Measuring plasma C4D to monitor immune complexes in lupus nephritis

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ABSTRACT

Objective Because currently available assays that measure circulating immune complexes (ICx) are suboptimal, a novel assay was recently developed measuring C4d, a stable product of activation of the classical complement pathway. The present study aimed to establish the value of measuring plasma C4d levels in a longitudinal cohort of patients with severe refractory SLE who were treated with a combination therapy of rituximab with belimumab (RTX+BLM).

Methods Fifteen patients with SLE who were treated with RTX+BLM in a phase 2A, open label study were included to sequentially measure plasma C4d levels and correlated to well-established markers of ICx formation, that is, autoantibodies against double-stranded (ds) DNA, autoantibodies against C1q and proteinuria. The performance of plasma C4d measurements, C4 measurements and the ratio of C4d over C4 (C4d:C4) was evaluated.

Results After establishing that on RTX+BLM treatment kinetics of C4d levels was distinct from traditional C3 and C4 levels, we found strong correlation of C4d:C4 with anti-dsDNA (R=0.76, p<0.001) and anti-C1q (R=0.65, p<0.001) autoantibody levels, which outperformed both stand-alone C4 and C4d levels. Additionally, changes in C4d:C4 over time correlated strongly with changes in proteinuria (R=0.59, p<0.001) as well as anti-dsDNA (R=0.46, p=0.003) and anti-C1q (R=0.47, p=0.002).

Conclusion In patients with severe SLE, plasma C4d levels in relation to C4 levels is useful for longitudinal monitoring after RTX+BLM treatment to reflect amelioration of classical complement activation by ICx as well as proteinuria.
In the present study, we aimed to further establish the value of measuring plasma C4d levels in longitudinal cohort of patients with severe, refractory SLE who were treated with a combination therapy of rituximab (RTX) with belimumab (BLM). The cohort was selected because based on previous studies, the most severe patients with SLE were anticipated to have high C4d levels and a novel, synergic B-cell targeted treatment was postulated to reduce, and even eradicate, ICx-forming autoantibodies. Therefore, optimising the clinical conditions to investigate sequentially measured plasma C4d levels in patients with SLE.

METHODS

Study participants

Patients with severe, refractory SLE who participated in a phase 2A, single-arm, proof-of-concept study investigating combination therapy of rituximab with belimumab (SynBioSe study, registered at ClinicalTrials.gov NCT02284984) were selected for this study. Patients were treated with 1000 mg rituximab at weeks 0 and 2 and with monthly 10 mg/kg belimumab from week 4 onwards. Only patients who completed 24 weeks of follow-up with serial serum sampling were selected. Details of the study participants, treatment and clinical efficacy were published elsewhere. All patients provided written informed consent.

ELISA detecting soluble C4d

Maxisorp 96-well plates (Nunc) were coated with rabbit anti-human C4d neoepitopes-specific antibody. After blocking with washing buffer (50 mM Tris–HCl, 0.15 M NaCl, 0.1% Tween, pH 7.5) supplemented with 3% fish gelatin (Norland Products), patient plasma as well as pooled plasma from healthy volunteers (lacking C4d) supplemented with *Escherichia coli* expressed C4d standard in serial dilutions, were diluted to 4% in phosphate-buffered saline+0.02% Tween-20+0.02 M Na2EDTA and added to the plate for 1 hour incubation at 37°C. Detection was achieved using mouse anti-human C4d antibody (Quidel, #253) followed by horseradish peroxidase-conjugated goat antimouse secondary antibody (Dako, #P0447). Plates were developed using OPD (Kem-En-Tec) as substrate and absorbance was measured at 490 nm using Cary50 microplate reader (Varian).

Reference values

In order to assess complement normalisation, clinical routine reference values were applied for C3 (normal >0.9 g/L) and C4 (normal >0.5 mg/L). For C4d, reference values were based on previous published C4d levels and were categorised according to receiver operating characteristic curve analyses in a longitudinal cohort for lupus nephritis (normal <1.1 mg/L).

Statistical analyses

For patients’ characteristics before and after treatment, descriptive analysis was used and median values reported on the basis of the small cohort size. Non-parametric, two-tailed, paired samples Wilcoxon tests were used to compare treatment effects. To investigate the assay’s performance, non-parametric Spearman’s correlation coefficients were analysed.

RESULTS

Table 1 describes the patients’ disease characteristics illustrating severe and refractory SLE and establishing that RTX+BLM indeed resulted in significant reductions in SLE relevant autoantibody levels. Anti-double stranded (ds) DNA levels declined from a median of 144 IU/mL (range 18–505) at baseline to 56.5 (1–374) at week 24 (p<0.001) and anti-C1q levels declined from 75.1 U/mL (25.1–135) to 27.6 (9.8–71.3) (p<0.001). With respect to immune complex formation, the solid-phase C1q binding test was positive in 5 of 15 patients. As a surrogate, serum levels of C3 and C4 were reduced at baseline and improved significantly on treatment from, respectively, median 0.6 g/L (0.3–1.3) to 1.0 g/L (0.5–1.4) (p<0.001) and 80 mg/L (21–260) to 151 (39–339) (p=0.008). Plasma C4d levels were high at baseline with a median of 2.05 mg/L (0.98–6.40), in comparison to previously published C4d levels in patients with SLE and decreased significantly on treatment to 1.25 mg/L.

### Table 1: Patient characteristics

| Characteristic          | N=15 patients |
|-------------------------|---------------|
| Age (years)             | 31 (19–51)    |
| Sex (F/M)               | 13/2          |
| Disease duration (years)| 9 (2–24)      |
| Relapses (n)            | 4 (1–6)       |
| Before treatment        |               |
| SLEDAI                  | 18 (6–29)     |
| Urine PCR (mg/mmol)     | 107 (11–852)  |
| C4 (mg/L)               | 79 (21–260)   |
| C3 (g/L)                | 0.6 (0.3–1.3) |
| C4d (mg/L)              | 2.05 (0.98–6.40) |
| Anti-dsDNA (IU/ml)      | 144 (18–505)  |
| Anti-C1q (U/ml)         | 75.1 (25.1–135) |
| After treatment (24 weeks)|             |
| SLEDAI                  | 2 (0–13)*     |
| Urine PCR (mg/mmol)     | 46 (9–134)*   |
| C4 (mg/L)               | 151 (39–339)* |
| C3 (g/L)                | 1.0 (0.5–1.4)*|
| C4d (mg/L)              | 1.25 (0.17–3.62)* |
| Anti-dsDNA (IU/ml)      | 56.5 (1–374)* |
| Anti-C1q (U/ml)         | 27.6 (9.8–71.3)* |

Depicted are median (range). *Statistically significant (p<0.05) difference from baseline. PCR, protein:creatinine ratio; SLEDAI, Systemic Lupus Erythematosus Disease Activity Index; dsDNA, double stranded DNA.
We observed that at baseline, C4d levels were abnormal in 14 out of 15 patients (93%) which was significantly more than C4 (abnormal in nine patients, equal to 60%) and comparable to C3 (abnormal in 13 patients, equal to 87%) (Table 2). There was no significant difference for normalisation within 24 weeks of C4d (occurred in six patients (43%)), of C4 (in six patients (67%)) and of C3 (in six patients (46%)). Of interest, out of six patients whose C4d normalised, four patients had normalised C3 levels. Vice versa, out of six patients whose C3 normalised only half had normalised C4d levels suggestive of continued classical complement activation despite normalisation of C3 levels.

We next correlated C4d levels to C3 and C4 levels to establish that C4d levels reflected distinct aspects of complement activation. Indeed, we found no correlation between levels of C4d with either C4 levels (R=0.25, p=0.38) nor C3 levels (R=−0.12, p=0.66). Also, reduction of C4d levels did not correlate to increases in C4 (R=0.02, p=0.90) or C3 (R=−0.20, p=0.20).

Since levels of total C4 showed individual variation, we calculated the ratio of C4d activation product and total C4 (C4d:C4). The performance of these complement markers as a reflection of ICx-mediated inflammation was determined by comparison with other well-established markers of ICx-formation, that is, antibodies against dsDNA (anti-dsDNA) and anti-C1q (anti-C1q) and proteinuria. We observed (figure 1A) the strongest correlations for C4d:C4 with anti-dsDNA (R=0.76, p<0.001) and anti-C1q (R=0.65, p<0.001) autoantibodies which outperformed both C4 (respectively, R=−0.43, p=0.001 and R=−0.62, p<0.001) and C4d (respectively, R=0.67, p<0.001 and R=0.35, p=0.01). Of note, none of the complement markers correlated with proteinuria. Additionally, we investigated whether changes in complement markers accompanied changes in markers of ICx-formation (figure 1B). Again we found that changes in C4d:C4 over time had the strongest correlation with changes in anti-dsDNA (R=0.46, p=0.003), anti-C1q (R=0.47, p=0.002) and proteinuria (R=0.59, p<0.001). Likewise, changes in C4d:C4 outperformed the stand-alone measurements of C4 and C4d.

**Discussion**

The present study demonstrated that in patients with severe SLE measuring plasma C4d levels, and quantifying C4d:C4, was well correlated with current widely employed, surrogate measurements for ICx-formation (ie, anti-dsDNA, anti-C1q levels and proteinuria). In a cohort of patients with SLE with severe lupus nephritis, changes in C4d:C4 was strongly correlated with the renal outcome criterion of proteinuria and outperformed stand-alone C4d and C4 measurements. Also, on treatment, in some cases, C4d levels normalised, while C3 levels were still abnormal. Taken together, this study suggests that C4d quantification could be of added value to conventional complement measurements and warrants further validation for its use in the evaluation of immune complex-mediated diseases, including the wide spectrum of patients with mild to severe SLE, cryoglobulinemic glomerulonephritis or antibody-mediated humoral rejection.

Thus far, assays for measuring circulating immune complexes are insufficient because of the multiplicity of different techniques available for detecting immune complexes, differences in results when using different techniques, lack of reproducibility between the same techniques performed in different laboratories, and lack of studies demonstrating clear clinical predictive value of immune complex measurements. As such, the lack of detectable circulating immune complexes does not exclude the presence of immune complex-mediated diseases, including involvement of complement activation. Therefore, there is a clinical need for the development of more specific biomarkers of classical pathway complement activation. The assay for quantifying plasma C4d levels employs neoantigen-specific antibodies to nascent cleavage epitope found only in C4d but not its parent molecules such as C4 and C4b. Of note, patients with a genetic variation at the C4 null allele can impact the rate of production and catabolism of C4 and C4d. To correct this phenomenon, the ratio of C4d:C4 can be calculated or serial measurements within the same patient can be performed, as exemplified in this study.

Previously, in patients with SLE, the deposition of C4d on reticulocytes and platelets was shown to associate with SLEDAI and was suggested as useful biomarker for monitoring disease activity. A prospective pilot study showed that levels of C4d on circulating cells correlated with C4d staining in lupus nephritis biopsies. An assay panel combining anti-dsDNA, ANA, anti-mutated citrullinated vimentin, erythrocyte C4d and B-cell C4d was shown to be sensitive and specific for the diagnosis of SLE. The present study is the first to add to this body of evidence that C4d levels in relation to C4 levels is a useful marker for longitudinal monitoring in the setting of treatment reflecting amelioration of classical complement activation by immune complexes. Future studies...
Figure 1  Significant association of plasma C4d levels with traditional, surrogate markers for immune complex formation. (A). Levels of C4d and C4 were measured simultaneously from which the C4d:C4 ratio was derived and correlated to circulating levels of autoantibodies directed against double stranded (ds) DNA and C1q and proteinuria (measured by the urine protein:creatinine ratio (UPCR)). The correlation plots of the ratio of C4d:C4 illustrate strong correlation with levels of anti-dsDNA and anti-C1q autoantibodies (R=Spearman’s correlation coefficient). The table reports the correlation coefficients separately for C4d, C4 and C4d:C4 where strong correlations (R≥0.5) are depicted in dark grey filled boxes; moderate correlation (0.3<R<0.5) are depicted in light grey filled boxes; no or weak correlations (R≤0.3) are depicted in non-filled boxes. P values were considered significant when below 0.05. (B). Change in levels of C4d, C4 and C4d:C4 in comparison to baseline levels were calculated. The relative changes of C4d, C4 and C4d:C4 were correlated to the relative changes in circulating levels of autoantibodies directed against dsDNA, C1q and proteinuria. Depicted graphs illustrate the correlation of changes in C4d:C4 with changes in anti-dsDNA (R=0.46; p=0.003) and anti-C1q (R=0.47; p=0.002) autoantibody levels and, most relevantly, changes in proteinuria (R=0.59; p<0.001). In the table, correlation coefficients are reported separately for changes in C4d, C4 and C4d:C4.

will need to elucidate how clinical decision-making can benefit from routine C4d:C4 measurements.

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Competing interests  AMB and MO are named as inventors in a patent application including claims to use of C4d as biomarker.

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