

Role of toll receptors in diabetic nephropathy

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ABSTRACT

Objectives: Diabetic nephropathy is the leading cause of chronic kidney disease. The pathogenesis of DN remains incompletely understood. It has been recently demonstrated that inflammatory processes play a significant role in the development and progression of DN. Toll-like receptors play a fundamental role in the innate immune system by triggering proinflammatory signaling pathways. Our aim is to evaluate the expression of TLRs on monocytes and relate their expression with inflammation in HD patients with & without diabetic nephropathy. **Method:** In a case control study (60) patients from Alkasr El Aini Hospital on hemodialysis were divided into two groups: Group 1, 30 patients on hemodialysis not due to diabetic nephropathy, Group 2, 30 patients on hemodialysis due to diabetic nephropathy, compared to Group 3, including 30 healthy controls. All participants were subjected to: Full medical history, complete physical examination, Serum creatinine, uric acid, A1C, fundus examination, detection of TLR2, TLR expression by real time PCR in peripheral blood mononuclear cells. Data were statically calculated using SPSS, comparison between groups was done using student T test comparing 2 groups, correlation using spearman's correlation. **Results:** Diabetic had significantly increased TLR2, TLR4 mRNA in peripheral blood mononuclear cells compared to controls and non diabetics patient on hemodialysis ($p < 0.001$), TLR2, TLR4 significantly correlated with dialysis duration in diabetic ($p < 0.001$), no correlation with A1C in relation to TLR2 ($p = 0.078$), TLR4 ($p = 0.163$). **Conclusion:** TLR2, TLR4 were significantly elevated in diabetic on dialysis initiating event in the patho-

genesis of DN, providing a link between hyperglycemia and hypoxia with inflammation and fibrosis within the kidney. Hence, therapeutic interventions aimed at targeting the inflammatory component through interruption of TLR signaling may be a novel strategy to target prevention and treatment of DN.

KEYWORDS

Diabetic Nephropathy; Toll Receptor 2, 4

1. INTRODUCTION

Diabetic nephropathy has become the most common cause of end stage renal disease (ESRD) in developed countries, which is mainly due to the increasing prevalence of Type 2 diabetes [1]. Although DN is traditionally viewed as a non-immune disease, emerging evidence suggests that inflammatory mechanisms play an important role in disease pathogenesis and progression.

Indeed, the decline of renal function in diabetic patients with proteinuria is positively associated with tubulointerstitial inflammation [2].

From a therapeutic perspective, blockade of the renin-angiotensin system coupled to strictglycemic and blood pressure control is the best documented treatment strategy for DN.

Besides, the initial promises shown in early clinical trials, indicate many patients still progress relentlessly to ESRD [2].

Therefore, investigations into the mechanisms underlying intrarenal inflammation may provide new therapeutic targets for anti-inflammatory strategies against DN.

Toll-like receptors (TLRs) are a conserved family of pattern recognition receptors that play a fundamental role in the innate immune system by triggering proinflammatory signaling pathways in response to microbial pathogens. In addition, TLRs are also activated by endogenous

agonists of no microbial origin and are involved in non-infectious inflammatory conditions [3].

Recently, TLRs have been implicated in the pathogenesis of acute and chronic renal disorders. [4]

Among the 11 human TLRs, TLR4 and/or TLR2 might be a molecular link between inflammation and diabetes as they promote tubulointerstitial inflammation during diabetic nephropathy [5].

Therefore we intend to study the relation and the levels of TLR2 and TLR4 in end stage diabetic nephropathy.

2. PATIENTS AND METHODS

2.1. Population of Study & Disease Condition

90 Patients from Kasr El Ani University Hospital on hemodialysis (HD) due to diabetic nephropathy, patients non-diabetic nephropathy & group of healthy people. All participants are divided into three groups: Group 1, 30 patients on hemodialysis not due to diabetic nephropathy, Group 2, 30 patients on hemodialysis due to diabetic nephropathy & Group 3, 30 participants including healthy control.

2.2. Exclusion Criteria

Patients receiving statins, or having cancer or autoimmune disease or recently hospitalized for infectious diseases.

2.3. Methodology in Details

All patients underwent full medical history, complete physical examination, Funds examination to all participants of Group 2, Serum creatinine, BUN, CRP, HbA1c, Membrane expression of TLR2 and TLR4.

Serum creatinine and blood urea. Glycosylated hemoglobin

Detection of TLR2&TLR4 gene expression by real time-PCR: Three ml peripheral venous blood was withdrawn by clean venipuncture on EDTA vacutainer tubes from cases as well as 20 healthy volunteers Plasma and serum samples were separated by immediate centrifugation at 4°C (1500 g for 10 min), aliquoted and stored at -80°C until analysis.

Isolation of mononuclear cells from whole blood was performed through use of Ficoll-Paque Premium (Biorchrom, Berlin, Germany).

-RNA Extraction:

Total RNA was extracted from cell pellet using SV Total RNA Isolation System kit, Promega, Madison VVI, USA .

It was designed for isolation of total intranuclear RNA from fresh, whole blood treated anticoagulated with

EDTA.

Quantitation and Storage of RNA:

*To determine the concentration and purity of RNA, absorbance at 260 nm and 280 nm was measured in a spectrophotometer.

cDNA Synthesis:

The extracted RNA was reverse transcribed into cDNA using: AMV Reverse Transcriptase, Promega, and Madison. VVI, USA

qPCR (quantitative real time PCR)

Statistical methodology

Data were statistically calculated using SPSS, comparison between groups was done using student t test, correlation using spearman's correlation

3. RESULTS

90 Patients from Kasr El Ani University Hospital on hemodialysis due to diabetic nephropathy, patients non-diabetic nephropathy & group of healthy people. All participants are divided into three groups: Group 1, 30 patients on hemodialysis not due to diabetic nephropathy, Group 2, 30 patients on hemodialysis due to diabetic nephropathy & Group 3, 30 participants including healthy control (**Table 1** showing Demographic and laboratory: data of patients and control groups).

There is statistically significant correlation between level of TLR2 in end stage diabetic nephropathy compared to end stage non diabetic nephropathy and control group (**Figure 1**).

There is statistically significant correlation between level of TLR4 in end stage diabetic nephropathy compared to end stage non diabetic nephropathy and control groups (**Figure 2**).

Table 1. Demographic and laboratory data of patients and control groups.

	Group 1 (non-DM)	Group 2 (DM)	Control group
AGE	50.8 ± 11.22	56 ± 10.9	42.1 ± 4.8
SEX M/F	17/13	17/13	30/0
D/HDx	5.4 ± 3.39	1.7 ± 1.6	
Associated diseases			
*HTN	73.3%	80%	0%
*IHD	33.3%	20%	0%
*Both	13.3%	30%	
Cr (mg/dl)	11.08 ± 2.26	9.8 ± 2.6	1.08 ± 0.28
Urea (mg/dl)	180.0 ± 39.6	168.7 ± 48.9	32.9 ± 5.9
HbA1c %	4.6% ± 0.4%	7.2% ± 0.7%	3.45 ± 0.65%
TLR2	0.28 ± 0.16	1.18 ± 0.54	0.15 ± 0.06
TLR4	0.5 ± 0.2	2.13 ± 0.9	0.4 ± 0.14

D/HDx (duration of hemodialysis)

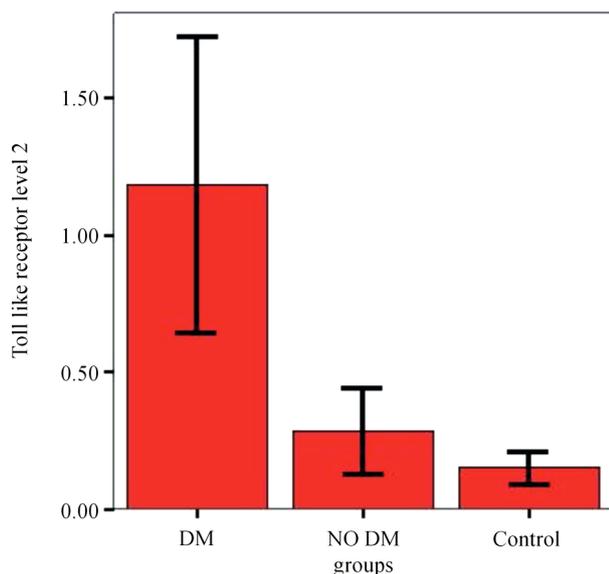


Figure 1. Comparison Group 1 (non-DM), Group 2 (DM) & Group 3 (control group) as regarding the level of toll like receptors 2.

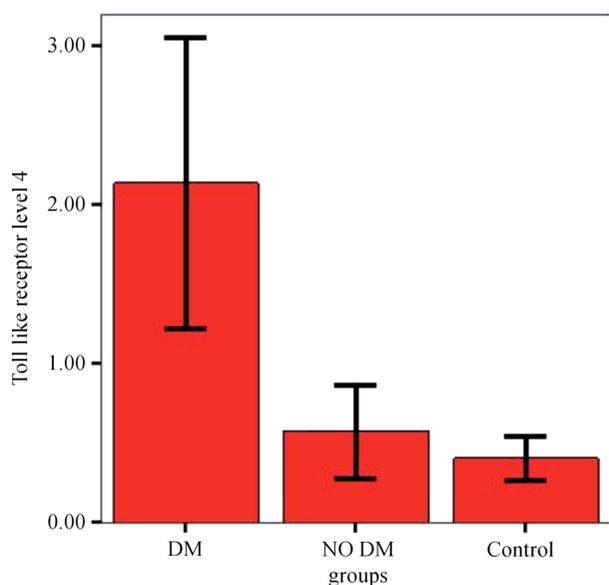


Figure 2. Comparison Group 1 (non-DM), Group 2 (DM) & Group 3 (control group) as regarding the level of toll like receptors 4.

Correlation between toll-like receptors with different clinical and laboratory data, showing a correlation between the duration of hemodialysis and TLR2 and TLR4 ($r = 0.661$, $p = 0.001$), ($r = 0.5$, $p = 0.003$) respectively which is statistically significant.

The correlation between glycosylated hemoglobin and TLR2 and TLR4 is ($r = 0.169$, $p = 0.372$) and ($r = 0.231$, $p = 0.219$) respectively which is not statistically significant shown in **Table 2**.

The correlation between the duration of diabetes and

TLR2 and TLR4 is ($r = 0.147$, $p = 0.734$) and ($r = 0.111$, $p = 0.559$) respectively which is not statistically significant.

There is correlation between the duration of hemodialysis and TLR2 and TLR4 ($r = 0.83$, $p = 0.0001$) and ($r = 0.712$, $p = 0.1$) respectively which is statistically significant.

The correlation between glycosylated hemoglobin and TLR2 and TLR4 is ($r = 0.326$, $p = 0.078$) and ($r = 0.261$, $p = 0.163$) respectively which is not statistically significant (results shown in **Table 3**).

There is statistically significant correlation between level of TLR2 and duration of hemodialysis in end stage diabetic nephropathy compared to end stage non diabetic nephropathy shown in **Figure 3**.

There is statistically significant correlation between level of TLR4 and duration of hemodialysis in end stage diabetic nephropathy compared to end stage non diabetic nephropathy shown in **Figure 4**.

4. DISCUSSION

Diabetes has become the most common single cause of end-stage renal disease worldwide, this is due to the facts that:

- Diabetes, particularly Type 2, is increasing in prevalence.
- Diabetes patients now live longer.

About 20% - 30% of patients with Type 1 or 2 diabetes develop evidence of nephropathy, but in Type 2 dia-

Table 2. Toll like receptors correlation with different clinical/laboratory data in Group 1 (non-DM).

clinical/laboratory data	TLR2		TLR4	
	r	p	r	p
age	-0.106	0.579	0.091	0.632
D/HDx	0.661	0.001	0.523	0.003
Cr (mg/dl)	-0.335	0.071	-0.154	0.416
Urea (mg/dl)	0.270	0.149	0.324	0.08
HbA1c%	0.169	0.372	0.231	0.219

Table 3. Toll like receptors correlation with different clinical/laboratory data in Group 2 (DM).

clinical/laboratory data	TLR2		TLR4	
	r	p	r	p
AGE	0.278	0.137	-0.299	0.108
D/DM	0.147	0.437	0.111	0.559
D/HDx	0.832	0.001	0.712	0.001
Cr (mg/dl)	0.110	0.564	0.92	0.630
Urea (mg/dl)	-0.062	0.746	0.022	0.906
HbA1c%	0.326	0.078	0.261	0.163

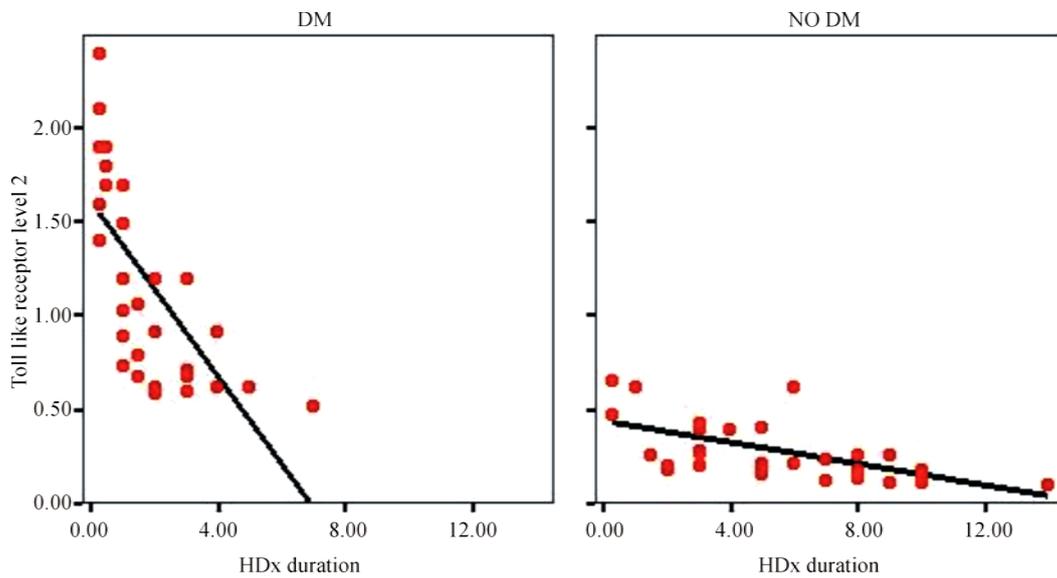


Figure 3. Correlation of TLR2 and duration of hemodialysis.

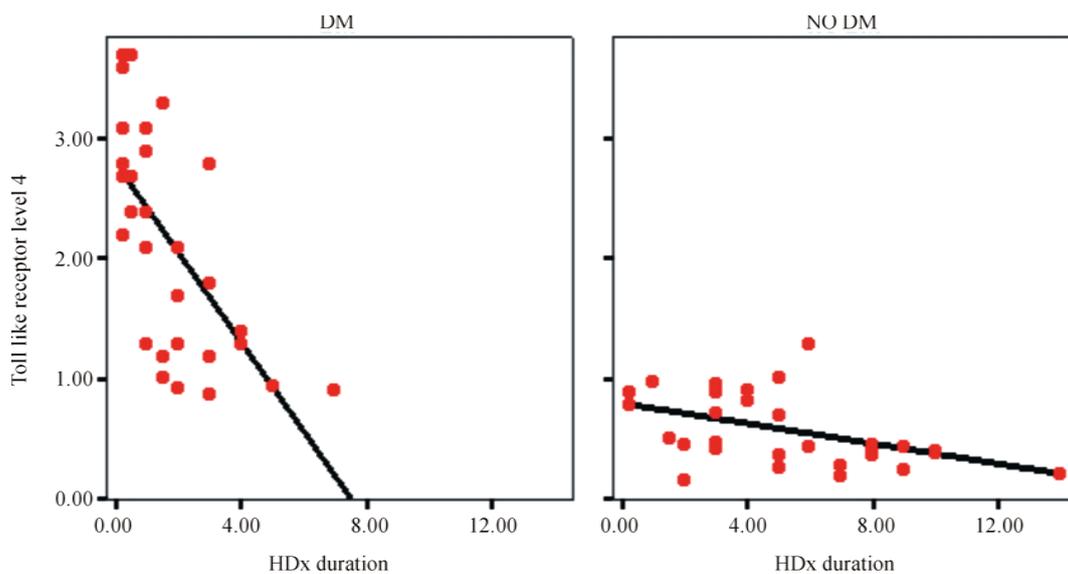


Figure 4. Correlation of TLR4 and duration of hemodialysis.

betes, a considerably smaller fraction of this progress to end stage renal disease (ESRD).

However, because of much greater prevalence of Type 2 diabetes, such patients constitute over half of those diabetic patients currently starting on dialysis [6].

Diabetes Mellitus is a major risk factor for heart, kidney diseases, and lower limb amputations. Emerging *in vitro* and *in vivo* data suggest that systemic inflammation plays a role in the pathogenesis of diabetes mellitus complications via innate immune receptors. Mainly nephropathy [7].

Diabetic nephropathy is traditionally viewed as no immune disease; emerging evidence suggests that inflammatory mechanisms play an important role in disease pa-

thogenesis and progression. Indeed, the deterioration of renal function in diabetic patients with proteinuria is positively associated with tubulointerstitial inflammation [5].

Inflammatory response is a recognized factor in the pathogenesis of diabetic nephropathy. Numerous experimental and clinical studies have shown the participation of different inflammatory molecules and pathways in the path physiology of this complication [5].

Although albuminuria is widely used and is considered the best marker for renal damage in diabetic patients, certain limitations should be taken into consideration. First, not all patients with proteinuria will develop progressive renal dysfunction [8]. Additionally, 30% of diabetic patients with renal impairment have normoalbu-

minuria [9]. Second, the cardiovascular and renal morbidity is elevated in the “high normal” range of UAE [10]. Third, a number of variables affecting UAE lack standardization including urine collection methods, reporting of test results, reference intervals for albumin-to-creatinine ratio and lack of a complete reference system for urine albumin and creatinine measurements [10]. Finally, multiple markers of renal dysfunction, either urinary or serum, can appear before the detection of microalbuminuria, suggesting that microalbuminuria occurs once significant kidney damage has already occurred [10].

Toll-like receptors (TLRs) recognize molecular patterns relating to a variety of microbial infections. Stimulation through TLRs leads to activation of antigen-presenting cells, production of inflammatory cytokines creating inflammation, and production of Type 1 interferons (IFNs) that include IFN- α and β , and exerts direct effects on regulatory cells. These effects can direct the immune response, dealing with the immediate problems of infection and activating more specific responses of the adaptive immune system [11].

The recognition of microbial components by mammalian TLRs plays an important role in the activation of the innate immune response and subsequent proinflammatory reactions. TLRs also interact with ligands generated at sites of injury [12].

TLRs present on the cell surface recognize bacterial and fungal components, whereas intracellular TLRs recognize viral or microbial nucleic acids. In addition, TLRs also interact with endogenous ligands, such as oxLDL (oxidized LDL), HSPs (heat-shock proteins) 60 and 70, fibrinogen and fibronectin, which are also elevated in Diabetes Mellitus [13]. Thus different TLRs are amenable to targeting by different types of agents [13].

Among the TLRs, TLR2 and TLR4 play a critical role in the pathogenesis of inflammation and diabetes, in both experimental and clinical conditions [14].

The interactions among increased glucose levels, elevated NEFAs [non-esterified (“free”) fatty acids] and resultant pro-inflammatory cytokines in diabetes have clear implications for the immune system [15].

Mohammad *et al.*, 2006 [16,17], (Creely *et al.*, 2007) [15] and (Song *et al.*, 2006) [18] reported increased TLR2 and TLR4 expression in bone marrow-derived macrophages of Type 1 diabetic NOD mice, correlating with increased NF- κ B (nuclear factor κ B) activation in response to end toxins and increased pro-inflammatory cytokines.

It was reported that the increase in TLR2 and TLR4 expression, intracellular signaling and TLR-mediated inflammation in monocytes has a significant correlation with HbA1c (glycated hemoglobin) levels in diabetic patients. [14]

Also, TLR2 and TLR4 play an important role in atherosclerosis.

TLR2 and TLR4 bind to components of the gram positive and gram negative bacteria, respectively [19]. They are expressed in multiple cells and tissues, primarily in monocytes. TLR2 and TLR4 expression is increased in atherosclerotic α plaque macrophages and in animal models of atherosclerosis [20].

Our study included 3 groups: Group 1: 30 patients with end stage renal disease on regular hemodialysis due to non-diabetic nephropathy, Group 2: 30 patients with end stage renal disease due to diabetic nephropathy on regular hemodialysis and Group 3: 30 healthy volunteers, Patients receiving statins, or having cancer or autoimmune disease or recently hospitalized for infectious diseases have been excluded.

We aimed in this study to evaluate the expression of TLRs on monocytes and relate their expression with inflammation in hemodialysis patients with & without diabetic nephropathy.

We took full history of all patients, complete physical examination, fundus examination of all patients, we measure serum creatinine, blood urea, level of HbA1c, level of toll-like receptors 2 and 4.

Our results included that:

- The level of TLRs 2 and 4 was higher in patients with end stage renal disease due to diabetic nephropathy ($p \leq 0.01$) than patients with end stage renal disease due to non-diabetic nephropathy and control group.
- The level of TLRs 2 and 4 has no correlation with demographic data of these patients independent on sex and age.
- The level of TLR2 and 4 has no correlation to clinical/laboratory data of patients including serum keratinize, blood urea, HbA1c, duration of diabetes mellitus. However, there was positive correlation between the level of TLRs 2 and 4 with duration of hemodialysis ($p \leq 0.01$).

This result partially agreed with Xanthi Zikou *et al.*, (2011) [21], his study concluded that chronic kidney disease CKD patients and patients with diabetic nephropathy are characterized by increased expression of TLRs 2 and 4, on monocytes, that may contribute to their increased inflammatory state. He divided the subjects into: Group 1 included 37 CKD patients, not having diabetes mellitus. Group 2 included 19 CKD patients with diabetic nephropathy. Both groups were compared with age matched controls, (control group). His results showed that Patients of Group 1 exhibited increased membrane expression only of TLR2 in monocytes compared with the control group ($p < 0.02$). Patients of Group 2 presented increased membrane expression of both TLR2 and TLR4 compared with the control group ($p < 0.003$ and $P < 0.001$, respectively) and increased expression of TLR4 compared with Group 1 ($p < 0.02$).

Our result also, partially agreed with (Miao Lin *et al.*,

(2012) [22], his study concluded overexpression of TLR4 in the human diabetic kidney suggesting a possible role for TLR4 but not TLR2 in mediating monocot/macrophage recruitment and tubulointerstitial inflammation in diabetic nephropathy. His study was on Kidney biopsy tissues were obtained from nine subjects with Type 2 diabetes and biopsy-proven DN, nine subjects with Type 2 diabetes lacking nephropathy, nine normal control subjects, and nine subjects with no diabetic nephrotic syndrome. His result showed that the Renal Cortical TLR4 but Not TLR2 was elevated and Correlated with Infiltrating CD68+ Monocytes/Macrophages in diabetic nephropathy Biopsies.

Also, our result partially agreed with (Mohan R. *et al.*, (2010) [23], his study concluded that TLR2 and TLR4 expression, their ligands, signaling, and functional activation are increased in recently diagnosed T2DM and contribute to the proinflammatory state. In his study he measured the freshly isolated monocytes from healthy human controls (n = 23) and T2DM subjects (n = 23) using real time RT PCR, Western blot, and Flow cytometric assays. His result showed that T2DM subjects had significantly increased TLR2, TLR4 mRNA and protein in monocytes compared to controls (p < 0.05). This result in his study agreed with our result. However, he showed that this increased TLR2, TLR4 mRNA and protein in monocytes was correlated with HbA1c which did not agree with our result.

Also, our result partially agreed with (Sridevi Devaraj *et al.*, (2011) [24], his study concluded that there is significant role for TLR2 in promoting inflammation and early changes of incipient diabetic nephropathy. His study was on mice, He induced T1DM in TLR2 knockout mice and wild-type littermates using streptozotocin (STZ). Fasting blood, peritoneal macrophages, and kidneys were obtained for flow cytometry, Western blot, microscopy, and cytokine assays at 6 and 14 weeks after induction of diabetes. His result was that Macrophage TLR2 expression and MyD88-dependent signaling were increased in diabetic mice (wild type_STZ) compared with non diabetic wild type mice.

5. CONCLUSIONS

Based on our findings that the level of toll-like receptors 2 and 4 is elevated in patients with end stage diabetic nephropathy compared to that of patients with end stage renal disease due to non-diabetic nephropathy, TLRs 2 and 4 are significant serum markers of diabetic nephropathy and play a critical role in pathogenesis of diabetic nephropathy. Thus, we recommend the following:

- We recommend further studies on the Toll-like receptors 2 and 4 to correlate them with the different stages of diabetic nephropathy.

- The use of anti-inflammatory therapeutic agents and their value on the change of the level of Toll-like receptors 2 and 4.

CONFLICT OF INTEREST

None.

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