

Prediction of prognosis and renal outcome in lupus nephritis

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ABSTRACT

Lupus nephritis (LN) is a severe manifestation of SLE, characterised by subendothelial and/or subepithelial immune complex depositions in the afflicted kidney, resulting in extensive injury and nephron loss during the acute phase and eventually chronic irreversible damage and renal function impairment if not treated effectively. The therapeutic management of LN has improved during the last decades. but the imperative need for consensual outcome measures remains. In order to design trials with success potentiality, it is important to define clinically important short-term and longterm targets of therapeutic and non-therapeutic intervention. While it is known that early response to treatment is coupled with favourable renal outcomes, early predictors of renal function impairment are lacking. The information gleaned from kidney biopsies may provide important insights in this direction. Alas, baseline clinical and histopathological information has not been shown to be informative. By contrast, accumulating evidence of pronounced discrepancies between clinical and histopathological outcomes after the initial phase of immunosuppression has prompted investigations of the potential usefulness of per-protocol repeat kidney biopsies as an integral part of treatment evaluation, including patients showing adequate clinical response. This approach appears to have merit. Hopefully, clinical, molecular or genetic markers that reliably reflect kidney histopathology and portend the long-term prognosis will be identified. Novel non-invasive imaging methods and employment of the evolving artificial intelligence in pattern recognition may also be helpful towards these goals. The molecular and cellular characterisation of SLE and LN will hopefully result in novel therapeutic modalities, maybe new taxonomy perspectives, and ultimately personalised management.

INTRODUCTION

Lupus nephritis (LN) is one of the most severe complications in SLE, affecting 35%–60% of the patients depending on ethnicity, sex and age of disease onset among other factors.^{1–4} Genetic and nephron cargo, SLE disease course and kidney toxicity, for example, drug-induced toxicity, contribute to long-term impairment of the renal function. Importantly, LN flares constitute a major cause of nephron loss, cumulatively contributing to renal function deterioration.

Current therapies for LN are not sufficiently efficacious in inducing remission or preventing new flares and not all patients show adequate treatment responses. In fact, fewer than 30% achieve complete remission within 6 months of therapy.^{5 6} Despite decelerating rates over the last decades, up to 20% of patients who have been afflicted by LN will ultimately develop end-stage kidney disease within the first decade of their disease course.7-11 For this reason, prediction of the long-term renal outcome at early stages of the disease is of vital importance. Thus, several studies have sought to identify early clinical features, laboratory tests and molecular mechanisms that are associated with unfavourable renal prognosis, in order to optimise the surveillance and interventions in these patients.

The targets of management of patients with LN can be divided into short-term (prevention of flares) and long-term targets (prevention of renal function impairment). Current research in the field focuses on identification of clinical, histological and molecular markers of activity, damage, treatment response and renal prognosis.

CLINICAL PREDICTORS

Baseline clinical and laboratory clinical data do not predict renal outcome

Different investigations have examined clinical and serological markers as predictors of renal prognosis. The information retrieved at the time of active renal disease (baseline), including proteinuria and conventional serological markers, has not shown ability to adequately predict the long-term kidney outcome, in terms of renal function. This was for example the case in two randomised clinical trials, the MAINTAIN Nephritis Trial¹² and the Euro-Lupus Nephritis Trial,¹³ as shown in post hoc analyses by Tamirou *et al*,¹⁴ and Dall'Era *et al*,¹⁵ respectively.

Early response predicts good long-term renal prognosis

By contrast, early decrease of proteinuria levels over 6 months of treatment has shown ability to predict a more favourable long-term renal outcome compared with patients with



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persistently high-grade proteinuria,¹⁶ and a recent series of three studies of similar design concluded that the most useful clinical predictor of good long-term renal outcome (defined using a cut-off for serum concentrations of creatinine at 7 years after the LN episode) is proteinuria levels <0.7–0.8 g/day at month 12 from treatment initiation.^{14 15 17} Nevertheless, while this clinical target could identify patients who had a good renal outcome with a high positive predictive value, it could not predict the outcome of patients who did not achieve the target since the negative predictive value of this proteinuria cut-off was low in two of the three investigations.^{14 15} Thus, the challenge to find appropriate predictors remains, at least predictors of unfavourable renal prognosis.

Persistent microscopic haematuria does not add prognostic value

Cellular casts in the microscopic analysis of the urinary sediment are included in SLE classification criteria sets¹⁸ ¹⁹ and constitute an important diagnostic tool to raise suspicion of kidney involvement. Importantly, findings in the urinary sediment differ substantially between proliferative and non-proliferative glomerulonephritis, with haematuria being mostly associated with proliferative disease,²⁰ pointing to another diagnostic utility of the urinalysis.

Urinalysis may also be useful in differential diagnostics between disease entities, based on, for example, whether proteinuria or haematuria is the predominant aberrance. For example, while proteinuria typically is the main finding in urinalysis in LN, haematuria is a more prominent abnormality in antineutrophil cytoplasmic antibody-associated vasculitis. In the latter, haematuria mainly derives from glomerular endothelial cell and surface layer injuries, resulting in necrotising and crescentic glomerulonephritis and ultimately a severe impairment of the glomerular filtration barrier.²¹

However, the usefulness of the urinalysis in clinical studies has in recent years been debated; the main reason for that being the fact that quantification of its components is characterised by a profound variability between assessors and between consecutive measurements. Moreover, urinalysis has been withdrawn in the newly proposed 2019 European League Against Rheumatism/American College of Rheumatology classification criteria for SLE.²²

More importantly, results from urinalysis, in particular haematuria either before the initial immunosuppressive therapy or at 1 year from treatment initiation, have not shown prognostic ability regarding the long-term kidney outcome, and addition of such results to the proteinuria response did not improve the prognostic value of proteinuria¹⁵ or even worsened it.¹⁴ Additionally, haematuria has in previous investigations shown weak or no correlation with activity features in kidney biopsies.²³

One of the reasons for this is undoubtedly the fact that different laboratories analyse the urine sediment differently, and thus, no homogenised data can be retrieved for use in trials. Moreover, the urine volume may affect the accuracy of the measurements of cellular counts, with large volumes resulting in greater dilutions of the cellular elements in the pellets. In the same manner, the time of the collection may affect the dilution grade for elements of importance, normally with higher concentrations in the first morning void compared with random collections during the rest of the day. While the aforementioned obstacles could be overcome with clear instructions and standardisation, other important barriers include the multiple reasons for haematuria, especially in female patients, and the high grade of subjectivity during the assessment. Indeed, not all red cells are indicative of glomerular injury, and accurate recognition of dysmorphic acanthocytes of glomerular origin contra red blood cells originating from the lower urinary tract needs special training.

Using more sophisticated approaches such as multivariable Cox proportional hazards regression models, Mackay *et al*²⁴ recently developed panels of routine clinical, laboratory and histological parameters (termed hazard index tools) to predict key long-term outcomes, that is, new or progressive chronic kidney disease, severe kidney injury and permanent kidney replacement. It is worth noting that proteinuria and serum creatinine at month 12 were found to be important components in these tools, whereas urinalysis was not shown to add value.

HISTOLOGICAL PREDICTORS Diagnostic kidney biopsy

At clinical suspicion of LN, it is important to perform a kidney biopsy in order to determine the LN class and exclude mimicking conditions, such as antiphospholipid antibody or antiphospholipid syndrome-associated nephropathy, IgA nephropathy, hypertensive nephrosclerosis, diabetic nephropathy and thin basement membrane disease. This information dictates the decision of treatment.²⁵⁻²⁷ Indeed, proliferative (2003 International Society of Nephrology/Renal Pathology Society (ISN/RPS) class III and IV) and membranous (2003 ISN/ RPS class V) LN differ with regard to the long-term renal outcome, thus demanding different surveillance and therapeutic management.²⁸ By contrast, a previous study of 98 patients with incident proliferative LN showed no difference across subclasses within the proliferative spectrum (2003 ISN/RPS class III, IV-S and IV-G) regarding long-term prognosis.¹⁰ In fact, the S (for segmental) and G (for global) designations have been suppressed in the new proposed revision of the ISN/RPS classification,²⁹ currently awaiting embracement.

The importance of the National Institutes of Health (NIH) chronicity index in the initial kidney biopsy in portending long-term prognosis was demonstrated by Austin *et al* as early as in 1983,³⁰ which however was not confirmed later by the Lupus Nephritis Collaborative Study Group³¹ or in more recent retrospective data from the LN database of the Université catholique de Louvain

(unpublished). This point is further discussed in the repeat biopsy section.

The initial kidney biopsy also provides information about the afflicted domains within the kidney, that is, the extent of the injury in the glomerular versus the tubulointerstitial compartment. Although current classification sets mainly focus on glomerular lesions, the importance of tubulointerstitial injury and damage in short-term and long-term prognosis has been repeatedly highlighted in the literature.^{32–37} Proteinuria, immune complex deposition in the interstitium, proinflammatory molecules on renal tubular cells and rupture of the Bowman's capsule and cryptic antigen presentation by juxtaglomerular cells are some of the insults resulting in interstitial infiltration by inflammatory cells and, ultimately, tubular atrophy,^{38–43} collectively constituting a strong rationale for inclusion of the tubulointerstitial compartment in classification sets, prognostic markers and outcome measures.

The role of the repeat kidney biopsy

The role of the repeat kidney biopsy in patients with LN has been discussed rigorously during the last decades, but consensus among researchers and physicians has yet to be established. Before elaborating on the role of the repeat biopsy, it is important to make clear distinctions between different scenarios in which such repeat biopsies can be performed and how nomenclature has been used in the literature. As discussed in a recent editorial by Anders,⁴⁴ five different scenarios could be described by the term 'repeat biopsy', that is, the per-protocol repeat biopsy at a predefined time point for treatment evaluation and new decision of therapy, the partial response repeat biopsy for distinguishing between residual activity and delayed healing and guide treatment accordingly, the flare repeat biopsy, the repeat biopsy to support withdrawal of the immunosuppressive treatment and the chronic kidney disease (CKD) progression repeat biopsy to determine the grade of nephrosclerosis contra treatable active injury.

Even if the nomenclature and definitions have not been used uniformly in studies of repeat kidney biopsies, several investigations have shown a discordance between clinical and histological outcome after the initial phase of immunosuppressive therapy for LN. More specifically, most studies reporting results from repeat biopsies have shown that residual renal activity may be evident in repeat biopsies from a considerable proportion of patients who have shown complete clinical responses to treatment, the latter mainly based on the proteinuric outcome.^{45–49} Again, as discussed above, haematuria levels have been demonstrated to yield weak or no correlations with activity components at the level of tissue in both initial and control kidney biopsies.²³

The discrepant patterns between clinical and histological data at the time of the repeat kidney biopsy have prompted investigations on the role of the tissue-level information in tailoring treatment and portending the long-term kidney outcome. While the former question has yet to be addressed in prospective studies, several studies have attempted to address the latter one.

Associations between chronic tissue damage in repeat kidney biopsies and long-term impairment of the renal function have been demonstrated in both European and Hispanic LN populations.^{45 47} Nevertheless, this was not confirmed in another study,⁵⁰ indicating a need for validation. The role of residual activity in repeat kidney biopsies as a marker of the long-term kidney outcome is even less clear. Thus, the idea of a prospective multicentric study of per-protocol repeat kidney biopsies to provide evidence for optimised surveillance and management receives indeed increasing embracement within the LN researcher community.⁴⁴ In this direction, a recent retrospective investigation of incident cases of proliferative LN demonstrated that different histological components in per-protocol repeat kidney biopsies showed ability to portend renal relapses and long-term renal function impairment (unpublished data). In this study, high NIH activity index scores in the repeat kidney biopsies were predictive of subsequent relapses, especially activity in the glomerular compartment, and high NIH chronicity index scores were associated with poor long-term renal prognosis, especially chronic damage in the tubulointerstitial compartment. The discrepancies with regard to the association between chronic damage in the initial kidney biopsy and long-term impairment of the renal function across studies may be due to differences in study design, as well as, importantly, due to improvements in the diagnosis and management of LN in the last decades, resulting in less kidney damage accrued in recent compared with earlier studies. Thus, the chronic changes in repeat kidney biopsies of recent studies, especially in patients who failed to respond to immunosuppressive therapy, may be similar in amount and prognostic attributes to the respective changes in initial biopsies of earlier investigations.

Altogether, accumulating evidence strongly supports the usefulness of repeat kidney biopsies as an integral part of treatment evaluation, including LN patients showing adequate clinical response. Thus, a new prospective study is currently being designed within the frame of the Lupus Nephritis Trials Network and will be entitled 'Per-protocol repeat kidney biopsy in incident cases of lupus nephritis' or, shortly, REBIOLUP. The objectives of the project will be to determine the percentage of LN patients in pathological remission after 12 months of standard of care immunosuppression, correlate histological and immunological (based on immune deposits) response to therapy with clinical response and evaluate whether therapeutic decisions steered by the results of a per-protocol repeat kidney biopsy improve renal outcomes compared with a matched control group of patients who will not undergo repeat kidney biopsy.

In brief, patients with an incident biopsy-proven proliferative or membranous LN, or combinations thereof, selected to be initiated at standard of care immunosuppressive therapy with either mycophenolate mofetil or intravenous cyclophosphamide according to the

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Euro-Lupus regimen¹³ (combined with glucocorticoids and ACE inhibitors or angiotensin II receptor blockers in both cases) will be eligible to be enrolled in the study. At baseline, patients will be randomised 1:1 to either undergo or not undergo a per-protocol repeat kidney biopsy at month 12 from baseline. In patients with 2003 ISN/RPS class III/IV (±V) at baseline and an NIH activity index score >3 (cut-off based on retrospective unpublished data) in the repeat kidney biopsy, the immunosuppressive therapy will be intensified based on the physician's and patient's shared decision. In patients with pure membranous (2003 ISN/RPS class V) LN at baseline, individual assessment of the repeat biopsy will steer the decision of treatment, based on, for example, evaluation of immune deposits in electron microscopy or spike formation. Results from this study, including centralised evaluation of electron microscopy in baseline and repeat kidney biopsies, are anticipated to generate data on how to evaluate response to therapy in pure membranous LN, as well as the value of the information retrieved from repeat kidney biopsies in portending long-term renal prognosis. Patients who have not undergone a repeat biopsy will be treated according to standard clinical parameters and, finally, percentages of complete renal response at month 24 and renal impairment at month 60 will be compared between the two study arms.

MOLECULAR BIOMARKERS AND TISSUE-BASED APPROACHES

In parallel with drug research, identification of serum and urinary non-invasive biomarkers that accurately reflect renal activity at the level of tissue and predict long-term renal outcome have been a topic of extensive study during the last decades.⁵¹ Investigations include single-molecule evaluations⁵²⁻⁵⁷ and large semi-unbiased proteomic screenings⁵⁸ of serum markers, while urinary biomarkers receive increasing interest⁵⁹⁻⁶² for two important reasons. First, molecules in urine are likely to be directly excreted from the kidney, which makes them highly relevant in the context of LN and other renal disorders. Second, urine samples are readily accessible for examination, and standardisation and commercialisation of urinary markers as diagnostic and/or predictive tools would therefore mean a paramount advance in the management of LN. Recent auspicious implications from single-cell RNA-sequencing pointed to strong correlations between gene expression of immune cells in urine specimens and corresponding kidney leukocytes,⁶³ indicating that urinary biomarkers may prove useful as a surrogate for kidney biopsies in the near future.

Identification of clinical markers that portend the longterm prognosis of the lupus kidney is anticipated to lead to treat-to-target approaches in the management of LN in the near future. However, the knowledge gap regarding LN pathogenesis and molecular pathways underlying clinical phenotypes remains a concern. Several recent efforts have therefore focused on deepening our understanding of the molecular events that occur in patients with LN during active and quiescent phases, as well as differences with other renal disorders and non-renal SLE. In a recent investigation, Pamfil *et al*⁶⁴ studied intrarenal molecular profiles associated with disease severity using high-throughput transcriptomic approaches on kidney biopsies from two independent cohorts of patients with LN and cadaveric donors. Adding important information to current knowledge, the researchers of this study found decreased kidney function in patients with histological evidence of tubular damage and transcriptomic patterns consistent with T and B cell activation in the kidney, which was corroborated using immunohistochemistry. By contrast, no association between decreased kidney function and systemic disease activity or activity and chronicity in the glomerular compartment was confirmed. In another study of intrarenal transcript expression by Parikh et al,⁶⁵ repeat kidney biopsies were used to associate differentially expressed genes in patients who showed good versus poor clinical response. The researchers found that transcripts associated with T cell activation, complement signalling pathways and type I and II interferon signature discriminated responders from non-responders and took advantage of the serial biopsies to generate a gene signature that discriminated responders from non-responders based on changes in transcript expression following treatment.

Using a more sophisticated approach, another recent study by Panousis *et al*⁶⁶ combined genetic and transcriptome analysis in whole blood samples from patients with SLE and age-matched and sex-matched healthy controls. In a subanalysis, the researchers compared patients with active renal versus active non-renal SLE and found differentially expressed genes enriched in granulocyte activation and antimicrobial humoural response, whereas combining differentially expressed genes in active versus inactive SLE, they could demonstrate a stepwise progression of transcriptome alterations from inactive non-renal SLE to active non-renal and active renal SLE.

Parallel to molecular mapping, different groups have attempted dissection of the cellular constellation of the lupus kidney. In a recent conference abstract, Crickx et al^{67} showed that B cells with plasmablast attributes infiltrate the kidney of patients with active LN before therapeutic intervention, whereas the kidney of patients who failed immunosuppressive therapy with mycophenolate mofetil was enriched with long-lived plasma cells, introducing a prospect of cellular biomarkers of response to therapy. Advancing cellular approaches, Der *et al*⁶⁸ for the Accelerating Medicines Partnership Rheumatoid Arthritis and SLE consortium recently published data from single-cell RNA sequencing on kidney biopsies from LN patients, demonstrating that high interferon response signature and fibrotic signature in tubular cells were associated with failure to respond to therapy and providing novel insights into histological differences across LN subtypes. Such techniques allow detailed characterisation of the cell populations in the kidney tissue, both infiltrating and resident immune cells as well as parenchymal cells, potentialising a deeper understanding of the pathophysiology underlying kidney inflammation in SLE in the forthcoming years and making the non-invasive biomarker mining in more accessible biological material, such as serum and urine, more relevant and more individualised.^{69 70}

PERSPECTIVES

To summarise, the management of LN has improved during the last decades, but the imperative need for consensual outcome measures remains. In order to design trials with success potentiality, it is important to define clinically important short-term and long-term targets of therapeutic and non-therapeutic intervention. Ideally, an international task force comprising rheumatologists, nephrologists and patient advocates will soon be formed to serve this aim. The information gleaned from kidney biopsies at active renal disease and after completion of treatment may provide important insights in this direction. Moreover, novel insights of SLE and LN pathogenesis as well as insights obtained from genetic studies, transcriptomics and proteomics will have to be incorporated.⁷¹⁷²

Hopefully, we will soon identify clinical, molecular or genetic markers, or combinations thereof, which reliably reflect renal histopathology and portend the long-term renal outcome. Novel non-invasive imaging methods and employment of the evolving artificial intelligence in pattern recognition may also be helpful towards these goals. Undoubtedly, the molecular and cellular characterisation of SLE and LN will result in novel therapeutic modalities, maybe new taxonomy perspectives, and ultimately personalised management of the patients.

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