

# Urinary Hypoxia-Inducible Factor-1alpha Levels Are Associated with Histologic Chronicity Changes and Renal Function in Patients with Lupus Nephritis

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**Purpose:** Tubulointerstitial hypoxia in the kidney is considered a hallmark of injury and a mediator of the progression of tubulointerstitial fibrosis. Hypoxia-inducible factor-lalpha (HIF-lalpha), a master transcription factor in cellular adaptation to hypoxia, regulates a wide variety of genes, some of which are closely associated with tissue fibrosis. The present study set out to characterize urinary HIF-1alpha expressions in patients with lupus nephritis (LN) and to explore whether urinary HIF-1alpha expressions are associated with histologic chronicity changes and renal function. Materials and Methods: Urinary HIF-1alpha levels were measured by enzymelinked immunosorbent assays in 42 patients with LN and in 30 healthy controls. Activity and chronicity indexes as well as tubular HIF-1alpha expressions were analyzed for each specimen. Results: Urinary HIF-1alpha levels were higher in LN patients than in healthy controls  $(3.977\pm1.696 \text{ vs. } 2.153\pm0.554 \text{ ng/mL}, p<0.001)$  and were associated with histologic chronicity indexes (r=0.463, p<0.01) and eGFR (r=-0.324, p<0.05). However, urinary HIF-1alpha levels showed no correlation with histologic activity indexes, anti-dsDNA, ANA, complement 3 and 4 levels, proteinuria, systemic lupus erythematosis disease activity index, and WHO pathological classification (p>0.05). Conclusion: Urinary HIF-1alpha levels were elevated in LN patients and were associated with histologic chronicity changes and renal function, indicating that HIF-1alpha might contribute to histologic chronicity in LN.

**Key Words:** Hypoxia-inducible factor-1alpha, lupus nephritis, histologic chronicity change, renal function

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#### INTRODUCTION

Lupus nephritis (LN) is one of the main causes of morbidity and mortality in systemic lupus erythematosus (SLE). Although the glomerular abnormalities of LN have attracted much attention, the mechanisms that lead to tubulointerstitial fibrosis and damage in LN remain obscure. Tubulointerstitial injury is characterized by tubular atrophy and interstitial fibrosis. Pathological analysis shows that the extent

of tubulointerstitial damage is better correlated with impaired renal function than the degree of glomerular injury in patients with chronic kidney disease (CKD) and it may be less amenable to treatment compared with glomerular lesions.<sup>3</sup>

Accumulating evidence emphasizes chronic hypoxia in the tubulointerstitium as a final common pathway to end-stage renal disease (ESRD), and the final common pathway of chronic hypoxia operates principally in the tubulointerstitium. As Renal hypoxia in early and late stage of CKD is secondary to imbalances in vasoactive factors, obliteration and loss of the interstitial capillary, reduced efficiency of oxygen diffusion, increased oxygen demand from hyperfiltration and tubular hypertrophy. Recent evidence shows that hypoxia may accelerate CKD progression by affecting renal cell survival, inflammatory cell infiltration, ECM deposition, and epithelial to mesenchymal transition (EMT). As Acceptable 1.

The family of hypoxia-inducible factors (HIFs) are central regulators of hypoxic responses.<sup>12</sup> HIFs regulate several target genes that have important functions in renal physiologic and pathophysiologic processes, including energy metabolism, vasomotor regulation, angiogenic growth, matrix metabolism, and apoptosis/cell survival. 13-17 HIF-1 is a heterodimer composed of a constitutively expressed HIF-1B subunit and an O2-regulated HIF-1alpha subunit. HIF-1alpha plays a central role in the hypoxic response of tubular epithelial cells. 10 Indeed, hypoxia-inducible gene expression in primary renal proximal tubular epithelial cells is almost completely blocked by inactivation of HIF-1alpha, suggesting that their response to hypoxia is largely dependent on HIF-1alpha.<sup>18</sup> In addition, genetic ablation of HIF-1alpha inhibited the development of tubulointerstitial fibrosis *in vivo*. <sup>19</sup> In summary, HIF-1alpha appears to be a critical contributor to the progression of tubulointerstitial fibrosis due to hypoxia.

In this study, we investigated urinary HIF-1alpha levels in patients with LN and analyzed the associations between urinary HIF-1alpha levels and the clinical features in LN patients.

# **MATERIALS AND METHODS**

#### **Subjects**

Forty-two patients with LN who underwent renal biopsy at Hainan Provincial People's Hospital from 2003 January to 2010 December were identified via an electronic medical

record system. All 42 patients fulfilled at least four of the 1997 American College of Rheumatology revised criteria for SLE and had persistent proteinuria greater than 0.5 g/day or cellular casts, as described in the criteria for renal involvement.<sup>20</sup> This study was approved by the Institutional Review Board of Hainan Provincial People's Hospital and informed consent from all patients was obtained.

# Histological classification of LN and morphologic evaluation

Kidney specimens from 42 patients with LN were stained with hematoxylin-eosin and then classified according to the International Society of Nephrology/Renal Pathology Society 2003 classification.<sup>21</sup> Activity indices (AI) and chronicity indices (CI) were calculated in proliferative forms.<sup>22</sup> Quantitation of interstitial inflammation and chronic tubulo-interstitial damage (TID) is based on the percentage of cortex involved by interstitial fibrosis and tubular atrophy. A semi-quantitative grading of TID and interstitial inflammation was described as follows: 0=none; 1=involvement of <25%; 2=involvement of 26-50%; 3=involvement of 51-75%; 4=involvement of >76%.

#### Urinary HIF-1alpha assays

Urinary HIF-1alpha levels were measured in 42 patients with LN and 30 healthy controls, using an enzyme-linked immunosorbent assay kit (Groundwork Biotechnology Diagnostics Inc., San Diego, CA, USA, Catalogue number: H108-9). All experiments were performed in duplicate.

#### Statistical analysis

Statistical analysis was performed using SPSS software version 11.0. Results are reported as means±SD Student's ttest was used to compare urinary HIF-1alpha levels between the LN group and the control group. Correlations between two data sets were assessed using Spearman's correlation test. *p*-values of <0.05 were considered significant.

# RESULTS

# Patients' demographic and clinical information

Table 1 summarizes the demographic and clinical information of the 42 patients with LN. Fifteen (35.71%) patients demonstrated nephrotic range proteinuria. The most frequent histological class was WHO class IV (20/42, 47.62%). There were 8 cases with mixed classes: 5 with III plus V and 3

Table 1. Patients' Demographic and Clinical Information

	LN patients	Normal control
Number	42	30
Age (mean±SD, yrs)	24.07±11.14	31.2±8.55
Gender, female	38 (90.48%)	18 (60%)
SBP (mm Hg)	122.71±28.06	109.33±9.8
DBP (mm Hg)	80.81±18.12	66.17±5.52
Complement 3 (g/L)	0.51±0.28	
Complement 4 (g/L)	$0.12 \pm 0.07$	
Positive rate of ANA, n (%)	32 (76.19%)	
Positive rate of anti-dsDNA, n (%)	21 (50%)	
Albumin (g/L)	25.98±6.44	
Proteinuria (g/d)	$3.08\pm2.07$	
eGFR (mL/min/1.73 m <sup>2</sup> )	82.80±35.75	
SLEDAI	13.17±5.56	
WHO classification		
II	2 (4.76%)	
III	5 (11.9%)	
IV	20 (47.62%)	
V	7 (16.67%)	
III plus V	5 (11.9%)	
IV plus V	3 (7.14%)	

LN, lupus nephritis; SD, standard deviation; SBP, systolic blood pressure; DBP, diastolic blood pressure; ANA, antinuclear antibody; eGFR, estimated glomerular filtration rate; SLEDAI, systemic lupus erythematosis disease activity index; WHO, World Health Organization.

Values are expressed as mean±SD or number of patients (percent).

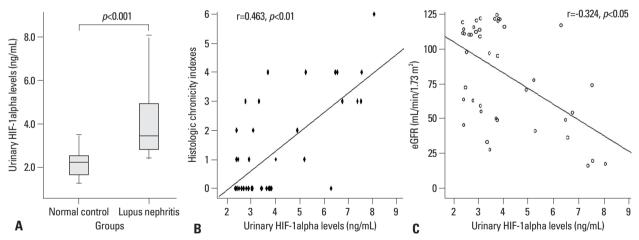


Fig. 1. Urinary HIF-1alpha levels in patients with lupus nephritis and in healthy controls. Urinary HIF-1alpha levels were significantly higher in patients with lupus nephritis than in healthy controls (3.977±1.696 vs. 2.153±0.554 ng/mL, p<0.001). The bars indicate group mean values (A). Urinary HIF-1alpha levels were associated with histologic chronicity indices (r=0.463, p<0.01) (B) and eGFR (r=-0.324, p<0.05) (C). HIF, hypoxia-inducible transchiption factor.

with IV plus V (Table 1).

# Urinary HIF-1alpha levels in patients with LN and in healthy controls and correlation between urinary HIF-1alpha levels and the clinical parameters of LN

Urinary HIF-1alpha levels were significantly higher in patients with LN than in healthy controls (3.977 $\pm$ 1.696 vs. 2.153 $\pm$ 0.554 ng/mL, p<0.001) (Fig. 1A). Urinary HIF-1alpha levels were associated with histologic CI (r=0.463,

p<0.01) (Fig. 1B) and eGFR (r=-0.324, p<0.05) (Fig. 1C), but not with anti-dsDNA, ANA, C3, C4, daily proteinuria amounts or histologic AI.

# **DISCUSSION**

LN is a severe organ manifestation of SLE and a major cause of morbidity. Whereas the immunopathogenesis of

glomerular lesions in LN has been extensively studied, less is known of tubulointerstitial disease and its strong association with less favorable long-term renal prognoses.<sup>3</sup> It has been suggested that hypoxia plays a critical role in the progression of CKD to ESRD,<sup>4,7</sup> and tubulointerstitial hypoxia in the kidney has been considered a hallmark of injury and a mediator of disease progression.<sup>4,5</sup> Hypoxia is a profibrogenic stimulus of tubular cells<sup>23</sup> and changes the ECM metabolism of resident renal cells.<sup>24,25</sup> In addition to ischemic acute renal failure, hypoxia also plays a crucial role in the development of nephrotoxic acute kidney injury, radiocontrast nephropathy, and acute glomerulonephritis.<sup>26,27</sup>

The chief mediators of this hypoxic response are hypoxiainducible factor-1 (HIF-1) and its oxygen-sensitive component HIF-1alpha. HIF activity and the effect of HIF activation on CKD progression remain controversial. Chronic hypoxia accompanied by the up-regulation of HIF-regulated genes has pivotal roles in the progression of renal damage from ischemia/reperfusion injury,<sup>31-33</sup> the early stage of the uninephrectomized Thy1 nephritis,<sup>34</sup> the remnant kidney model<sup>35-37</sup> and an obese, hypertensive type 2 diabetes rat.<sup>38</sup>

Nevertheless, long-term overactivation of HIF-1α in chronic kidney diseases is implicated to be pathogenic and is associated with the progression of chronic renal injuries. 4,28-30,32,39-41 A profibrotic role of cross talk between the transforming growth factor (TGF)-\(\beta\)1 and HIF-1alpha signaling pathways has been reported in human renal tubular epithelial cells, as HIF-1α was shown to be necessary for hypoxic, normoxic and TGF-β1-stimulated renal cell fibrogenesis. 42 Wang, et al. 43 demonstrated that HIF-1α mediates angiotensin II-induced profibrotic effects through activation of cell transdifferentiation. Recently, Higgins, et al.<sup>19</sup> also showed that hypoxia enhanced EMT of primary tubule cells through HIF-1alpha in vitro, and deletion of HIF-1alpha in tubule cells in a murine model of CKD inhibited the development of tubulointerstitial fibrosis. Kimura, et al.41 also demonstrated that stable expression of HIF-1alpha in tubular epithelial cells promoted interstitial fibrosis. HIF-1 alpha has also been reported to be up-regulated in chronic renal diseases. 4,28,29,32,44 In summary, overactivation of the HIF signaling pathway might promote the progression of tubulointerstitial fibrosis, and normalization of overactivated HIF-1α under pathological conditions should antagonize excessive HIF-1alpha activity and restore normal physiological regulation. Therefore, inhibition of HIF-1alpha may be used as a treatment strategy to reduce fibrogenesis as a result of chronic kidney damage. 45 In our present study, we found that urinary HIF-1alpha levels in LN patients were significantly higher than that of healthy controls, demonstrating that HIF-1alpha is increasingly expressed in the tubulointerstitial compartment vs. healthy subjects.

Our study also found the good correlations between urinary HIF-1alpha levels and the histological chronicity index eGFR, suggesting that expressed HIF-1alpha might participate in the chronicity and renal function of LN. Although the chronicity index is known to represent fibrosis, the individual components of the chronicity index indicate the degree of fibrous tissue accumulation. Considering that HIF-1alpha is expressed in renal tubular epithelial cells, our finding that urinary HIF-1alpha levels might be more reasonable to correlated with chronicity index than activity index. Thus, it would be interesting to investigate whether urinary HIF-1alpha levels can help predict renal outcome.

In summary, urinary HIF-1alpha levels were elevated in LN patients and associated with histologic chronicity changes and renal function, indicating that HIF-1alpha might contribute to histologic chronicity and renal function in LN.

The limitations of our present study include the small number of patients enrolled and lack of control for renal disease groups, as cases of different types of glomerulone-phritis (i.e. primary and secondary glomerulone-phritis), as well as other tubulointerstitial diseases (i.e. allograft ne-phropathy, drug-induced tubulointerstitial nephritis) or glomerular diseases without interstitial damage are present in all chronic renal damage cases. In addition, it remains to be proven that there is a positive correlation between urinary and renal HIF-1alpha levels and whether urinary HIF-1alpha levels accurately represent renal HIF-1alpha levels.

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