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ORIGINAL RESEARCH

Association of Mitochondrial Pyruvate Carrier with the Clinical and Histological Features in Lupus Nephritis

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Background: Mounting evidence suggests that mitochondrial dysfunction contributes to lupus nephritis (LN) pathogenesis. Mitochondrial pyruvate carrier 1 (MPC1) and mitochondrial pyruvate carrier 2 (MPC2) mediating pyruvate transport from the cytoplasm to the mitochondrial matrix, determines the cell survival and cellular energy supply. Here, we aimed to investigate the association of mitochondrial pyruvate carrier expression with the clinical and histological features in LN.

Methods: Patients with biopsy-proven proliferative LN (class III and class IV, n=18) and membranous LN (class V, n=18) were included. Expression of MPC1 and MPC2 were examined by immunohistochemistry. MPC protein levels in the two groups were evaluated by the Student's *t*-test. Correlation analysis between MPC levels and clinicopathological features was performed by Spearman's rank correlation.

Results: Both MPC1 and MPC2 were exclusively expressed in renal tubules of enrolled LN. Significantly lower MPC1 and MPC2 were observed in patients with proliferative LN compared to membranous LN. In addition, the MPC1 and MPC2 were negatively correlated with SLEDAI-2K score, renal function, and renal pathology activity index.

Conclusion: Both MPC1 and MPC2 were localized in renal tubules, and decreased MPC content was more pronounced in proliferative LN than membranous LN. MPC levels were significantly correlated with renal functions and renal pathology activity. **Keywords:** lupus nephritis, mitochondrial pyruvate carrier, proliferative, membranous, tubulointerstitial lesions

Introduction

Lupus nephritis (LN) is one of the most common severe manifestations of systemic lupus erythematosus (SLE) and a key driver of mortality and morbidity in SLE.^{1–3} Numerous studies demonstrated that the proportion of patients with SLE developed lupus nephritis in their disease course, the prevalence was approximately 40%.⁴ Despite increased knowledge of disease pathogenesis and improved treatment options, a significant portion of patients with severe LN (10–30%) still progress to end-stage renal disease (ESRD) within 15 years even with aggressive immunosuppressive therapy.^{5,6} It is imperative to identify sensitive and precise indicators for reflecting disease activity and stratifying LN patients at risk of progressing to rapid renal impairment.

The role of mitochondria in lupus has attracted extensive attention, and functional alternation of mitochondria would take part in systemic lupus erythematous.^{3,7,8} Mitochondria is the main endogenous source of reactive oxygen species (ROS), therefore, plays a crucial role in generating oxidative stress. ROS modulates cytokine production, T cell activation and inflicts DNA damage, which facilitates the emergence of SLE.⁹ Moreover, mitochondria DNA damage may cause ATP reduction, ROS generation, and tissue damage in systemic lupus erythematous.¹⁰ Mitochondrial dysfunction-provoked excessive oxidative stress is a crucial downstream contributory factor for lupus pathogenesis and the dysregulation of upstream genetic/epigenetic functions.⁷ Pyruvate, a hub metabolite for glucose, lipid, and amino acid, is critical for maintaining the stability of mitochondrial function.^{11,12} Pyruvate drives ATP production by intersecting the citric acid cycle, while ROS levels can dramatically increase when the respiratory chain is dysfunctional, and pyruvate promotes oxidative stress resistance through hormetic ROS signaling.¹³ Mitochondrial pyruvate carrier (MPC), is located in the inner mitochondrial membrane, mediating the transport of pyruvate from the cytoplasm to mitochondria.^{14,15} MPC comprises two subunits, MPC1 and MPC2, and the expression of each is essential for mitochondrial pyruvate transport.^{7,16} Our previous study explored the expression of MPC in diabetic nephropathy (DN), primary glomerulone-phropathy (PGN), PGN with diabetes mellitus (PGN-DM), and their correlation with clinical features.¹⁷ However, the role of MPC in the kidneys of lupus nephritis patients was still unclear. Therefore, the present study was aimed to investigate the significance of MPC for clinical and pathological features in proliferative LN and membranous LN.

Methods

Subjects

A total of 36 patients with renal biopsy-proven lupus nephritis were recruited from January 2016 to October 2019 were retrospectively reviewed. All patients fulfilled the 2012 SLE international collaborating clinics classification criteria.¹⁸ Renal biopsies were performed following the indications recommended by the ACR,¹⁹ and LN pathological classification was based on 2003 International Society of Nephrology/Renal Pathology Society (ISN/RPS) pathological classification criteria, modified in 2018.^{20,21} The exclusion criteria were as follows: (1) patients aged at 18 and younger; (2) patients with mixed proliferative LN (class III+V or IV+V); (3) coexistence of other renal diseases (such as minimal change disease and thrombotic microangiopathy) or diabetes. Based on their renal biopsy reports, the patients were categorized into the proliferative LN (class III, class IV) and membranous LN (class V) groups.

Measurements

Clinical parameters and laboratory data were obtained from the medical records, including gender, age, SLE duration, blood pressure, serum creatinine, blood urea nitrogen (BUN), serum cystatin C, uric acid, albumin, serum complement 3 (C3), serum complement 4 (C4), antinuclear antibodies (ANA), anti-double-stranded DNA (anti-dsDNA), 24-h urine protein, and urinary acidification function. Plasma samples and 24-h urine samples were obtained from patients within three days before the renal biopsy. GFR was estimated using the chronic kidney disease epidemiology collaboration (CKD-EPI) equation.²² The SLE disease activity index 2000 (SLEDAI-2K) and renal SLE disease activity index (rSLEDAI) were used to assess SLE activity and kidney disease activity, respectively.^{23,24} The rSLEDAI consists of four parameters: hematuria, proteinuria, pyuria, and urinary casts, each accounting for four scores. The evaluation of SLEDAI-2K scores and rSLEDAI scores were obtained one day before renal biopsy.

Immunohistochemistry

Paraffin sections were deparaffinized and rehydrated. Then, the sections were subject to antigen retrieval with citrate buffer and washed with phosphate-buffered saline. After blocking by 5% bovine serum albumin for 1 h at room temperature, the sections were incubated with MPC1 antibody (1:900, Sigma, HPA045119) and MPC2 antibody (1:20, Sigma, HPA056091) overnight at 4 °C. The sections were washed with PBS and then incubated with the second antibody for 1 h at 37 °C. The sections were then stained with 3, 3'-diaminobenzidine tetrahydrochloride for 10 min and counterstained with hematoxylin, dehydrated and mounted. Immunohistochemistry was performed on tissue sections from every subject for each antibody, which was performed with positive and negative control. And immunohistochemical parameters were assessed through integrated optical density (IOD), which means integral calculus of the stained area times the intensity of stain in each pixel in the area, indicating the total amount of staining material in that area. For each specimen, 8–10 images of stained sections were randomly selected and captured, and the Image-Pro Plus 6.0 mage analysis software (media controls, Silver Spring, MD, USA) was used to measure the IOD value. Under the same parameter conditions, the IOD value of all the slices was analyzed and presented.

Renal Pathology

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Renal biopsy specimens were evaluated by two experienced pathologists following the modified ISN/RPS pathological classification criteria.²¹ Activity indices consisted of endocapillary hypercellularity, neutrophils/karyorrhexis, fibrinoid

necrosis, hyaline deposits, cellular/fibrocellular crescent and interstitial inflammation. Chronicity indices consisted of total glomerulosclerosis score, fibrous crescents, tubular atrophy, and interstitial fibrosis. And glomerular sclerosis, cellular crescents, and fibrous crescents were calculated as percentages of the total number of glomeruli. The fibrinoid necrosis and cellular crescents were weighted by a factor of 2.

Statistical Analysis

SPSS 22.0 software (SPSS, Chicago, IL, USA) was used for data analysis. Variables were expressed as the mean \pm standard deviation, median (interquartile range), or percentage. The non-significant values in Shapiro–Wilk and Levene's tests indicate normal distribution and homogeneity in variance, respectively. Comparisons for normal distribution groups were performed using Student's *t*-test. Differences among non-normal distribution groups were analyzed using the Mann–Whitney *U*-test. Correlations between MPC protein expression levels and clinical characteristics were assessed by using Spearman's rank test. Significance was defined as p < 0.05.

Results

Baseline Characteristics of the Enrolled Patients with LN

A total of 36 patients with biopsy-proven LN were included in the study. The clinical characteristics, laboratory parameters, and pathological features of patients were shown in Table 1. The patients were predominantly women (86.1%), with a mean age of 39.56 ± 13.39 years. The mean duration of SLE was 2.00 (1.00,4.50) months. 36.10% of patients presented with nephrotic syndrome and 41.70% of patients had hematuria. The mean levels of eGFR, urine protein, serum creatinine and C3 were 90.79 ± 35.72 mL/min/1.73 m², 2.08 (1.90, 3.71) g/24 h, 69.60 (54.00, 133.55) umol/L and 0.70 ± 0.35 g/L, respectively. Among the patients, 18 patients (50%) had proliferative LN (class III, 6 patients; class IV, 12 patients), and 18 patients (50%) had membranous LN.

The comparison of characteristics between proliferative LN and membranous LN were shown in Table 1. Age, gender distribution, SLE duration, and the proportion of patients with hypertension, nephrotic syndrome or hematuria did not

	Total (n=36)	Proliferative	Membranous	P value
		LN (n=18)	LN (n=18)	
Clinical features				
Female, n (%)	31 (86.10)	15 (83.33)	16 (88.9)	0.635
Age (years)	39.56±13.39	38.89±15.19	40.22±11.72	0.770
SLE duration (months)	2.00 (1.00, 4.50)	2.50 (1.00, 6.00)	2.50 (1.00, 4.00)	0.518
Hypertension, n (%)	10 (27.80)	6 (33.33)	4 (22.22)	0.463
SBP (mmHg)	130.89±17.21	128.33±15.86	133.44±18.55	0.381
DBP (mmHg)	84.42±10.85	83.17±11.05	85.67±10.81	0.497
Nephrotic syndrome, n (%)	13 (36.10)	4 (22.22)	9 (50.00)	0.087
Hematuria, n(%)	15 (41.70)	9 (50.00)	6 (33.33)	0.310
SLEDAI-2K	10.72±4.98	12.83±5.54	8.61±3.31	0.009
rSLEDAI	8 (4.00, 8.00)	8.00 (4.00, 8.00)	4.00 (4.00, 8.00)	0.098
Cutaneous lupus, n (%)	15 (41.67)	8 (44.44)	7 (38.89)	0.739
Nonscarring alopecia, n (%)	3 (8.33)	3 (16.67)	0 (0.00)	0.074
Oral/nasal ulcers, n (%)	I (2.78)	0 (0.00)	l (5.56)	0.317
Arthritis, n (%)	12 (33.30)	7 (38.89)	5 (27.78)	0.486
Serositis, n (%)	0 (0.00)	0 (0.00)	0 (0.00)	0.999
Neurological disorder, n (%)	0 (0.00)	0 (0.00)	0 (0.00)	0.999
Hemolytic anemia, n (%)	26 (72.22)	16 (88.89)	10 (55.56)	0.028
Lymphopenia/leukopenia, n (%)	11 (30.56)	8 (44.44)	3 (16.67)	0.074
Thrombocytopenia, n (%)	6 (16.67)	3 (16.67)	3 (16.67)	0.999

Table I Comparison of the Clinical and Histologic Features Between the Patients with Proliferative LN and Membranous LN

(Continued)

Table I (Continued).

	Total (n=36)	Proliferative	Membranous	P value
		LN (n=18)	LN (n=18)	
Laboratory assessment				
WBC (10^9/L)	5.32±2.71	4.54±2.14	6.09±3.05	0.087
Hemoglobin (g/L)	110.47±19.82	101.78±18.63	119.17±17.37	0.007
Platelet (10 ⁹ /L)	185.69±60.87	184.50±53.37	186.89±69.11	0.908
eGFR (mL/min/1.73 m ²)	90.79±35.72	77.58±45.32	104.01±14.26	0.028
Serum cystatin C (mg/L)	1.74±1.11	2.05±1.23	1.11±0.38	0.172
Serum creatinine (µmol/L)	69.60 (54.00, 133.55)	90.35 (56.00, 166.20)	64.50 (51.30, 73.60)	0.009
BUN (mmol/L)	6.16 (4.12, 10.70)	6.53 (4.70, 12.95)	4.98 (3.50, 7.70)	0.103
Uric acid (µmol/L)	374.59±110.83	421.62±92.39	327.57±109.86	0.009
Albumin (g/L)	26.47±7.75	27.11±6.02	25.83±9.30	0.628
Anti-Sm antibodies (+), n (%)	19 (52.80)	10 (55.56)	9 (50.00)	0.742
Anti-dsDNA antibodies (+), n (%)	18 (50.00)	11 (61.11)	7 (38.89)	0.189
Serum IgG (g/L)	13.65±6.26	16.28±6.28	11.03±5.17	0.010
Serum C3 (g/L)	0.70±0.35	0.59±0.35	0.82±0.31	0.047
Serum C4 (g/L)	0.14±0.11	0.12±0.12	0.16±0.10	0.386
Urine protein (g/24h)	2.08 (1.90, 3.71)	2.19 (1.76, 4.23)	4.46 (1.07, 7.31)	0.268
Urine pH	6.07±0.61	6.02±0.74	6.11±0.46	0.667
Urine titratable acid (mmol/L)	17.22±11.68	16.39±13.10	18.06±10.39	0.675
Urine ammonia (mmol/L)	30.81±19.05	26.56±22.35	35.06±14.48	0.185
Biopsy index				
Modified NIH activity index	5.33±3.52	6.83±3.854	3.83±2.431	0.046
Endocapillary hypercellularity, n (%)	5 (13.89)	3 (16.67)	2 (11.11)	0.635
Neutrophils/karyorrhexis, n (%)	13 (36.11)	8 (44.44)	5 (27.78)	0.298
Fibrinoid necrosis, n (%)	4 (11.11)	4 (22.22)	0 (0.00)	0.037
Hyaline deposits, n (%)	32 (88.89)	18 (100.00)	14 (77.78)	0.097
Cellular/fibrocellular crescents, n (%)	15 (41.67)	5 (27.78)	10 (55.56)	0.091
Interstitial inflammation, n (%)	16 (44.44)	10 (55.56)	6 (33.33)	0.359
Modified NIH chronicity index	2.06±2.52	2.39±3.05	1.72±1.87	0.435
Total glomerulosclerosis score, n (%)	16 (44.44)	7 (38.89)	9 (50.00)	0.597
Fibrous crescents, n (%)	9 (25.00)	6 (33.33)	3 (16.67)	0.268
Tubular atrophy, n (%)	18 (50.00)	8 (44.44)	10 (55.56)	0.958
Interstitial fibrosis, n (%)	8 (21.05)	6 (33.33)	2 (11.11)	0.117
ISN/RPS class				
III, n (%)	6 (15.79)	6 (33.33)	0 (0.00)	<0.001
IV, n (%)	12 (31.58)	12 (66.67)	0 (0.00)	<0.001
V, n (%)	18 (50.00)	0 (0.00)	18 (100.00)	<0.001

Notes: Values for categorical data were given as a number (percent); values for continuous variables were expressed as mean \pm standard deviation (normally distributed data) or median (interquartile range) (non-normally distributed data); values for renal histopathology indices were expressed as median (minimum, maximum). A two-tailed P<0.05 was considered statistically significant. Bold values denote statistical significance at the P<0.05 level.

Abbreviations: SLE, systemic lupus erythematosus; SBP, systolic blood pressure; DBP, diastolic blood pressure; SLEDAI-2K, systemic lupus erythematosus disease activity index; WBC, white blood cell; eGFR, estimated glomerular filtration rate; BUN blood urea nitrogen; IgG, immunoglobulin G; C3, complement C3; C4, complement C4; NIH, National Institutes of Health; ISN/RPS, International Society of Nephrology/Renal Pathology Society.

differ between the two groups. The SLEDAI-2K was significantly higher in the proliferative LN group (12.83 ± 5.54 vs 8.61 ± 3.31 , p=0.009). Patients with proliferative LN had a significantly higher prevalence of hemolytic anemia than those with membranous LN, and there were no other disease activity indicators showing the difference between the two groups. Moreover, patients with proliferative LN had significantly higher serum creatinine, uric acid, and IgG levels, but lower hemoglobin, eGFR and C3 levels than those with membranous LN. In addition, in the comparison of renal histopathological features, higher levels of modified NIH activity index and the proportion of patients with fibrinoid necrosis were observed in proliferative LN.

MPC Expression in the Renal Tissue of LN

To study the role of MPC in patients with LN, we compared the levels of MPC1 and MPC2 between proliferative LN (n=18) and membranous LN (n=18) by immunohistochemical staining (Figures 1 and 2). Both MPC1 and MPC2 were exclusively detected in renal tubules in enrolled LN patients. Renal biopsies from patients with membranous LN showed more intense staining of MPC1 and MPC2 in the tubules compared to those with proliferative LN (p=0.002, p<0.001, respectively) (Figures 1C and 2C).

Association Between MPC and Features of LN

The associations between tubular MPC expression and the clinical and histologic features of LN were further analyzed in Table 2. In respect of clinical characters, the MPC1 was negatively correlated with SLEDAI-2K scores and rSLEDAI scores. And the MPC2 was only negatively correlated with SLEDAI-2K scores. Regarding the laboratory data, the MPC1 was positively associated with eGFR and urine ammonia, whereas it was inversely correlated with serum creatinine, BUN, uric acid, and anti-dsDNA antibodies. There were significant negative correlations between the MPC2 and serum creatinine, BUN, uric acid and anti-dsDNA antibodies. MPC2 was positively associated with eGFR and urine associated with eGFR associated with e

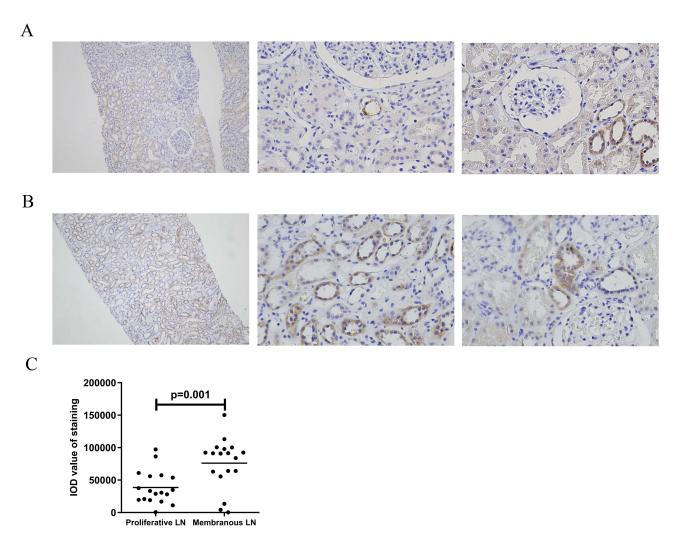


Figure 1 MPC1 expression in kidney tissues of LN. Representative immunohistochemical staining of MPC1 in proliferative LN ((**A**), n=18), and membranous LN ((**B**), n=18) groups, respectively. Left column (Magnification×100). Right two columns (Magnification×400). (**C**) shows a comparison of quantitative analysis of MPC1 expression between proliferative LN and membranous LN using Image-Pro Plus 6.0 software. **Abbreviations:** LN, lupus nephritis; IOD, integrated optical density.

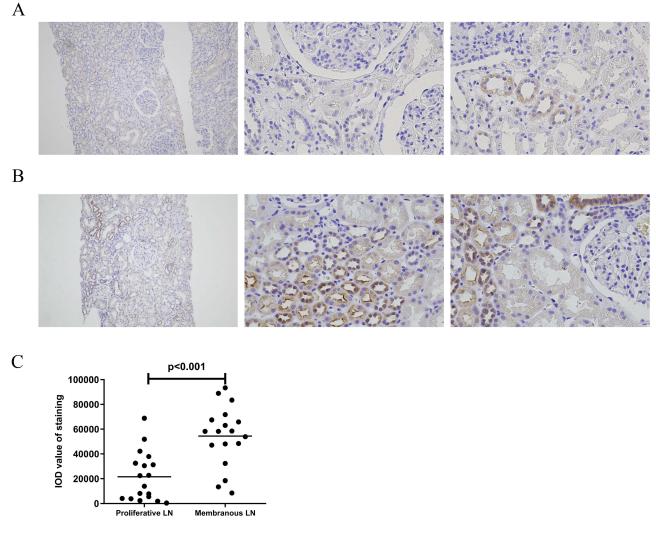


Figure 2 MPC2 expression in kidney tissues of LN. Representative immunohistochemical staining of MPC2 in proliferative LN ((**A**), n=18), and membranous LN ((**B**), n=18) groups, respectively. Left column (Magnification×100). Right two columns (Magnification×400). (**C**) shows a comparison of quantitative analysis of MPC2 expression between proliferative LN and membranous LN using Image-Pro Plus 6.0 software. **Abbreviations:** LN, lupus nephritis; IOD, integrated optical density.

As for renal pathological features of LN, both MPC1 and MPC2 had significant negative correlations with modified NIH activity index and neutrophils/karyorrhexis. However, there was no association between MPC1/MPC2 and other biopsy indexes (Table 2).

	MPCI		MPC2	
	r	P value	r	P value
Clinical features				
Gender	0.329	0.050	0.259	0.127
Age (years)	0.098	0.571	0.252	0.137
SLE duration (months)	0.107	0.536	0.057	0.074
Hypertension	-0.209	0.211	-0.078	0.653
Nephrotic syndrome	-0.03 I	0.859	0.092	0.594
SLEDAI-2K	-0.510	0.001	-0.364	0.029
rSLEDAI	-0.424	0.011	-0.234	0.176

Table 2 Associations Between MPC and Features of Lupus Nephritis (n=36)

(Continued)

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	MPCI		MPC2	
	r	P value	r	P value
Laboratory assessment				
eGFR (mL/min/1.73 m2)	0.484	0.003	0.343	0.041
Serum cystatin C (mg/L)	-0.448	0.145	-0.140	0.665
Serum creatinine (µmol/L)	-0.633	<0.001	-0.525	<0.001
BUN (mmol/L)	-0.487	0.003	-0.526	0.001
Uric acid (µmol/L)	-0.576	<0.001	-0.397	0.017
Albumin (g/L)	0.239	0.161	0.055	0.749
Anti-Sm antibodies (+)	0.340	0.052	0.121	0.484
Anti-dsDNA antibodies (+)	-0.380	0.022	-0.390	0.019
Serum IgG (g/L)	0.028	0.870	0.031	0.857
Serum C3 (g/L)	0.159	0.354	0.130	0.451
Serum C4 (g/L)	0.121	0.481	0.039	0.823
Urine protein (g/24h)	-0.122	0.478	-0.005	0.977
Urine pH	0.244	0.151	0.235	0.167
Urine titratable acid (mmol/L)	0.174	0.309	0.056	0.746
Urine ammonia (mmol/L)	0.391	0.018	0.385	0.020
Biopsy index				
Modified NIH activity index	-0.422	0.010	-0.372	0.025
Endocapillary hypercellularity	-0.159	0.356	-0.244	0.152
Neutrophils/karyorrhexis	-0.508	0.002	-0.451	0.006
Fibrinoid necrosis	-0.289	0.087	-0.086	0.619
Hyaline deposits	-0.160	0.351	-0.062	0.720
Cellular/fibrocellular crescents	-0.253	0.137	-0.301	0.075
Interstitial inflammation	-0.167	0.331	-0.133	0.440
Modified NIH chronicity index	-0.082	0.636	-0.060	0.728
Total glomerulosclerosis score	0.012	0.944	0.052	0.761
Fibrous crescents	-0.032	0.853	-0.260	0.125
Tubular atrophy	-0.059	0.734	-0.028	0.870
Interstitial fibrosis	-0.235	0.167	-0.247	0.147

Table 2 (Continued).

Notes: A two-tailed P<0.05 was considered statistically significant. Bold values denote statistical significance at the P<0.05 level.

Abbreviations: MPC, Mitochondrial pyruvate carrier; SLE, systemic lupus erythematosus; SLEDAI-2K, systemic lupus erythematosus disease activity index 2000; rSLEDAI, renal systemic lupus erythematosus disease activity index; eGFR, estimated glomerular filtration rate; BUN, blood urea nitrogen; IgG, immunoglobulin G; C3, complement C3; C4, complement C4; NIH, National Institutes of Health.

Discussion

This retrospective study showed that MPC1 and MPC2 were uniquely expressed in renal tubules in LN patients. Both MPC1 and MPC2 expression were significantly decreased in proliferative LN compared to membranous LN. Moreover, tubular MPC1 levels were positively correlated with eGFR, while it was inversely associated with serum creatinine, BUN and uric acid, SLEDAI-2K, rSLEDAI, modified NIH activity index, and neutrophils/karyorrhexis in enrolled LN patients, suggesting that MPC1 expression was significantly associated with renal function and disease activity. The analogous correlation was also observed in the MPC2 group.

Accumulating data has demonstrated that mitochondrial dysfunction plays an essential role in the pathogenesis of LN.⁸ It has been reported that injured renal tubular cells in LN have a defect in fatty acid oxidation, which causes mitochondrial dysfunction, reprograms them to a pro-fibrotic phenotype, and contributes to their death.^{25,26} Abundant mitochondria provide energy in tubular epithelial for the reabsorption of various solutes.²⁷ Pyruvate is critical for mitochondrial ATP generation and energy metabolism, while MPC activity is necessary for several processes that require the presence of pyruvate inside mitochondria. Hence, we focus our attention on the significance of MPC,

a gatekeeper in mitochondrial metabolism for transporting pyruvate, which is a central substrate in the metabolism of fatty acid, carbohydrate, and amino acid. We observed that both MPC1 and MPC2 expression were exclusively detected in renal tubules in proliferative LN and membranous LN. MPC was positively associated with renal tubular damage marker urine ammonia, which reinforces the potential association of mitochondrial metabolism with renal tubular function in LN. Furthermore, MPC expression was significantly decreased in proliferative LN compared to membranous LN, and the mitochondrial dysfunction caused by decreased expression of MPC in kidney may be involved in the progression of LN disease. However, the results in LN are not unique, and similar results have been seen in other glomerular nephritis. In our previous study, the results of immunohistochemical staining indicated the expression of MPC1 and MPC2 in the DN were similar to those seen in LN. The expression of MPC in minimal change disease, membranous nephropathy, IgA nephropathy and focal segmental glomerulosclerosis served as positive controls.

Mitochondrial dysfunction is not only one of the pathological mediators in renal diseases but also an important participant in the progression. Previous studies have shown that increased production of ROS, up-regulated expression of cytochrome C oxidase I and IV, and inactivated mitochondrial respiratory chain complex IV in patients with CKD IV-V stage, indicating that mitochondrial dysfunction was closely related to the progression of CKD.^{28,29} Han et al showed that Niclosamide ethanolamine salt, a mitochondrial uncoupler, attenuated lupus nephritis in MRL/lpr mice through decreasing urinary excretion of tubular injury biomarkers, inhibiting tubular proliferation and suppressing renal interstitial inflammation and fibrosis.³⁰ In the present research, MPC was correlated with markers of renal function, including eGFR, serum creatinine, cystatin C, BUN. Hyperuricemia is an independent risk factor for the progression of renal function in CKD patients, and its mechanisms include the induction of epithelial-mesenchymal transition, the activation of the renin-angiotensin system, and endothelial dysfunction.³¹ Both MPC1 and MPC2 were negatively correlated with serum uric acid levels in LN. Mitochondrial dysfunction caused by reduced expression of renal tissue MPC may be involved in LN progression. Another important finding was that the MPC correlated significantly with the disease activity of LN. Both the MPC was correlated with renal SLEDAI score and anti-dsDNA antibodies. More importantly, the MPC is significantly associated with renal pathology activity index rather than the chronicity index, which is concordant with the observed correlation with clinical renal disease activity. A growing body of research has documented that the activity index in renal histopathology is a predictor of long-term renal outcomes.^{32,33} The correlation between the MPC and the activity index suggests a prediction potential of the MPC for a long-term renal outcome, which needs further large-scale studies.

There are several limitations in the present study that should be considered. Firstly, it shares all the limitations of observational, single-center studies. The relatively small sample size limited us from performing subgroup analyses or more advanced statistics. Subsequent studies should consider expanding the sample size in order to improve the statistical power and verify the results. Secondly, the MPC expression was semi-quantified by immunohistochemistry and this study mainly focused on the results of LN, which makes the current study more limited. Further studies on MPC expression in multiple types of kidney diseases should be conducted to obtain more meaningful results. Thirdly, long-term follow-up data is missing for further investigation of the impact of the MPC on the renal prognosis in LN. Last but not least, the interrelationship and precise role of the MPC-pyruvate-mitochondria axis in the tubular damage in LN are worthy of mechanistic studies in the future.

In conclusion, our observations indicate that both MPC1 and MPC2 were exclusively expressed in renal tubules of LN, and decreased levels were significantly associated with unfavorable clinicopathological features.

Data Sharing Statement

The datasets analyzed during the current study are available from the corresponding author on reasonable request.

Ethics Approval and Consent to Participate

All experiments and methods were performed in accordance with relevant guidelines and regulations. This study protocol was reviewed and approved by the Ethics Committee of the First Affiliated Hospital of Nanjing Medical University and was conducted in accordance with the Declaration of Helsinki. Informed consent was obtained from all individual participants included in the study.

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Disclosure

The authors declare that they have no competing interests.

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