

Urinary monocyte chemotactic protein-1 and transforming growth factor- β in systemic lupus erythematosus

S. Torabinejad^{1,2}, R. Mardani², Z. Habibagahi³, J. Roozbeh^{1,3}, P. Khajedeji^{1,3}, M. Pakfetrat^{1,3}, M. A. Banihashemi^{1,3}, S.J. Banihashemi^{1,3}

¹Shiraz Nephrology Urology Research Center, ²Departments of Pathology and ³Medicine, Shiraz Medical School, Shiraz University of Medical Sciences, Zand Avenue, Shiraz, Iran

ABSTRACT

The purpose of this investigation was to assess the correlation of two biomarkers with the occurrence of renal flares in systemic lupus erythematosus (SLE). Urine levels of monocyte chemotactic protein-1 (MCP-1) and transforming growth factor beta (TGF- β) were measured at baseline, and at two and four months in five groups of patients: 25 lupus nephritis patients with active disease (active LN), 10 lupus nephritis patients with SLE in remission (remission LN), 25 patients with clinical active SLE and without nephritis (active NLN), 10 patients without nephritis with SLE in remission (remission NLN) and 10 healthy controls. We used repeated measurement and ANOVA with Duncan's post hoc to analyze the data; the urine level of the two proteins could distinguish the groups based on the existence of lupus nephritis and/or activity of SLE disease. Furthermore we performed receiver operating curve analysis to identify a cutoff point with a good sensitivity and specificity to diagnose lupus nephritis with either one of the urine proteins. Finally the samples from active LN were grouped according to whether they were Class IV or other classes. Baseline urinary MCP-1, but not TGF- β , was significantly different between the classes. Further investigation into the use of these cytokines in a prospective study is needed to determine their capacity as diagnostic tools for renal flares.

Key words: Lupus nephritis, monocyte chemotactic protein-1, systemic lupus erythematosus, transforming growth factor beta

Introduction

Systemic lupus erythematosus (SLE) is a chronic inflammatory disease of unknown cause. It has a reported prevalence of 20 to 150 cases per 100,000 in the US population.^[1-3] Renal involvement as shown by abnormal urinalysis and/or renal function is present in as many as 75% cases at some point.^[4] In some, the symptoms may not be evident early in the course of disease,^[5-7] therefore clinical findings underestimate the true prevalence of renal involvement. Within the first one to two years after diagnosis of SLE, renal abnormalities appear with

nephropathy occurring in 30% in the next decade.^[1,8-12] A large portion of these patients will progress to end-stage renal disease.^[13]

Renal biopsies have remained the “gold standard” of assessing lupus nephritis (LN) patients not only at diagnosis but also to assess the efficacy of treatment. But this may not always be feasible due to the invasive nature of the procedure. That is why replacements have been sought in the plasma and urine. The ability of these markers to predict the “gold standard”, such as renal biopsy scores or longer-term outcome, is imperative to make them reliable.^[14]

Two promising candidates were taken into account for this study: monocyte chemotactic protein 1 (MCP-1), also known as chemokine CCL2, and transforming growth factor beta (TGF- β). They have been shown to correlate with disease activity in SLE and renal flare associated with the disease, and a role in its pathogenesis has been proposed.^[14-26] We assessed the urine levels of these two proteins and compared them to findings on renal biopsies where available. The aim was to assess the levels of these two cytokines in these five different groups.

Address for correspondence:

Dr. Simin Torabinejad, Shiraz Nephrology Urology Research Center, Shiraz Medical School, Shiraz University of Medical Sciences, Zand Avenue, Shiraz, Iran. E-mail: torabins@yahoo.com

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Materials and Methods

This study was done on five populations, composed of 25 LN patients with active disease (active LN), 10 LN patients with disease in remission (remission LN), 25 patients with clinical active disease but without nephritis (active NLN), 10 patients without nephritis with disease in remission (remission NLN) and 10 healthy controls. Patients with co-morbid conditions such as diabetes mellitus, hypertension, or those on antibiotic therapy at the time were excluded from the study groups. Lupus nephritis and/or active SLE received treatment.

Renal flare in LN was defined as ≥ 300 mg protein in a 24-h urine collection and/or a 25% increase in serum creatinine.^[27,28] Patients without LN in the past two years were considered NLN.

Diagnosis of SLE was based on having four or more criteria according to the American College of Rheumatology criteria for SLE.^[29] Active disease was based on having a score of 6 or more according to the systemic lupus erythematosus disease activity index (SLEDAI).^[15] Those without manifestations of activity in the past six months were considered in remission.

Biopsy samples were placed in 10% formalin and were later dehydrated and embedded in paraffin in an automated machine (Autotechnicon Duo[®] MOD. 2A, Technicon Corporation, USA); 5- μ m-thick sections were cut using a microtome (Leitz 1512 Microtome (WS-LEITZ1512), Ernst Leitz Wetzlar GmbH, Germany).

Twenty-two patients with LN and active disease and nine with LN in remission had biopsy specimens for this study. Classification was done according to the criteria from ISN/RPS.^[17-19] The histological activity and chronicity index of each biopsy sample was calculated after preparation.^[30] The histological activity index was the sum of semiquantitative manual scores (each ranging from 0-3) for the following parameters: endocapillary hypercellularity, leukocyte infiltration, subendothelial hyaline deposits, interstitial inflammation, necrosis, and cellular crescents. Scores of the last two parameters were counted double, therefore the activity index ranged from 0-24. The chronicity index was the sum of semiquantitative scores (each ranging 0-3) for the following parameters: Glomerular sclerosis, fibrous crescents, tubular atrophy, and interstitial fibrosis. The score ranged from 0-12. All biopsy specimens were scored and evaluated by one board-certified pathologist. Biopsies were performed as part of their standard clinical indications in patients suspected of having renal involvement related to SLE.

Patients were asked to provide freshly voided, whole-stream, early-morning urine samples three times, at baseline and later at two-month intervals on each visit. The urine samples were provided in the lab and were soon centrifuged to remove supernatants and frozen in small aliquots without further manipulation at -80°C for chemokine analysis at the end of the study. It has been shown that the freezing procedure protects the samples from deterioration and they can be used after thawing.^[14,27] Urine samples were cultured to exclude the possibility of infection.

All patients with LN received pulse methylprednisolone 500-1000 mg/day for three consecutive days followed by one dose of cyclophosphamide 500-1000 mg on the fourth day. From the fifth day onwards therapy was changed to prednisolone 1 mg/kg/day for two months and was gradually tapered throughout one month and later continued at 5-10 mg/day. Cyclophosphamide was administered 500-1000 mg IV once per month for six to eight months, which was later changed to every three months for at least one year.

Both groups with SLE activity regardless of their kidney involvement received maintenance steroid therapy, hydroxychloroquine sulfate and calcium D supplement.

Urine MCP-1 (uMCP-1) was measured using the human MCP-1 kit (BMS281, manufactured by Bender MedSystems GmbH[®], Vienna, Austria, Europe). Limit of detection for the kit was 2.31 pg/ml. Urine TGF- β 2 (uTGF- β 2) protein was measured using human TGF- β 2 Elisa kit (BMS254, manufactured by Bender MedSystems GmbH[®], Vienna, Austria, Europe). Limit of detection for the kit was 6.6 pg/ml. Both cytokines were measured by specific enzyme-linked immunosorbent assay (ELISA) according to the manufacturer's instructions. This assay employed the quantitative sandwich enzyme immunoassay technique. Results were reported as pg/ml. According to the manufacturer's guide samples would remain stable when frozen at -20°C and then thawed for analysis.

Statistical analysis was performed using SPSS for Windows, version 15.0 (SPSS software for Windows, Chicago, IL, USA). The results are presented as mean \pm standard deviation (SD) unless specified otherwise. Urine MCP-1 and uTGF- β 2 were compared between groups and at different time intervals by repeated measurement. They were also compared between groups at baseline by one-way ANOVA and Duncan's post hoc to determine difference of means between groups. Pearson's correlation between activity index and uMCP-1 and uTGF- β 2 was used. We used the receiver operating characteristic (ROC) curve to determine the best value

of uMCP-1 and uTGF-β2 with the highest sensitivity and specificity to differentiate patients with LN from those without LN. *P* values less than 0.05 were considered significant. All probabilities were two-tailed.

Results

We studied 70 SLE patients in total. The baseline demographic and characteristics of these groups have been provided in Table 1. Among the two groups with LN (active LN and remission LN) 31 underwent kidney biopsy (22 and nine respectively). Groups did not differ regarding age (*P* = 0.585).

The maximum mean level of both uMCP-1 and uTGF-β2 was in the active LN group, while the minimum belonged to the control group. The mean levels of these substances in the urine at baseline were significantly different in the five groups [Table 1]. The five groups of patients were classified according to similarity in their mean uMCP-1 and uTGF-β2 [Table 2].

The groups were arranged into three subsets according to their uMCP-1 values. Subset 1 were those with LN regardless of their SLE activity; they had the highest amount of uMCP-1 in their urine compared to other groups, but they did not differ from each other. Subset 2 was those without nephritis regardless of disease activity. The mean value of uMCP-1 in the first subset (572.54 pg/ml) was more than twice that of the mean value of the second subset (276.76 pg/ml). The control group formed a third subset which showed a similar value to the Active NLN group. This caused an overlap between the last two subsets.

In assessing the uTGF-β2 level three subsets appeared again. The first subset was the Active LN group, which had the highest level of uTGF-β2. The second subset consisted of those with SLE in remission regardless of their nephritis. This group overlapped with the third subset which consisted of patients without nephritis regardless of the clinical activity of SLE and healthy controls. The mean value of uTGF-β2 in the first subset (86.52 pg/ml) was more than twice that of the combined mean value of the second subset (42.17 pg/ml).

Table 1: Demographics and characteristics of the study groups*

Group	Active LN [†]	Remission LN [†]	Remission NLN	Active NLN [†]	Controls
No. of subjects	25	10	10	25	10
Age (years)	30.1 ± 6.7	33.9 ± 9.4	31.0 ± 6.2	30.1 ± 6.7	31.5 ± 1.4
Gender (F/M)	24/1	10/0	7/2	24/1	8/2
Chronicity index	8.7 ± 7.0	7.1 ± 4.2	-	-	-
Activity index	1.4 ± 1.4	2.3 ± 2.5	-	-	-
Class of LN					
I	0	0	-	-	-
II	4	1	-	-	-
III	5	2	-	-	-
IV	10	5	-	-	-
V	3	1	-	-	-
VI	0	0	-	-	-
uMCP-1					
Baseline	592.9 ± 319.0	521.6 ± 118.9	319.0 ± 183.2	171.2 ± 71.1	138.8 ± 42.6
(range)	(273 - 1360)	(322 - 763)	(85 - 789)	(78 - 278)	(85 - 221)
95% CI	461.3 - 724.6	436.5 - 606.6	243.4 - 394.9	120.4 - 222.0	108.3 - 169.2
2 months	424.5 ± 197.6	403.7 ± 117.6	251.9 ± 131.8	151.2 ± 81.2	-
(range)	(171 - 1000)	(200 - 620)	(80 - 524)	(65 - 272)	-
95% CI	343.0 - 506.1	319.6 - 487.8	197.5 - 306.3	93.0 - 209.3	-
4 months	417.7 ± 194.0	535.8 ± 99.1	210.0 ± 140.3	102.1 ± 58.1	-
(range)	(200 - 1100)	(185 - 462)	(51 - 640)	(50 - 245)	-
95% CI	337.7 - 497.8	282.9 - 424.65	152.1 - 267.9	60.6 - 143.7	-
uTGF-β					
Baseline	86.5 ± 32.5	48.7 ± 15.6	39.6 ± 16.2	26.3 ± 7.7	25.4 ± 10.2
(range)	(35 - 153)	(28 - 82)	(8 - 82)	(15 - 38)	(10 - 41)
95% CI	73.117 - 99.915	37.509 - 59.851	32.883 - 46.245	20.789 - 31.771	18.061 - 32.699
2 months	76.8 ± 31.1	41.1 ± 14.2	34.7 ± 14.3	25.5 ± 6.3	-
(range)	(30 - 140)	(22 - 73)	(12 - 70)	(15 - 35)	-
95% CI	64.0 - 89.6	30.9 - 51.3	28.8 - 40.6	20.9 - 30.0	-
4 months	62.4 ± 25.0	40.8 ± 8.9	32.5 ± 16.0	20.4 ± 5.9	-
(range)	(25 - 145)	(30 - 62)	(10 - 68)	(12 - 30)	-
95% CI	52.1 - 72.7	34.5 - 47.2	25.9 - 39.1	16.1 - 24.6	-

*Values are mean ± SD, except where indicated otherwise; [†]Indicates groups receiving treatment; From the Active LN group only 22 and from the Remission LN group only nine had biopsy specimens available; CI = Confidence interval; LN = Lupus nephritis; uMCP-1 = Urine levels of monocyte chemotactic protein-1; uTGF-β = Urine level of transforming growth factor beta; NLN = No lupus nephritis

We combined the groups Active LN and Remission LN into a group 35-strong and assessed the internal correlation between uMCP-1 and uTGF-β2. Urine MCP-1 at four months correlated positively with uTGF-β2 at all time intervals (baseline: $r = 0.458, P = 0.006$; 2 mo: $r = 0.395, P = 0.019$; 4 mo: $r = 0.483, P = 0.003$). Urine TGF-β2 at four months also correlated positively with uMCP-1 at all time intervals (baseline: $r = 0.443, P = 0.008$; 2 mo: $r = 0.418, P = 0.12$; 4mo: $r = 0.483, P = 0.003$). The only time when both correlated with each other in the same cross-sectional time interval was at four months after treatment. In the rest of time intervals there was no significant correlation between the two cytokine levels at corresponding months in the urine.

Using receptor operating curve analysis, we calculated a cutoff point with maximum sensitivity and specificity for urine levels of cytokines to discriminate between patients with LN and those without it regardless of the activity of SLE disease. Urine MCP-1 [Figure 1] had the higher sensitivity and uTGF-β2 [Figure 2] had the higher specificity. We recalculated these values excluding the control group, the results appeared similar (data not shown).

Both urine cytokines at baseline correlated positively with the class of LN (uMCP-1: $r = 0.454, P = 0.010$; uTGF-β2: $r = 0.421, P = 0.018$). They also correlated positively with the histological activity index (uMCP-1: $r = 0.409, P = 0.022$; uTGF-β2: $r = 0.371, P = 0.040$) at baseline. They did not, however, correlate with the chronicity index. Neither histological index correlated with the class of LN. The two indices also did not correlate with each other. Specimens with a histological activity score of >12 had a significantly higher level of uMCP-1 and uTGF-β2 at baseline than those ≤ 12 (uMCP-1: 795.100 ± 392.4 vs. 470.5 ± 148.6 , respectively, $P = 0.029$; uTGF-β: 96.0 ± 27.4 vs. 64.2 ± 30.7 , respectively, $P = 0.009$) [Figures 3 and 4]. Only 31 patients were included as only 22 patients from the active LN group and nine from the remission LN group had biopsies available for the aforementioned analyses and only baseline values of urine cytokines were used.

We categorized patients with LN in groups according to whether they were Class IV or not. Their characteristics have been outlined and compared in Table 3.

Figures 5 and 6 show the trend of decrease of uMCP-1 and

Table 2: Mean urine levels of cytokines in each group compared at baseline

Groups	Number	uMCP-1			uTGF-β2		
		1	2	3	1	2	3
Active LN	25	592.940			86.516		
Remission LN	10	521.550				48.680	
Remission NLN	25		318.980			39.564	39.564
Active NLN	10		171.200	171.200			26.280
Healthy controls	10			138.750			25.380
<i>P</i> value*		0.395	0.081	0.699	1.000	0.284	0.117

Groups are divided into three subsets, each with some overlap according to their urine cytokine levels. **P* values represent the difference observed between values in one subset. Combined mean value of each subset differed from other subsets for each urine cytokine ($P < 0.05$); LN = Lupus nephritis; uMCP-1 = Urine levels of monocyte chemotactic protein-1; uTGF-β = Urine level of transforming growth factor beta; NLN = No lupus nephritis

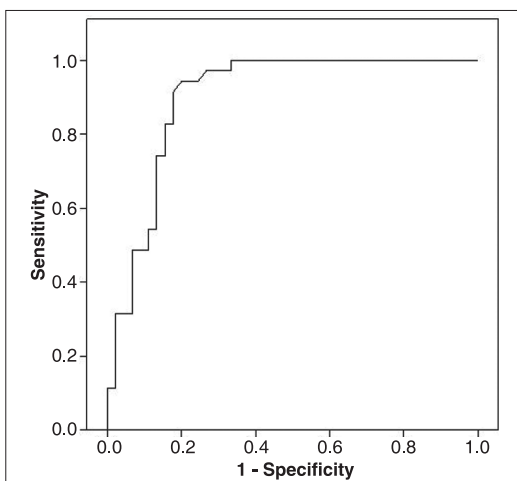


Figure 1: Receiver operating characteristic curve for baseline uMCP-1. Cutoff point to diagnose LN regardless of SLE activity was 304.5 pg/ml with a sensitivity of 94.3 (95% CI 80.8%-99.1%), specificity of 80% (95% CI 65.4%-90.4%), and AUC = 0.900

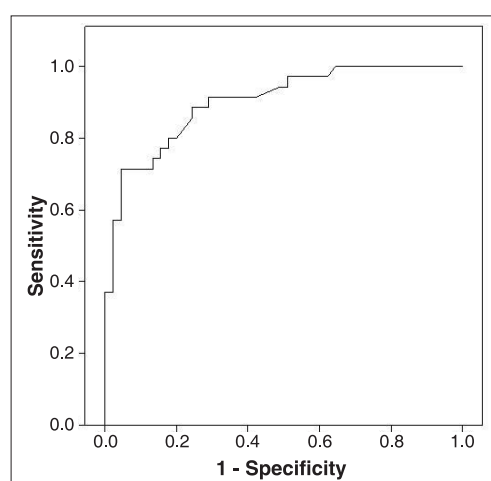


Figure 2: Receiver operating characteristic curve for baseline uTGF-β2. Cutoff point to diagnose LN regardless of SLE activity was 54.2 pg/ml with a sensitivity of 71.4% (95% CI 53.7%-85.3%), specificity 95.6% (95% CI 84.8%-99.3%), and AUC = 0.902

uTGF-β2 in patients in the four groups, at two months' intervals (healthy controls were not included in this analysis as only baseline values were available). When we took time and grouping into consideration simultaneously, the trend of decline in uMCP-1 amongst the four groups

Table 3: Patient characteristics according to their lupus class*

Group	Class IV	Other classes	P value
No. of subjects	15	19	
Age (range)	31.7 ± 7.6 (22-46)	30.6 ± 8.0 (19-52)	0.570
Gender (F/M)	15/0	18/1	0.999
Chronicity index (range)	2.5 ± 1.9 (0-8)	0.9 ± 1.3 (0-7)	0.008
Activity index (range)	13.5 ± 4.6 (3-20)	3.3 ± 2.3 (0-4)	<0.001
uMCP-1			
Baseline (range)	689.7 ± 358.0 (273.5-1360.5)	486.5 ± 158.2 (300.0-820.0)	0.034
95% CI	(491.4-887.9)	(410.3-562.8)	
uTGF-β			
Baseline (range)	82.9 ± 31.1 (32.1-144.0)	71.8 ± 34.9 (28.0-153.0)	0.340
95% CI	(65.7-100.1)	(54.9-88.6)	

*Values are mean ± SD, except where indicated otherwise; CI = Confidence interval; uMCP-1 = Urine levels of monocyte chemotactic protein-1; uTGF-β = Urine level of transforming growth factor beta

was similar ($P = 0.111$), but the trend of decline in uTGF-β2 in the groups was different ($P = 0.001$). This was because uTGF-β2 in the active NLN group from baseline to two months decreased only $0.2 \pm 22.3\%$ while the decline in the other three groups was $11.5 \pm 7.8\%$, $15.6 \pm 10\%$ and $5.9 \pm 34.0\%$ respectively, for active LN, remission LN and Remission NLN. From two months to the end of the study the groups active LN and active NLN decreased $16.3 \pm 19.3\%$ and $18.0 \pm 23.2\%$ respectively; this was in contrast to the decline observed in remission NLN at only $6.3 \pm 28.8\%$ and the rise of uTGF-β2 by $5.4 \pm 28.7\%$ in the remission LN group.

When groups with LN were made into two groups according to their class (Class IV versus Other classes) and their response to treatment was assessed, the results showed that the trend of decline was not significantly different amongst them (data not shown).

Discussion

Both MCP-1 and TGF-β have been proposed as candidates in the pathogenesis of renal damage in inflammatory states

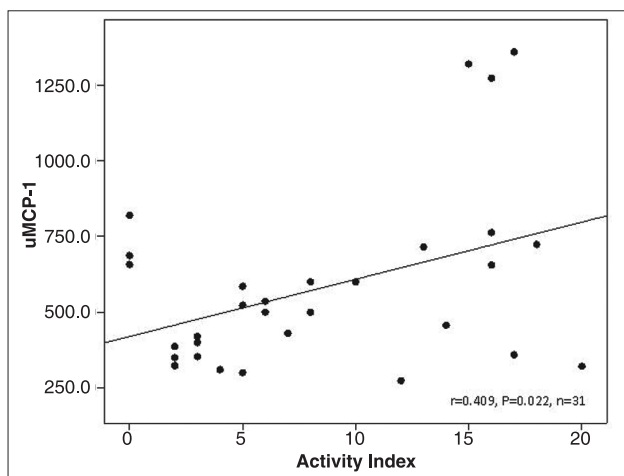


Figure 3: Correlation between baseline uMCP-1 (pg/ml) and histological activity index on the kidney biopsy specimens of patients with LN

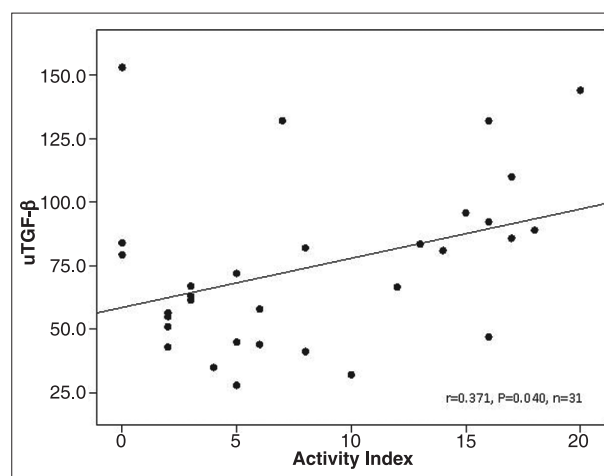


Figure 4: Correlation between baseline uTGF-β2 (pg/ml) and histological activity index on the kidney biopsy specimens of patients with LN

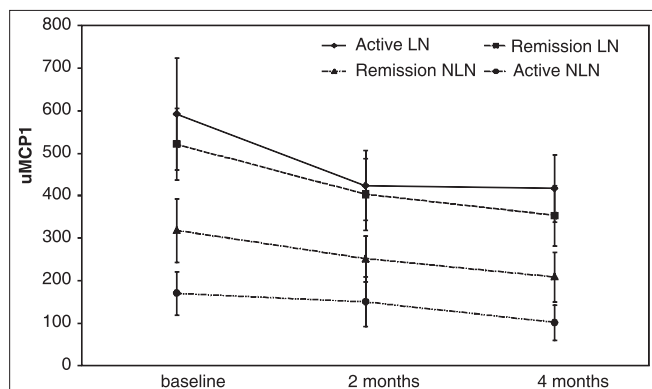


Figure 5: Trend of decrease in uMCP-1 levels from baseline to four months

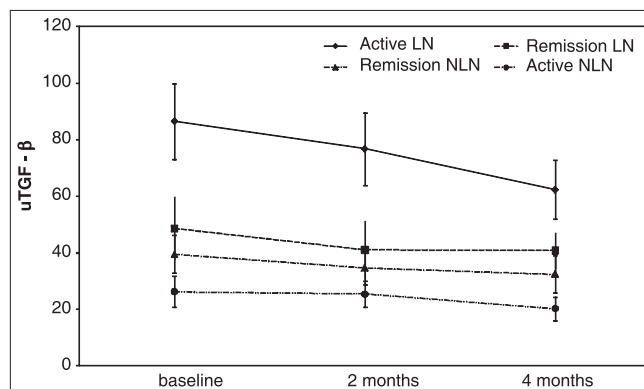


Figure 6: Trend of uTGF-β2 level from baseline to four months

such as SLE.^[16] In a previous study, mRNA expression of MCP-1 and TGF- β in urinary sediments of patients with LN decreased with treatment, correlating with clinical activity of the disease and SLEDAI score and histological activity index. The protein product level of MCP-1, but not TGF- β , correlated with the SLEDAI score but neither correlates with the histological activity index.^[23]

Transforming growth factor beta (TGF- β) levels in the plasma and urine correlate with SLEDAI^[14] in children. Decreased plasma levels and increased urinary excretion of TGF- β has been shown in children with active LN.^[21] The glomerular or tubular gene expression of TGF- β has no correlation with serological markers of lupus activity in the kidney; however, its tubulointerstitial expression may be indicative of chronic renal damage as it correlates with the chronicity index.^[20]

Monocyte chemotactic protein-1 (MCP-1) has been shown to increase in gene expression and urine excretion in patients with LN and those with LN flare. It has proven to be sensitive and specific for renal flare in contrast to other renal diseases. It remains a valid biomarker even for patients on maintenance immunosuppression. The amount found in urine does not correlate with the plasma level, leading to the conclusion that it is local production in the kidney, not simply filtration, that increases the urine levels in SLE flare. It is thought to be involved in the pathogenesis of LN by mediating leukocyte infiltration.^[16,22,26,27] Glomerular expression of MCP-1 correlates with serological markers of lupus activity such as C3, C4 as well as with the activity index on histological examination^[17-19] while the tubular expression correlates with the chronicity index and the percentage of cortical fibrosis.^[20] Urine MCP-1 was associated with LN in an experimental model.^[31]

By dividing the groups according to their urine cytokine levels we attempted to see if these could discriminate between the different groups. Interestingly uMCP-1 values could discriminate between groups according to whether they had LN or not regardless of their SLE activity, while uTGF- β 2 values only usefully discriminated between groups with LN according to their SLE activity [Table 2]. Just as intrarenal gene expressions of these cytokines are indicative of the disease involving the kidneys and not systemic activity,^[20] urine protein levels seem to have the same discriminating ability.

We also tried a cutoff point for both cytokines which had a very good sensitivity, specificity and area under curve (AUC) for uMCP-1 level to separate patients with and without LN. For uTGF- β 2 the test was very specific but lacked a satisfactory sensitivity. This is important in achieving the goal of excluding renal biopsies for select

patients while being able to identify those who are highly likely to have renal involvement in flare episodes.

Both cytokines correlated positively and significantly with the class of LN and the histological activity index. This also shows that they are indicative of how advanced the disease is and since they did not correlate with the histological chronicity index it can be assumed that they are indicators of acute involvement rather than long-term damage as seen on the renal biopsy specimens. There is a significant correlation between the glomerular expression of MCP-1 and histological activity and its tubulointerstitial expression and the chronicity index.^[20] Our findings are in line with the current notion that MCP-1 and TGF- β are involved in the pathogenesis of renal involvement in SLE.^[20,32,33] A previous study had observed a similar trend for urine levels of mRNA of these cytokines but urine protein levels did not correlate with either histological index.^[23] Their limit of detection for uMCP-1 and uTGF- β was $\times 2$ and $\times 2.3$ times less sensitive than that of ours respectively. This could also explain why they could not detect uTGF- β in some samples. Our results show that the urine level of these cytokines can be a useful marker and does correlate with indices related to LN. Also, the method utilized is less expensive and laborious in comparison to real-time polymerase chain reaction (PCR). In another study uMCP-1 was shown to be significantly associated with manifestations of severe renal injury^[34] and it was also shown that if patients develop active SLE while on immunosuppressive therapy, rise of this marker can be used as an indicator of LN as it correlates with local kidney involvement and not systemic increase in SLE activity. It was also a sensitive and specific biomarker of renal flare and the severity of renal injury in SLE. More interestingly this rise was observed a few months in advance suggesting a pathogenic and predictive role.

We also observed a correlation between the cytokine levels themselves. Only at the end of the study did both correlate positively at the same cross-section. We could not find an explanation for the observed significant correlations of uMCP-1 levels at four months with uTGF- β 2 levels observed at baseline and two months, and vice versa. Whether this shows that these cytokine expressions affect the production and subsequent rise in urine of the other can only be speculated. Another study, which had included the mRNA levels in the urine of both cytokines, had however found a significant internal correlation between the two.^[20]

Urine levels of both cytokines decreased in response to treatment. Urine TGF- β 2 levels steadily declined over the course of four months and were substantially

different from other groups at all time intervals. Urine MCP-1 levels had a sharper decrease in the first two months and could differentiate the two groups with LN (Active LN and Remission LN) from the rest, but not from each other.

Finally, Class IV of LN showed a distinctively higher chronicity and activity as well as baseline uMCP-1, but not uTGF- β . As Class IV is known to have a worse prognosis, the difference in the histological indices was not a surprise. Activity index differed more prominently indicating that Class IV cases experience more inflammation during flares. As to why uTGF- β at baseline did not differ, an explanation could not be offered.

A number of other biomarkers have been studied with regard to renal SLE flares. Increased plasma and urine levels of adiponectin have been detected in patients with SLE, particularly in those with renal flare. Interestingly, urinary levels of adiponectin, particularly the high molecular weight isoform, rise in renal SLE flare and not nonrenal SLE flare.^[35,36] Within a four-month period before and after renal flare, urine concentrations of hepcidin 20 and 25 correlated with the occurrence of a flare.^[37] Neutrophil gelatinase-associated lipocalin also increases in urine and plasma, and can predict the worsening of renal childhood-onset SLE up to three months prior to the exacerbation.^[38]

Our study had a number of shortcomings. Firstly, we used a spot urine sample for determining the protein levels; ideally we should have included urine creatinine to create a protein-to-creatinine ratio to adjust the values for fluctuating levels of urine cytokines. Secondly, a more populous study should be done to be able to focus on different aspects, such as different classes of LN and to increase the certainty of cutoff points mentioned in this study. Lastly, we lacked a follow-up on our control group at two and four months which could have provided a better comparison with the trend of decline in the other groups.

Conclusion

This study shows that MCP-1 and TGF- β increase in renal flare of SLE. Their capacity to be used as a diagnostic tool to identify a renal flare in patients should be assessed in a prospective longitudinal follow-up starting from a few months before the occurrence of flare.

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