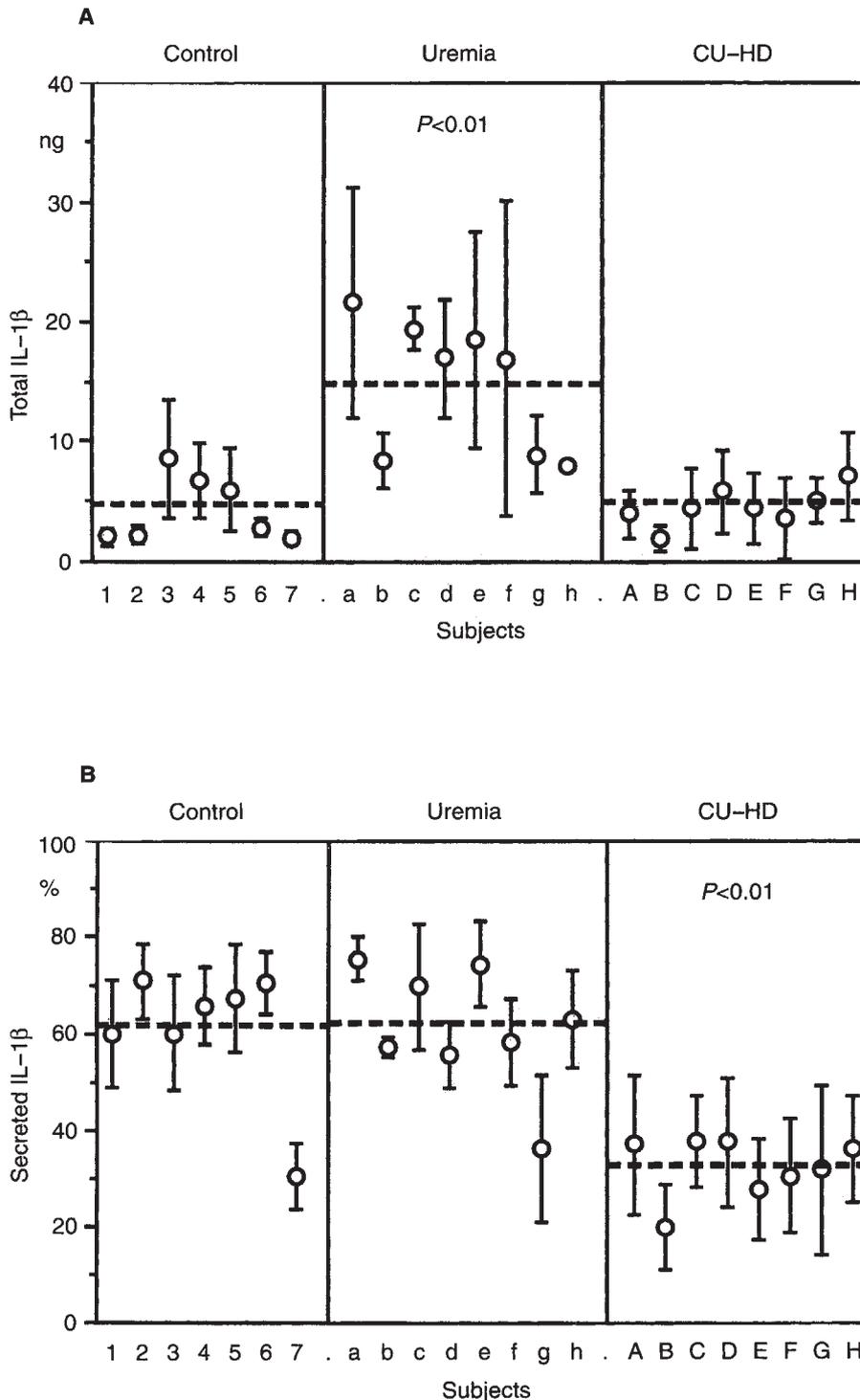


Fig. 1. Total LPS (100 ng/ml)-induced production (A) and release of IL-1 β (B) from PBMC. Each number or letter under the horizontal axis represents a different individual (subject). Symbols represent the mean \pm SD of 6 determinations per subject in controls, 3 determinations per subject in uremic patients (uremia), and 6 to 8 determinations per subject in ESRD patients on Cuprophane hemodialysis (CU-HD). Broken horizontal lines indicate the means per study group: Controls $N = 7$, uremia $N = 8$, and CU-HD $N = 8$. Total IL-1 β production is expressed as ng/ 2.5×10^6 PBMC/18 hours of incubation; secreted amounts are depicted as percent of total production. Total IL-1 β production was higher in uremia compared to controls and CU-HD ($P < 0.01$); IL-1 β secretion was significantly reduced in CU-HD ($P < 0.01$) compared to controls and uremia.

measurements by radioimmunoassays the 24 well plates were subjected to three freeze- (-70°C) thaw (37°C) cycles in order to lyse the PBMC for measurement of cell-associated cytokines.

Cytokine measurements

IL-1 β and TNF α were measured by radioimmunoassay (RIA) in cell culture supernatants and cell lysates as described previously [7, 19]. The polyclonal antibodies used in each RIA (rabbit

anti-human TNF α and rabbit anti-human IL-1 β) were gifts from Prof. C.A. Dinarello (New England Medical Center, Boston, MA, USA). The anti-human IL-1 β antiserum detects the mature IL-1 β peptides (17 kD and 21 kD) as well as 50% of the IL-1 β precursor which is not recognized by monoclonal antibodies produced against the mature form of IL-1 β [20, 21]. The sensitivity of both RIAs is approximately 80 pg/ml in PBMC culture supernatants and cell lysates.

To validate the cytokine assays for repeated measurements, we determined the intra- and inter-assay variability expressed by the coefficient of variation ($V = \text{sd}/\text{mean} \times 100$). Aliquotes of the same PBMC supernatant and corresponding cell-lysate were assayed 10 times in one assay, yielding the intra-assay variability of $V_{\text{intra}} = 11.4\%$; testing the same sample in ten different assays, the inter-assay variability was $V_{\text{inter}} = 11.7\%$. Hence, to obtain the total cytokine production level, we added the cell-associated levels to the extracellular levels. This number was not significantly different from the actual measured total production in PBMC when supernatants and cells were lysed together (calculated total IL-1 $\beta = 9.9 \pm 1.0$; measured total IL-1 $\beta = 10.9 \pm 1.3 \text{ ng}/2.5 \times 10^6 \text{ PBMC}/18 \text{ hr}$; $N = 10$).

Prostaglandin E₂ (PGE₂) in the supernatants of LPS-stimulated PBMC was measured by specific ELISA as previously described [22]. The detection limit of this assay is 0.25 ng/ml.

Statistics

Results are given as means \pm standard deviation (sd). Statistical significance was assessed using one Factor ANOVA analysis. Post-hoc method of data evaluation was Fisher's PLSD test. Differences were considered to be significant when $P < 0.05$.

Results

Decreased release of LPS-induced IL-1 β from PBMC associated with hemodialysis

Figure 1 depicts LPS-induced IL-1 β production as well as release from PBMC in the three study groups. The total LPS-induced IL-1 β production (Fig. 1A) was similar to that in controls ($4.3 \pm 2.7 \text{ ng}/2.5 \times 10^6 \text{ PBMC}/18 \text{ hrs}$) and ESRD patients (4.5 ± 1.6) but higher in uremic patients (14.8 ± 5.5 , $P < 0.01$). Secretion of LPS-induced IL-1 β (Fig. 1B) was significantly reduced in PBMC from ESRD patients on chronic hemodialysis using low-flux Cuprophan dialyzer membranes ($32.5 \pm 6.3\%$; $P < 0.01$) compared to that of controls ($60.8 \pm 14.1\%$) and uremic patients ($61.4 \pm 12.6\%$).

There was intra-individual variability in total IL-1 β production in each of the three study groups (Fig. 1A). The coefficient of variation was significantly higher ($P < 0.02$) in ESRD patients using Cuprophan dialyzers ($62.1 \pm 16.7\%$) compared to uremic patients ($43.4 \pm 23.9\%$) and healthy controls ($40.1 \pm 13.5\%$). The intra-individual variability was significantly reduced in each study group when the levels of extracellular IL-1 β were expressed as percent of total LPS-induced IL-1 β production. Similar to the differences in total IL-1 β production, the intra-individual variability for secreted amounts of IL-1 β was higher in ESRD patients ($36.3 \pm 6.34\%$; $P < 0.01$) than in normal controls ($15.5 \pm 5\%$) and in uremic patients ($15.7 \pm 11.8\%$). In six out of seven control individuals and seven out of eight uremic patients IL-1 β release from PBMC was greater than 50%. However, despite the higher intra-individual variability, PBMC from patients on hemodialysis with Cuprophan dialyzers consistently secreted less than 50% of LPS-induced IL-1 β .

LPS-induced TNF α production and release from PBMC

In contrast to IL-1 β , neither the total production nor the release of LPS-induced TNF α from PBMC differed significantly between the three study groups (Fig. 2). Total TNF α production was $6.8 \pm 3.9 \text{ ng}/2.5 \times 10^6 \text{ PBMC}/18 \text{ hours}$ in controls, 5.2 ± 3.0

ng in uremic patients and $8.8 \pm 3.7 \text{ ng}$ in ESRD patients. TNF α release from PBMC was $91.1 \pm 4.6\%$ in controls, $88.9 \pm 2.5\%$ in uremic patients and $88.6 \pm 4.7\%$ in ESRD patients on Cuprophan dialyzers.

The intra-individual variability of LPS-induced TNF α production (Fig. 2A) was not significantly different from the one described for IL-1 β . The coefficient of variation was $53.3 \pm 22.1\%$ in healthy controls, $32.3 \pm 18.8\%$ in uremic patients and $58.5 \pm 14.4\%$ in ESRD hemodialysis. When the variability was calculated for secreted TNF α (Fig. 2B), the levels were in the range of 10%, which falls into the variability of the method used.

LPS-induced PGE₂ production from PBMC

As shown in Figure 3, LPS-induced PGE₂ production was decreased in PBMC from uremic ($20.5 \pm 8.3 \text{ ng}$; $P < 0.05$) and ESRD patients on hemodialysis with Cuprophan dialyzers ($15.2 \pm 9.7 \text{ ng}$; $P < 0.01$) or AN 69 dialyzers ($3.9 \pm 5.2 \text{ ng}$; $P < 0.001$) compared to healthy controls (30.7 ± 9.7). The intra-individual variability was higher for PGE₂ than for IL-1 β and TNF α with coefficients of variation of $74.4 \pm 27.0\%$ in controls, $55.8 \pm 43.3\%$ in uremic patients, $78.8 \pm 25.7\%$ in Cuprophan hemodialysis, and $61.4 \pm 26.1\%$ in AN 69 hemodialysis.

The suppression of LPS-induced IL-1 β release is reversible

The ESRD patients on Cuprophan dialyzers were switched to AN 69 dialyzers and were studied again after a washout period of eight weeks. Figure 4 depicts the average of eight values for each patient for total LPS-induced IL-1 β and TNF α production and secretion as well as PGE₂ synthesis. The intra-individual variability was not significantly different between the two hemodialysis periods. The total IL-1 β and TNF α production decreased in six out of eight patients but increased in the same two patients, resulting in a non-significant change when comparing the Cuprophan with the AN 69 period. Furthermore, TNF α release did not change under the two treatment conditions. After 8 to 16 weeks of hemodialysis using AN 69 dialyzers, IL-1 β release increased consistently in each of the eight patients ($P < 0.05$). This reversal in IL-1 β release coincided with a decrease in PGE₂ production in seven out of eight patients ($P < 0.02$) when Cuprophan dialyzers were replaced by AN 69 (Fig. 4).

As there was no time related trend to increased IL-1 β release and decreased PGE₂ production during the AN 69 period, the washout period of eight weeks was sufficient to reach stable conditions on AN 69 hemodialysis. Therefore, it was legitimate to average the eight values per study period.

Inhibition of endogenous PGE₂ production by indomethacin enhances IL-1 β release from PBMC

Although the absolute concentrations of PGE₂ are reduced in PBMC supernatants from ESRD patients compared to normal controls, the data shown in Figure 4 suggest that relative changes in PGE₂ synthesis influence IL-1 β secretion. In order to test this hypothesis, PBMC from healthy volunteers were incubated in the absence and presence of indomethacin (Table 2). Inhibition of PGE₂ did not enhance total IL-1 β production in response to LPS (Fig. 5). However, in these same cultures, IL-1 β secretion increased in the presence of indomethacin in response to 10 ng/ml

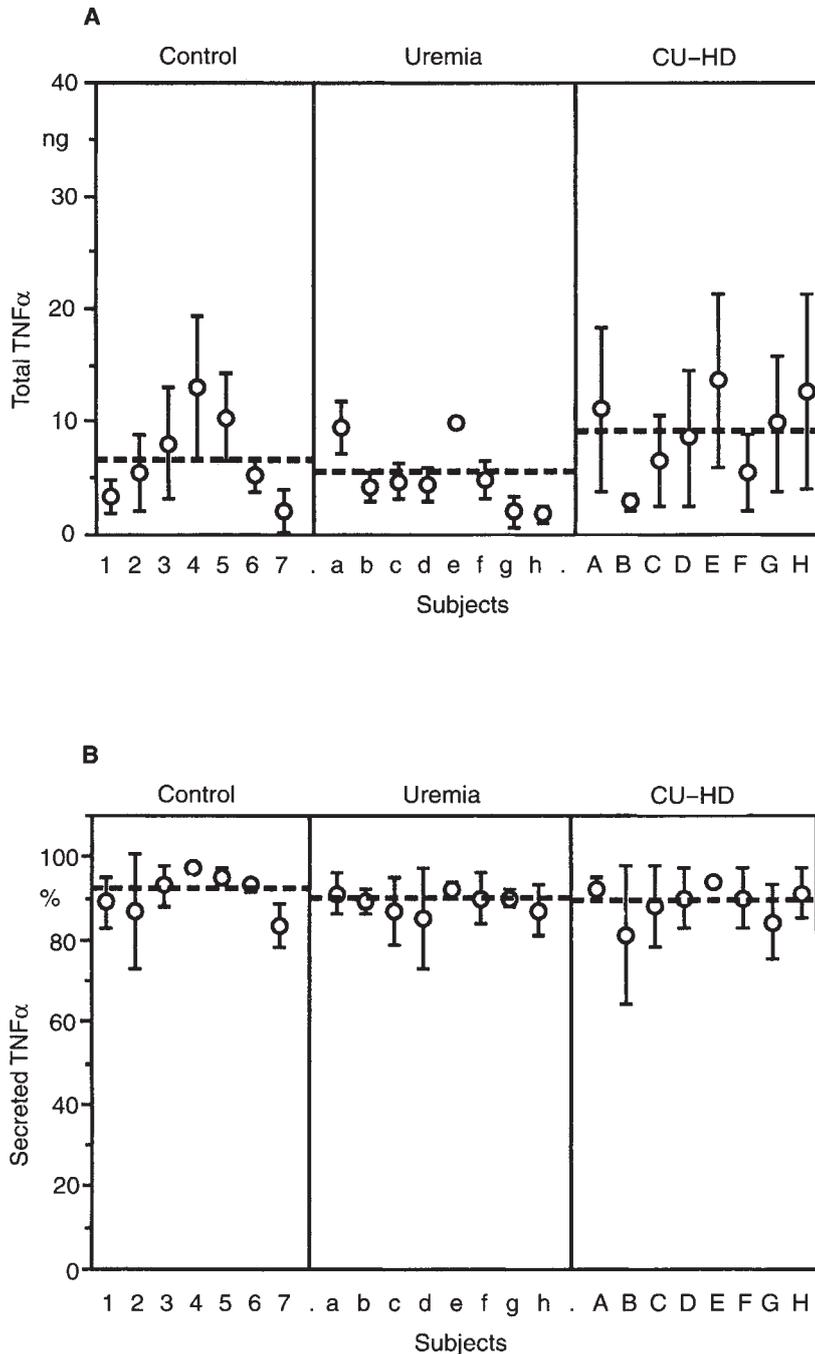


Fig. 2. Total TNF α production (A) and TNF α release (B) from LPS (100 ng/ml)-stimulated PBMC of healthy controls (N = 7), uremic patients (N = 8) and ESRD patients on Cuprophane hemodialysis (CU-HD, N = 8). Symbols represent the mean \pm SD of 6 determinations per control subject, 3 determinations per uremic patients (uremia), and 6 to 8 determinations per ESRD patient on Cuprophane hemodialysis (CU-HD). Broken horizontal lines indicate the means per study group. Differences in total TNF α production or TNF α secretion between the three groups were not statistically significant.

LPS from $53.1 \pm 17.6\%$ to $63.1 \pm 15.7\%$ ($P < 0.02$). In response to 100 ng/ml LPS, IL-1 β secretion increased from $57.7 \pm 19.1\%$ to $71.1 \pm 16.2\%$ ($P < 0.01$; Fig. 5). In response to 100 ng/ml LPS, the total TNF α production increased from 10.7 ± 3.47 ng to 14.6 ± 5.03 ng in the presence of indomethacin ($P < 0.01$), but secretion of TNF α did not change significantly.

When PBMC from ESRD patients on Cuprophane hemodialysis were studied, total IL-1 β production in response to 100 ng/ml LPS was not enhanced by indomethacin, but IL-1 β secretion increased from $44.2 \pm 17\%$ to $57.9 \pm 18\%$ ($P < 0.01$) in the presence of indomethacin (Fig. 5).

Discussion

Suppressed IL-1 β secretion associated with Cuprophane hemodialysis

LPS-stimulated PBMC from ESRD patients on chronic hemodialysis with low-flux Cuprophane dialyzers are suppressed in their ability to secrete IL-1 β , but not TNF α . These data confirm previous observations [14] and are in agreement with other studies describing reduced extracellular levels of IL-1 β in the supernatants of LPS-stimulated PBMC from ESRD patients [15, 23, 24].

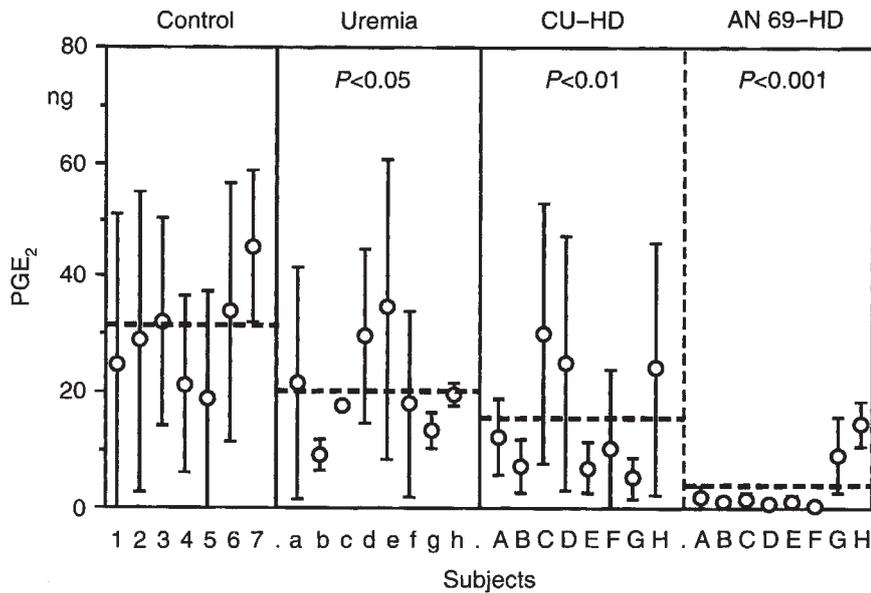


Fig. 3. LPS-induced PGE₂ synthesis by PBMC from healthy controls, uremic patients and ESRD patients on hemodialysis with Cuprophan and AN 69 dialyzers. Symbols represent the mean \pm SD of 6 determinations per control subject, 3 determinations per uremic patient, and 6 to 8 determinations per ESRD patient on Cuprophan hemodialysis (CU-HD). Broken horizontal lines indicate the means per study group. Compared to controls, PGE₂ synthesis was reduced in uremia ($P < 0.05$), Cuprophan hemodialysis ($P < 0.01$) and AN 69 hemodialysis ($P < 0.001$). Switching the same ESRD patients from Cuprophan to AN 69 decreased *in vitro* LPS-PGE₂ synthesis ($P < 0.02$).

On the other hand, PBMC from uremic patients not on hemodialysis are not different from those of healthy controls. Thus, the impaired IL-1 β release is not caused by uremia *per se* but is hemodialysis-dependent. In the present study, suppressed IL-1 β secretion was reversed when low-flux Cuprophan dialyzers were replaced by highly permeable AN 69 dialyzers for at least eight weeks. When four out of eight ESRD patients were studied again after one year of AN 69 hemodialysis, their PBMC secreted 60% of LPS-induced IL-1 β which is similar to that in healthy or uremic PBMC (unpublished observation). More IL-1 β secretion occurred without a change in total LPS-induced IL-1 β production and this reversal was associated with a decrease in PGE₂ synthesis. Similarly, inhibition of endogenous PGE₂ by indomethacin increased IL-1 β secretion from LPS-stimulated PBMC of healthy controls and ESRD patients on Cuprophan. These data suggest that in ESRD patients as well as in healthy volunteers, down-regulation of endogenous PGE₂ increases IL-1 β secretion rather than total IL-1 β production by PBMC. However, as PGE₂ synthesis is reduced in PBMC from ESRD patients compared to normal controls, the relative changes in PGE₂ production as opposed to the absolute amounts seem to influence IL-1 β secretion.

We studied PBMC *in vitro* approximately 2.5 days after termination of the preceding hemodialysis session. Therefore, our data indicate that blood-dialyzer interactions during hemodialysis may cause functional changes in PBMC which persist as long as two days after the end of the hemodialysis procedure. Switching the dialyzer membrane may persistently modify the PBMC response to LPS-stimulation. It remains to be established, however, whether the increased IL-1 β secretion observed under AN 69 hemodialysis reflects an improved or even a normalized PBMC function in ESRD patients.

Mechanisms involved in the suppression of IL-1 β secretion from PBMC of ESRD patients on long-term Cuprophan hemodialysis are unknown. However, as the dialyzer clearance may play a role, it is important to state that serum urea levels were not different comparing the two hemodialysis conditions. Therefore, it is unlikely that the modified PBMC function during the AN 69

phase was due to better dialyzer clearance of these small molecular substances. We cannot exclude, however, a beneficial effect of an increased clearance for undefined uremic toxins with a molecular weight of up to 15,000 Daltons which may have an inhibitory effect on PBMC and which are cleared by the high-flux AN 69 but not by the low-flux Cuprophan membrane.

The role of PGE₂

The question whether PGE₂ synthesis from PBMC in uremic patients and ESRD patients is altered is controversial. There are reports of increased [25], unchanged [24] and decreased [26] PGE₂ production in PBMC from ESRD patients compared to cells of healthy controls. In the present study, PGE₂ production was reduced in cells from patients during Cuprophan hemodialysis and decreased further during AN 69 usage in the same patients. This further reduction was associated with an increase in IL-1 β secretion. However, when individual data were analyzed, we could not demonstrate a significant inverse correlation between decreasing PGE₂ levels and increasing secretion of IL-1 β under the two dialysis conditions. In a second set of experiments, blockade of endogenous PGE₂ production by indomethacin resulted in increased IL-1 β secretion, not total IL-1 β production, from cells from normal subjects as well as from ESRD patients on Cuprophan hemodialysis. These data suggest that endogenous PGE₂ inhibits LPS-induced IL-1 β secretion but not total synthesis.

Previous reports described that PGE₂ down-regulates LPS-induced IL-1 β production [27]. In that study, the secretion of IL-1 into the cell supernatant was measured and, therefore, is consistent with our finding. As in response to 100 ng/ml LPS total TNF α production (but not IL-1 β) was significantly enhanced in the presence of indomethacin, the results presented here are in agreement with Endres et al who described that PGE₂ influences IL-1 β and TNF α production in normal PBMC differently [28].

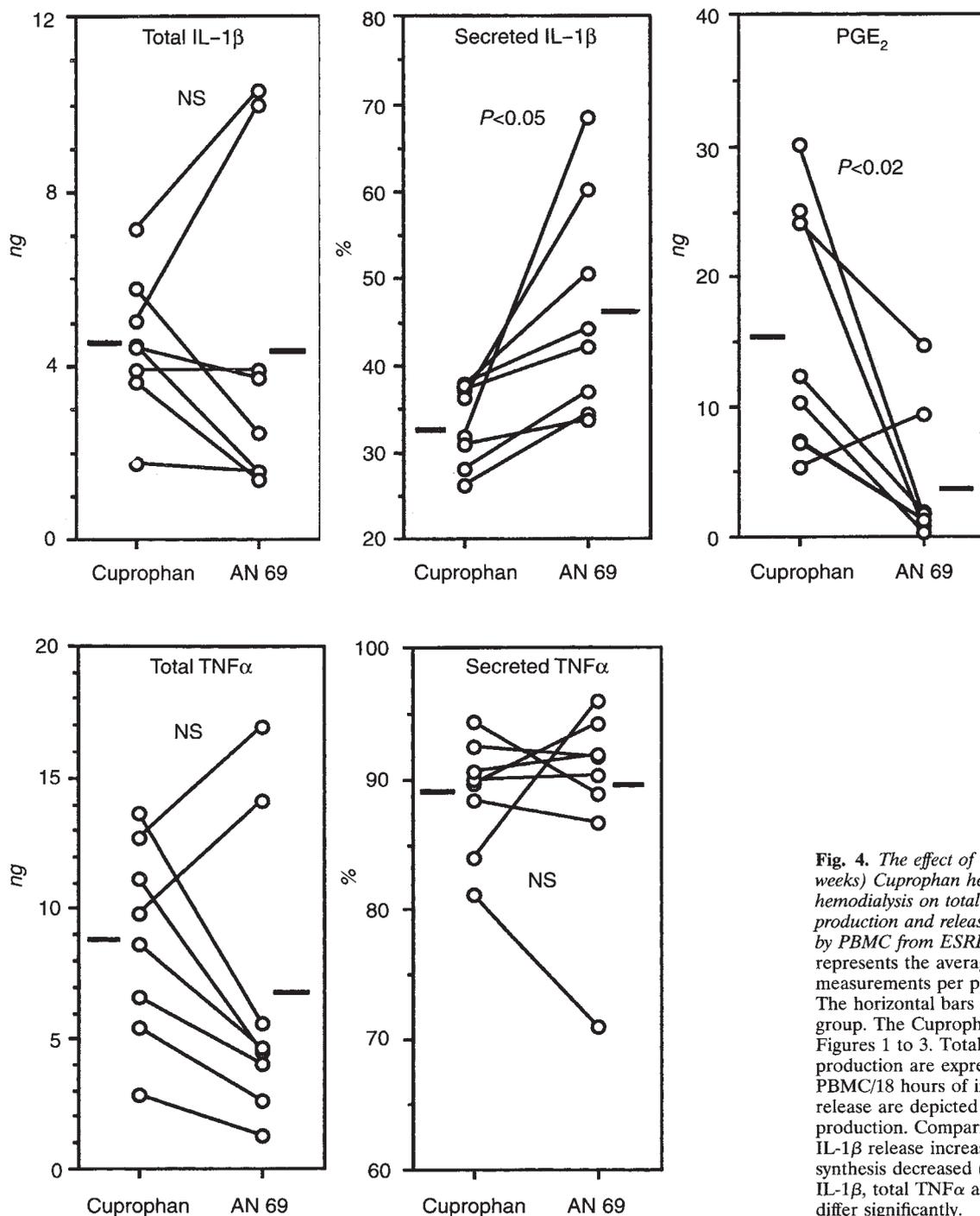


Fig. 4. The effect of chronic (a minimum of 8 weeks) Cuprophan hemodialysis versus AN 69 hemodialysis on total in vitro IL-1 β and TNF α production and release as well as PGE₂ synthesis by PBMC from ESRD patients. Each symbol represents the averaged value of 6 to 8 measurements per patient and study period. The horizontal bars represent the means per group. The Cuprophan data are the same as in Figures 1 to 3. Total IL-1 β , TNF α , and PGE₂ production are expressed as ng/2.5 \times 10⁶ PBMC/18 hours of incubation. IL-1 β and TNF α release are depicted as percent of total production. Comparing the two study periods, IL-1 β release increased ($P < 0.05$) and PGE₂ synthesis decreased ($P < 0.02$) whereas total IL-1 β , total TNF α and TNF α release did not differ significantly.

Mechanisms of IL-1 β secretion

Several mechanisms of IL-1 β release from LPS-stimulated PBMC have been introduced. IL-1 β is bound in the cytosol, processed by a specific enzyme and translocated to the plasma membrane. Release from the cell may involve a process which probably involves tissue plasminogen activator and serum enzymes such as plasmin [29, 30]. A role of protein kinase C (PKC) has been proposed [31] because PBMC activated by LPS in the presence of PKC inhibitors do not secrete IL-1 β , whereas total

IL-1 β production is unaffected. Furthermore, IL-1 β secretion seems to be regulated by temperature-sensitive short lived proteins, because preincubation of PBMC for 60 minutes at 42°C inhibits secretion but only reduces total production of IL-1 β [32]. In contrast to an active process of secretion, other studies favor the concept that IL-1 is released from LPS-stimulated PBMC by a non-specific pathway as a consequence of cellular injury [33]. It remains to be elucidated whether any of these mechanisms are different in PBMC from ESRD patients.

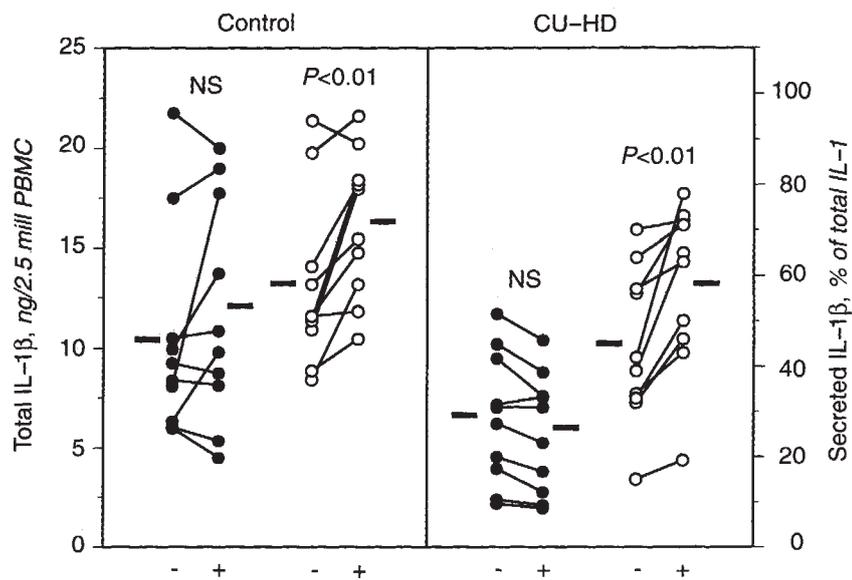


Fig. 5. The effect of indomethacin (0.5 μ g/ml) on total IL-1 β production and IL-1 β secretion by PBMC from healthy controls (Control) and from ESRD patients on Cuprophane hemodialysis (CU-HD). PBMC were stimulated with 100 ng/ml LPS. Total IL-1 β (●) is depicted on the left vertical axis as ng/2.5 \times 10⁶ PBMC/18 hours of incubation; secreted IL-1 β (○) is depicted on the right vertical axis as percent of total IL-1 β production. The symbols represent individual data of 10 controls and 10 CU-HD patients. The control data are the same as those depicted in Table 2. The horizontal bars represent the means per group. $P < 0.01$ compared to PBMC incubations without indomethacin; NS = not significant. - is no indomethacin; + is 0.5 μ g/ml indomethacin.

Table 2. Effect of indomethacin on LPS-induced IL-1 β and TNF α production and secretion

LPS ng/ml	Indomethacin 0.5 μ g/ml	Total IL-1 β ng/2.5 \times 10 ⁶ PBMC	Secreted IL-1 β % of total	Total TNF α ng/2.5 \times 10 ⁶ PBMC	Secreted TNF α % of total
0	-	<0.08	ND	0.26 \pm 0.1	ND
	+	<0.08	ND	0.32 \pm 0.1	ND
1	-	3.9 \pm 2.3	51.2 \pm 18.1	8.2 \pm 2.3	94.9 \pm 2.4
	+	3.7 \pm 2.4	46.6 \pm 21.6	8.8 \pm 2.8	96.1 \pm 0.4
10	-	11.3 \pm 8.7	53.1 \pm 17.6	10.1 \pm 2.9	94.1 \pm 6.2
	+	10.3 \pm 6.2	63.1 \pm 15.7 ^a	10.8 \pm 3.0	94.4 \pm 2.7
100	-	10.4 \pm 5.2	57.7 \pm 19.1	10.7 \pm 3.5	94.2 \pm 3.2
	+	11.8 \pm 5.6	71.1 \pm 16.2 ^b	14.6 \pm 5.0 ^a	96.0 \pm 1.2

Data are means \pm SD.

^a $P < 0.02$, ^b $P < 0.01$ compared to PBMC incubated without indomethacin

Processing of IL-1

Our data are also consistent with an impaired processing of the IL-1 β precursor molecule into the mature 17 kD IL-1 β , which is the major extracellular form of IL-1 β . Impaired processing of IL-1 β results in the accumulation of the precursor in the cytosol. As the precursor is partially recognized by the polyclonal anti-human IL-1 β antiserum used in this study, accumulation of the IL-1 β precursor molecule would result in increased cell-associated levels of IL-1 β . On the other hand, impaired processing will reduce the appearance of the mature IL-1 β molecule in the extracellular compartment. The enzyme involved in the processing of pro-IL-1 β is the interleukin-1 β converting enzyme (ICE) [34]. This IL-1 convertase is pH sensitive, is optimally active at low (5.7) external pH and is inactivated at pH \geq 8.0 [32]. One could propose from the data in this study that ICE is less active in PBMC from ESRD patients compared to healthy controls. It is possible that the increase in PGE₂ synthesis suppresses either the synthesis or activation of ICE.

Human ICE is homologous to the Nematode *c-elegans* 3 (*ced-3*) gene which is required for developmental programmed cell death

or apoptosis [35]. Cells undergoing apoptosis express the *ced-3* gene. By down-regulating of ICE activity during Cuprophane hemodialysis, impaired programmed cell death of monocytes/macrophages could result in longer survival of monocytes in the circulation. These monocytes are likely to express a cytokine profile which is different from that of young monocytes. Further studies are necessary to investigate whether these altered monocytes are present in the circulation of ESRD patients and whether these cells could account for the immune defect described in these patients.

Variability in IL-1 β and TNF α production

This study describes an intra-individual variation as high as 40 to 60% in LPS-induced production of IL-1 β and TNF α in all study groups. This variability indicates that determination of cytokine production using single point measurements is not reliable. When the release of IL-1 β from LPS stimulated PBMC was expressed as percent of total production, the coefficients of variation were reduced to the level of the assay variability. These data suggest that IL-1 β release as opposed to total IL-1 β production is a stable and reliable parameter to describe PBMC function in longitudinal studies.

Our observation that the inter-subject variation was highest in the control group is in agreement with the previously made distinction between high and low responders for LPS-induced cytokine production [9]. Studying IL-1 β release, the inter-individual variability was high in controls and uremic patients due to one individual in each group who had consistently reduced amounts of secreted IL-1 β . These data suggest that the mechanism of IL-1 β release from PBMC may be, in part, genetically determined. However, as IL-1 β secretion was less than 50% in PBMC from all ESRD patients on Cuprophane treatment, our data suggest an influence of the hemodialysis procedure on PBMC function.

Neither total TNF α production nor TNF α release was significantly different in uremic patients and hemodialysis patients compared to controls. These data indicate that neither uremia *per se* nor the hemodialysis procedure changes the ability of PBMC to synthesize and release TNF α in response to LPS. In contrast, total

LPS-induced IL-1 β production was higher in PBMC from uremic patients compared to those from healthy controls and ESRD patients. Explanations for this observation remain speculative; it is possible that uremia and/or an active kidney disease causes a preactivation of PBMC leading to a higher IL-1 β production. However, as different individuals were studied in the uremic group versus the ESRD group and because of the large intra-individual variability, the importance of differences in total IL-1 β production, although statistically significant, should be viewed with caution.

To our knowledge, our study provides the first evidence that the impaired ability to secrete IL-1 β is predominantly associated with the hemodialysis procedure itself and not by uremia *per se*. If one assumes that the reduced ability of PBMC to release IL-1 β plays a role in the impaired immune response described in hemodialysis patients [36–38], the results of the present study suggest that long-term hemodialysis therapy with synthetic high-flux membranes such as AN 69 could improve the ability of PBMC to secrete IL-1 β in ESRD patients. Whether this modified PBMC response observed *in vitro* is accompanied by an improved immune response remains to be established. However, as the functional changes of PBMC seem to be induced by the dialysis procedure, the reduced IL-1 β secretion under the conditions of long-term Cuprophane hemodialysis may just reflect an adaptation of the PBMC to protect the patient against the inflammatory effects of repeated cytokine induction during the extracorporeal procedure.

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