

学 位 論 文

Hydroxychloroquine modulates elevated expression of
S100 proteins in systemic lupus erythematosus

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21 Abstract

22 Objectives: We investigated the effect of hydroxychloroquine (HCQ) on S100A8 and
23 S100A9 serum levels in systemic lupus erythematosus (SLE) patients with low disease
24 activity receiving immunosuppressants.

25 Methods: SELENA-SLEDAI, Cutaneous Lupus Erythematosus Disease Area and
26 Severity Index (CLASI) and serum levels of complement factors, anti-double stranded
27 DNA antibodies, and white blood cell, lymphocyte and platelet counts were used to
28 evaluate disease activity, cutaneous disease activity and immunological activity,
29 respectively. Serum S100A8 and S100A9 were measured at HCQ administration and
30 after 3 or 6 months using ELISA.

31 Results: S100A8 and S100A9 serum levels were elevated at baseline and the magnitude
32 of decrease from baseline at 3 and 6 months after HCQ administration was greater in
33 patients with renal involvement than in those without (baseline: S100A8, $p=0.034$;
34 S100A9, $p=0.0084$; decrease: S100A8, $p=0.049$; S100A9, $p=0.023$). S100-modulating
35 was observed in patients with (n=17; S100A8, $p=0.0011$; S100A9, $p=0.0002$) and
36 without renal involvement (n=20; S100A8, $p=0.0056$; S100A9, $p=0.0012$), and was
37 more apparent in patients with improved CLASI activity scores (improved: S100A8,
38 $p=0.013$; S100A9, $p=0.0032$; unimproved: S100A8, $p=0.055$; S100A9, $p=0.055$). No
39 associations were observed for immunological biomarkers.

40 Conclusion: HCQ may improve organ involvement in SLE by modulating S100 protein
41 levels, especially in patients with renal or skin involvement.

42 Keywords

43 Systemic lupus erythematosus (SLE), Lupus Nephritis, Skin, Hydroxychloroquine,
44 S100A8, S100A9

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49 Introduction

50 Lupus nephritis (LN) is the major important organ involved in systemic lupus
51 erythematosus (SLE) and is caused by immune complexes [1]. Macrophage infiltration
52 can be observed histologically in LN and plays an important role in the pathogenesis of
53 glomerulonephritis; however, the details underlying this process remain unclear.
54 Cellular infiltrates in LN may be induced via responses coordinated between the innate
55 and adaptive immune systems.

56 Little is known about the role of the innate immune response in LN. Myeloid-
57 related proteins (MRPs) such as MRP-8 and -14 have been reported to play important
58 roles in the development of autoimmune nephritis in mice. Both of these proteins are
59 ligands of Toll-like receptor 4 (TLR4) expressed by monocytes, and they also
60 participate in the development of autoreactive CD8+T cells [2]. These proteins have
61 been identified as “danger signals” and are also known as DAMPs (damage-associated
62 molecular patterns). According to Heizmann's group, MRP8 is also known as S100A8,
63 and MRP14 as S100A9 [3]. These proteins are expressed in the cytoplasm of
64 neutrophils and macrophages [4]. Upon activation of these cells through TLR4 and
65 receptor for advanced glycation endproducts (RAGE) stimulation, S100A8 and S100A9
66 are secreted [5,6]. The secreted MRP-8/14 complex can bind to the activated
67 endothelium, assisting phagocytes in migrating anywhere in the body [7,8]. The
68 heterodimeric MRP-8/14 complex can also induce upregulation of CD11b/CD18
69 (MAC-1) on neutrophils, which results in enhanced adhesion to the endothelium [9].

70 Phagocytes expressing MRP-8 and -14 were reported to be involved in the
71 pathogenesis of multiple autoimmune and malignant conditions [10-14]. Serum levels
72 of these protein in SLE patients were positively correlated with disease activity,
73 especially lupus glomerulonephritis [15,16]. Additionally, these proteins were
74 demonstrated histologically to be expressed in renal tissues, with levels proportional to
75 the severity of LN [17,18]. The authors suggested that MRP-8 and -14 can be used as
76 diagnostic and prognostic markers for the progress of LN.

77 Tydén *et al.* observed that serum levels of S100A8 and S100A9 were associated
78 with disease activity in SLE [19]. Moreover, serum levels of these proteins were
79 reduced by treatment of SLE patients with several immunosuppressants. However, there

80 have been no reports on the effect of HCQ, which is used as an adjunctive therapy in
81 SLE patients, on expression of these proteins.

82 On the other hand, HCQ has approved for SLE treatment in Japan since July
83 2015. After the approval of HCQ treatment, many SLE patients with
84 immunosuppressant treatment, especially for SLE in women of child bearing age,
85 started to be treated with additional HCQ in Japan.

86 In this study, we investigated the effect of HCQ treatment on serum levels of
87 S100A8 and S100A9 proteins in SLE patients with low disease activity being treated
88 with immunosuppressants.

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90

91 Patients and Methods

92 Patients

93 This was a single-center, retrospective study. All subjects were diagnosed with SLE
94 using the American College of Rheumatology criteria [20] or the SLICC criteria [21],
95 and started additional HCQ treatment for the first time from September 2015 to
96 December 2017. All patients enrolled in this study had been receiving oral HCQ sulfate
97 (Plaquenil; Sanofi-Winthrop, Paris, France) continuously for at least 3 months.
98 There was no need for additional immunosuppressive treatments, including
99 glucocorticoids, in any patient during the 3 months prior to starting HCQ because of
100 sustained low disease activity in these patients. Low disease activity was defined as
101 SELENA-SLEDAI score of 8 or less with no activity in major organ systems such as
102 renal involvement, NPSLE, cardiopulmonary and vasculitis, current prednisolone or
103 equivalent dose of 10mg per day or less and well-tolerated maintenance doses of
104 immunosuppressant. Informed consent was obtained from all participants. The study
105 was approved by the ethical committee of Kagawa university.

106 HCQ was administered at a dose based on ideal body weight (IBW) calculated
107 using the modified Broca's method: 200 mg daily for IBW < 46 kg; 200 mg and 400 mg
108 on alternate days for IBW \geq 46 kg and < 62 kg; and 400 mg daily for IBW \geq 62 kg.
109 Clinical parameters (age, gender, HCQ dose, immunological biomarkers, disease
110 activity index and skin score) were investigated before and after HCQ treatment.
111 Disease activity was evaluated using the SELENA-SLEDAI 2011 criteria. Cutaneous
112 disease activity was evaluated using the Cutaneous Lupus Erythematosus Disease Area
113 and Severity Index (CLASI). According to CLASI improvement criteria reported by
114 Klein R, et al [22], CLASI disease activity was classified by the principle investigator
115 as improved, unchanged, or worse compared to the previous visit, as described above.
116 Those classified as improved were defined as CLASI responders and those classified as
117 unchanged or worse were defined as CLASI non-responders. Immunologic activity was
118 determined via serum levels of complement factors (C3, C4, CH50), anti-double
119 stranded DNA (dsDNA) antibodies, and counts of white blood cells, lymphocytes and
120 platelets. Serum levels of S100A8 and S100A9 were measured at the time of HCQ
121 administration as well as 3 or 6 months later using ELISA (CircuLex ELISA Kit, MBL)
122 according to the manufacturer's instructions. We investigated whether serum levels of
123 S100A8 and S100A9 proteins were associated with disease activity.

124

125 Statistical analysis

126 Immunological biomarker and serum levels of S100A8 and S100A9 were compared
127 using the Student's t test for continuous variables (or Wilcoxon signed-rank test for
128 non-normally distributed data). Comparisons between groups were performed using the
129 Wilcoxon rank sum test. All p values were two-sided, and p values <0.05 were
130 considered significant. For the statistical analyses, we used the following abbreviations:
131 * p <0.05, ** p <0.01, *** p <0.001 and **** p <0.0001. Data were analyzed using JMP®
132 13 software (SAS Institute Inc., Cary, NC, USA).

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135 Results

136

137 Baseline

138 Pregnant women and patients who started anti-thrombotic therapy or added
139 immunosuppressant after starting HCQ administration were excluded in this study.
140 Thirty-seven SLE patients (33 women, 4 men) with sustained low disease activity for
141 at least 3 months prior to administration of HCQ were enrolled in this study. Their
142 baseline characteristics are shown in Table 1. The mean age was 40.7 ± 13.7 years, the
143 mean disease duration was 14.4 ± 11.5 years, the mean SELENA-SLEDAI score was
144 3.7 ± 1.9 and the mean CLASI activity score was 3.2 ± 3.2 . Seventeen of 39 SLE
145 patients had a history of LN. All 17 patients have been in complete remission for over
146 one year. Serum levels of S100A8 and S100A9 proteins at baseline were significantly
147 higher in SLE patients with a history of renal involvement compared with those
148 without renal involvement ($p=0.034$ and $p=0.0084$, respectively; Figure 1).
149 Involvement of other organs such as skin or neuropsychiatric SLE was not associated
150 with serum levels of S100A8 and S100A9 proteins.

151 No significant correlations between S100 protein levels and CLASI score,
152 SELENA-SLEDAI score, complement levels or anti-dsDNA antibody levels were
153 identified (data not shown).

154

155 Effect of HCQ treatment on levels of S100A8 and S100A9 156 proteins

157 We analyzed changes in S100A8 and S100A9 protein levels prior to HCQ treatment as
158 well as 3 or 6 months after HCQ treatment in 37 SLE patients enrolled in this study.
159 However three of the 37 subjects were excluded at 6 month because of a lack of data
160 during observation for 6 months after HCQ administration. Serum levels of S100A8 and
161 S100A9 proteins decreased significantly 3 and 6 months after HCQ treatment compared

162 with those at baseline (Figure 2). Additionally, the modulating effect of HCQ treatment
163 on serum S100A8 and S100A9 levels was observed in both SLE patients with (n=17;
164 S100A8, $p=0.0011$; S100A9, $p=0.0002$) and without a history of renal involvement
165 (n=20; S100A8, $p=0.0056$; S100A9, $p=0.0012$; Supplementary Material Figure S1).
166 The magnitude of the changes in serum S100A8 and S100A9 levels in SLE patients
167 with renal involvement were significantly higher than in those without renal
168 involvement (S100A8, $p=0.049$; S100A9, $p=0.023$; Figure 3).

169 As for the correlation with skin involvement, there was no significant difference
170 in the magnitude of changes in serum S100A8 and S100A9 levels in SLE patients with
171 or without skin involvement (S100A8, $p=0.224$; S100A9, $p=0.072$; Supplementary
172 Material Figure S2). However, the modulating effect of HCQ treatment on serum
173 S100A8 and S100A9 levels was much more apparent in CLASI responders (CLASI
174 responders: S100A8, $p=0.013$, S100A9, $p=0.0032$; CLASI non-responders: S100A8,
175 $p=0.055$, S100A9, $p=0.055$; Figure 4).

176 In regard to SLE disease activity, SELENA-SLEDAI scores, CLASI activity
177 scores and anti-dsDNA antibody decreased and serum levels of C3 increased
178 significantly in all patients after 3 months' treatments. However, no association between
179 changes in S100A8 and S100A9 levels and immunological biomarkers or SELENA-
180 SLEDAI scores was identified. Next we focused on the association of autoantibody
181 titers with changes in S100A8 and S100A9 protein levels. The effect of HCQ treatment
182 on serum S100 protein levels differed between patients with and without detectable
183 anti-Sjögren's syndrome type B (SS-B) antibody titers. Serum S100A8 and S100A9
184 levels decreased significantly during HCQ treatment only in anti-SS-B antibody-
185 negative SLE patients (data not shown). Additionally, changes in serum S100A9 levels
186 in SLE patients positive for lupus anti-coagulant (LAC) were significantly less apparent
187 than in LAC-negative patients (Supplementary Material Figure S3, Figure 5).

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191 Discussion

192 Several biomarkers, including anti-dsDNA antibodies and/or complement factors, have
193 been suggested to reflect the disease activity of SLE. Recently, it has been suggested
194 that TLRs signaling may play an important role in the pathogenesis of SLE.
195 Additionally, several TLR ligands have been reported to be involved in the
196 pathophysiology of SLE. TLR7 or TLR9 signaling are important for production of the
197 interferons which drive SLE pathogenesis. Additionally, TLR4 has also been reported to
198 participate in the pathogenesis of SLE [23,24]. The S100A8 and S100A9 proteins,
199 which are well-known DAMPs, are also ligands of TLR4. Some reports have indicated
200 that serum levels of S100A8 and S100A9 were related to disease activity or serum titers
201 of anti-dsDNA antibodies [19]. Tantivitayakul *et al.* demonstrated that cells infiltrating
202 the glomeruli and peritubular capillaries in LN expressed S100A8/A9 using
203 immunohistochemical staining [18]. Our study showed that serum S100A8 and S100A9
204 levels significantly increased in SLE patients with renal involvement at baseline in
205 comparison with patients without renal involvement. However, serum levels of S100A8
206 and S100A9 proteins were not associated with disease activity indices such as SLEDAI
207 and CLASI (skin lesion activity). The reason for the difference from previous reports is
208 probably related to the disease activity of the study population. The current study
209 analyzed SLE patients with sustained low disease activity, and many other studies
210 examined patient populations with high disease activity. However, our data indicated
211 that elevation of S100 proteins was observed even in SLE with LN remission (data not
212 shown). This suggested the possibility that S100 proteins are produced from renal
213 lesions in SLE with remission of glomerulonephritis. In other words, we hypothesize
214 that many patients, even during clinical remission of LN, retain some LN activity
215 sufficient to increase the expression of S100 proteins, which is related to the
216 pathogenesis of LN.

217

218 We also investigated changes of S100A8 and S100A9 protein levels during
219 HCQ treatment. Previous reports demonstrated changes in S100 proteins during
220 immunosuppressive treatments, including glucocorticoid (GC), cyclophosphamide and

221 mycophenolate mofetil (MMF), in SLE patients with high disease activity. Tydén *et al.*
222 reported that serum S100A8/A9 levels decreased upon treatment of SLE patients with
223 active glomerulonephritis with immunosuppressive drug such as MMF, azathioprine
224 and GC [19]. However, there have been no reports describing the effect of HCQ
225 treatment on S100 protein levels in SLE patients with low disease activity. We
226 demonstrated that HCQ treatment in SLE patients with sustained low disease activity
227 reduced the levels of S100 proteins significantly. The magnitude of changes of serum
228 S100A8 and S100A9 levels in SLE patients with a history of renal involvement was
229 significantly higher than in patients without renal involvement (S100A8, $p=0.083$;
230 S100A9, $p=0.036$; Figure 3). This result indicated that HCQ could provide additional
231 benefits for LN through reduction of S100 protein expression at the site of
232 glomerulonephritis. Many reports indicated that HCQ use, combined with conventional
233 immunosuppressive therapy for LN, improved renal prognosis [25]. We speculate that
234 HCQ's modulating effect on S100 protein levels in SLE patients with low disease
235 activity might have a beneficial effect on renal prognosis. However, the modulating
236 effect of HCQ on S100 protein levels was observed to some extent among SLE patients
237 regardless of past renal involvement. That is to say, HCQ reduced the levels of S100
238 proteins in SLE cases without glomerulonephritis. Furthermore, a significant
239 modulating effect of HCQ on S100 proteins was observed in SLE patients with skin
240 involvement. Recently, Elloumi *et al.* demonstrated that TLR4, a sensor for DAMPs
241 such as S100 proteins, had an important pathogenic role in cutaneous lupus
242 erythematosus inflammation and LN. Upregulation of TLR4 was demonstrated in skin
243 and renal lesions, and played a role in the pathogenesis of cutaneous and renal disorders
244 in SLE [26]. Our data indicated that the mechanism in reducing renal and/or skin lesions
245 may be due to effects on TLR4 signaling through modulation of S100 protein
246 expression. Furthermore, there were some reports that HCQ reduced the expression of
247 TLR4 in several cell types including trophoblastic cells or macrophages [27,28].
248 According to our data and these previous reports, we suggest that the mechanism of
249 HCQ's effect on skin and or renal lesions in SLE patients may involve modulation of
250 TLR4 signaling by downregulation of both S100 proteins and TLR4 expression.

251 However, the overall mechanisms through which HCQ modulates S100 protein
252 levels remain unclear.

253 Taken together, our results indicate that HCQ had a modulating effect on S100
254 proteins which resulted in inhibition of TLR4 signaling. This represents another
255 example of an effect of HCQ on skin or renal disease in SLE. Furthermore, HCQ
256 reduced the levels of S100 proteins in patients with clinical remission of LN. This may
257 contribute to improved prognosis for LN. However, it remains unclear how HCQ
258 reduced the expression of S100 proteins in SLE. Further studies are needed to clarify
259 this issue.

260 There were some limitations in this study. First, there was no monitoring of
261 HCQ adherence through pharmacological dosage of blood HCQ levels. However,
262 disease activity and immunological biomarker were improved significantly 3 months
263 after HCQ treatment compared with those at baseline. Second, sample size was too
264 small in this study since pregnant women and patients who started anti-thrombotic
265 therapy or added immunosuppressant after starting HCQ administration were excluded
266 in this study.

267 In conclusion, S100 protein levels have been reported to correlate with the
268 pathogenesis of organ involvement in SLE patients. Our findings suggest that HCQ
269 improves organ involvement in SLE through modulation of S100 protein levels,
270 especially in patients with renal or skin involvement. Further investigation is needed to
271 clarify the mechanisms underlying S100 protein modulation by HCQ.

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275 **Declaration of conflicting interests**

276 The authors declare that there is no conflict of interest.

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364

365 Figure legend.

366 Figure 1. Association between serum S100 protein levels and renal involvement in SLE
367 patients prior to HCQ treatment.
368 S100A8 and S100A9 serum levels were increased in SLE patients with renal
369 involvement.
370 For statistical analyses * $p < 0.05$, ** $p < 0.01$
371 P value: Wilcoxon rank sum test

372 Figure 2. The change of serum level S100 proteins by HCQ treatment.
373 Serum S100A8 and S100A9 levels at baseline were compared with levels after 3 and 6
374 months of HCQ treatment with Bonferroni correction. Serum S100A8 and S100A9
375 levels significantly decreasing during HCQ treatment in SLE patients.
376 For statistical analyses **** $p < 0.0001$, ** $p < 0.015$, NS: Not significant P value: t
377 test

378 Figure 3. Degree of change in serum S100 protein levels in SLE patients with renal
379 involvement (n=17) or without renal involvement (n=20).
380 The magnitude of changes in serum S100A8 and S100A9 levels in SLE patients with
381 renal involvement were significantly higher than in patients without renal involvement.
382 For statistical analyses * $p < 0.05$.
383 P value: Wilcoxon rank sum test

384 Figure 4. Changes of serum S100 protein levels in CLASI responders (n=17) and
385 CLASI non-responders (n=8).
386 Serum S100A8 and S100A9 levels at baseline were compared with levels after 3
387 months of HCQ treatment in patients classified into CLASI responders and CLASI non-
388 responders. Decreases in serum S100A8 and S100A9 levels were much more apparent
389 in CLASI responders.
390 For statistical analyses * $p < 0.05$, ** $p < 0.01$. Wilcoxon signed-rank test

391 Figure 5. Changes in serum S100 protein levels during HCQ treatment in SLE patients
392 positive (n=6) or negative (n=31) for lupus anticoagulant(LAC).
393 Changes in serum S100A9 levels in LAC-positive SLE patients were significantly
394 lower than in LAC-negative patients.
395 For statistical analyses * $p < 0.05$.
396 lupus anticoagulant: LAC.
397 P value: Wilcoxon rank sum test

Table 1 Characteristics of SLE patients enrolled in this study

Characteristics	n=37
Female, no(%)	33(89)
Age, years, mean±SD	40.7±13.7
Disease duration, years, mean±SD	14.4±11.5
Complication	
APS	4 (11)
Past involvement	
Skin involvement	31 (84)
Renal involvement	17 (46)
Duration of CR free, years	5.2±3.3
NPSLE	2 (5)
Concomitant immunosuppressive treatments	
Prednisone	32 (86)
No.(%)	4.8 (1-10)
Median Dosage, mg/day (range)	24 (62)
Other immunosuppressant* ¹	
Tacrolimus	13 (35)
Mycophenolate mofetil	6 (16)
Cyclosporine A	2 (5)
Mizoribine	1 (3)
Methotrexate	1 (3)
Azathioprine	1 (3)
Anti-thrombotic therapy	
Anti-thrombotic therapy* ²	18 (49)
Antiplatelet agent	12 (32)
Anticoagulant agent	7 (19)
Positive rate of autoantibody	
Anti-Sm	8 (22)
Anti-RNP	17 (46)
Anti-SS-A	18 (49)

Anti-SS-B	7 (19)
Lupus anticoagulant	6 (16)
Anti-cardiolipin	15 (41)
Anti-β2GPI	2 (5)
<hr/>	
Disease activity	
<hr/>	
SELENA-SLEDAI score	3.7±1.9
CLASI activity score	3.2±3.2 (n=25)
CLASI damage score	0.6±1.3 (n=25)
anti-dsDNA positive, no(%) * ³	14 (38)
anti-dsDNA (IU/mL)	16.2±19.1
C3 (mg/dL)	81.5±21.1
C4 (mg/dL)	16.4±8.1
CH50 (U/mL)	35.6±8.9
low complement, no(%) * ⁴	18 (49)
White Blood Cell (/μL)	4978.1±1639.5
Lymphocytes (/μL)	1183.9±649.7
Platelet (× 10 ⁴ /μL)	22.1±7.2
<hr/>	

*¹ Two patients received multiple immunosuppressants.

*² One patient received antiplatelet agent and anticoagulant agent.

*³ Anti-dsDNA positive means anti ds-DNA titer increases over 12 IU/ml

*⁴ Low complement means any of C3, C4 and CH50 decreases to less 68mg/dl, less 12mg/dl, 30U/ml.

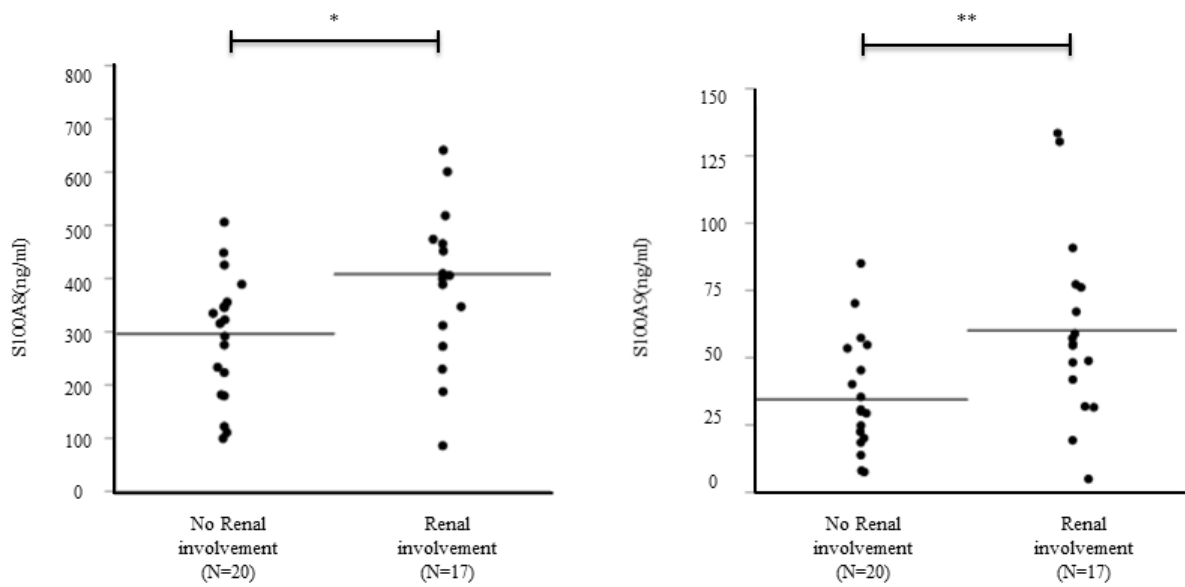


Figure 1

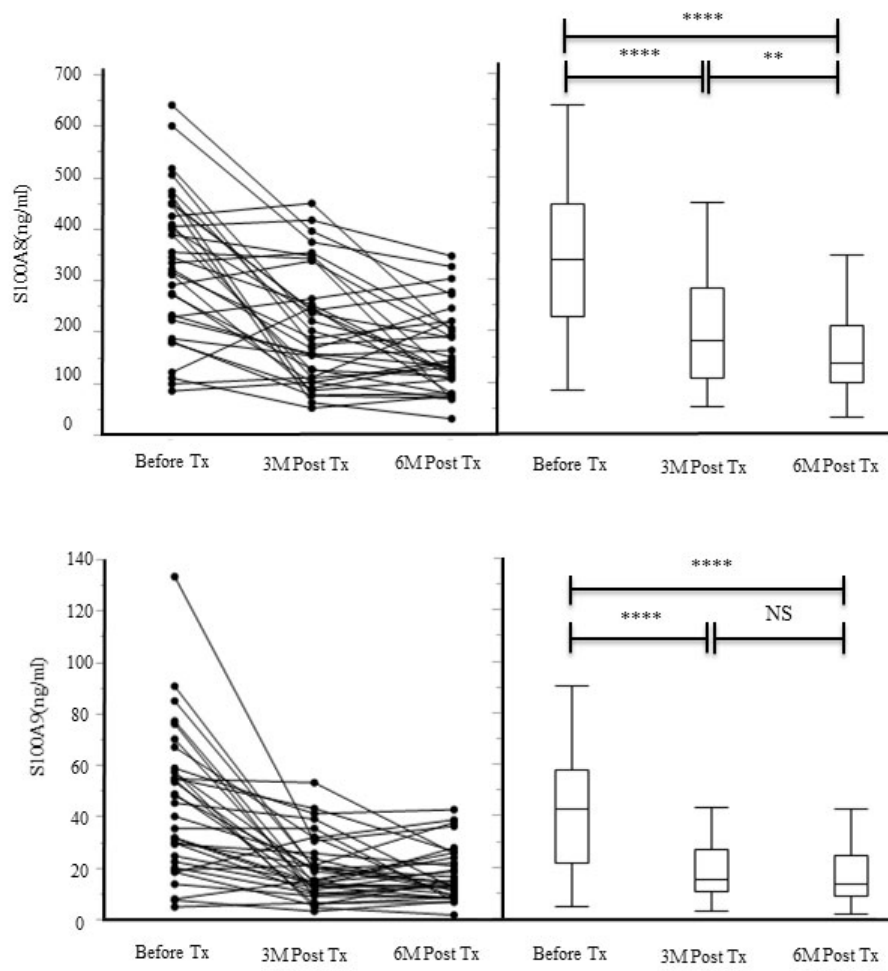


Figure 2

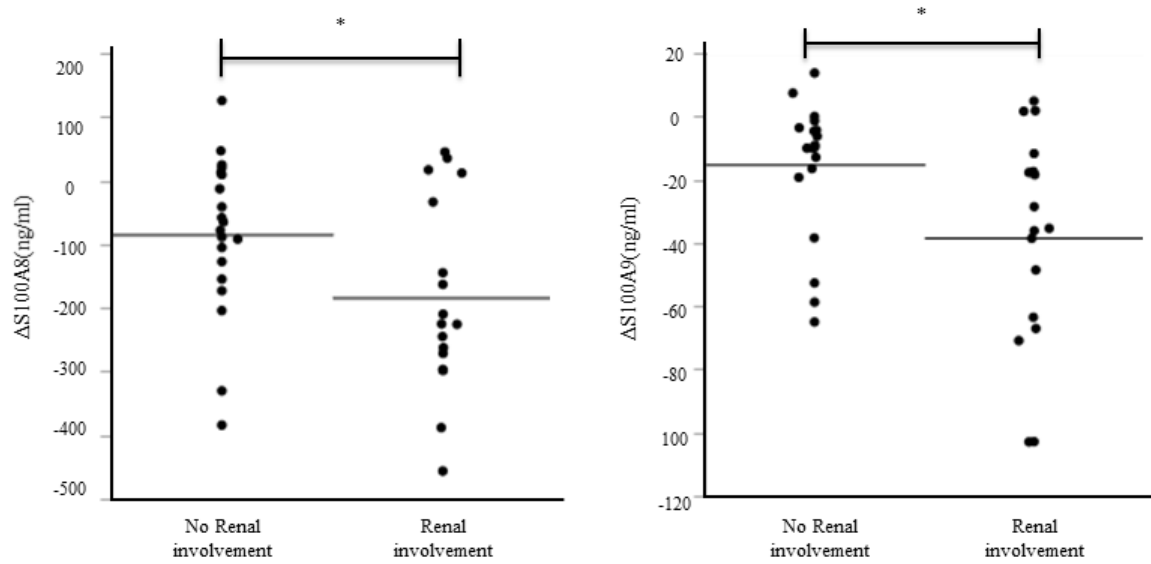


Figure 3

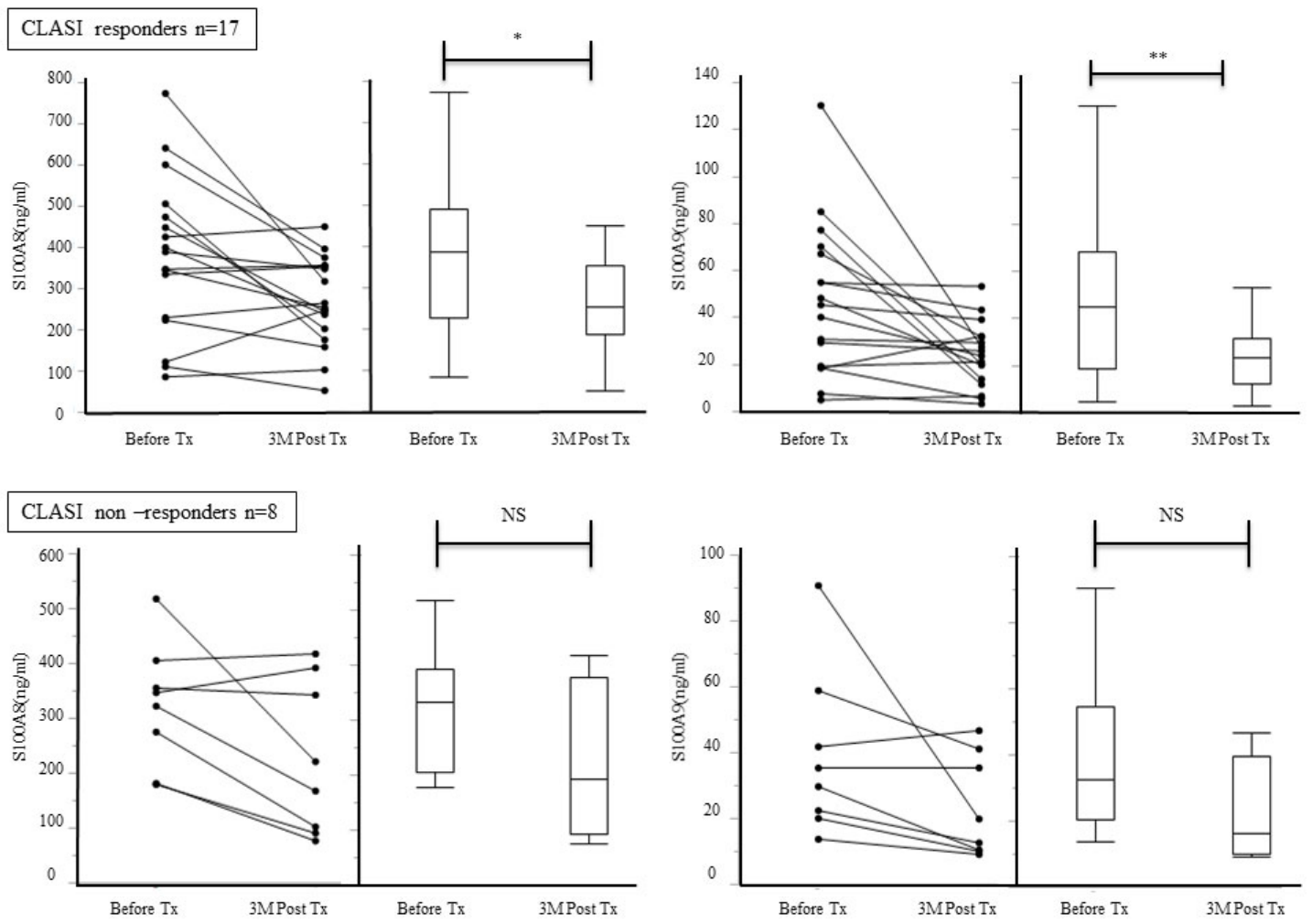


Figure 4

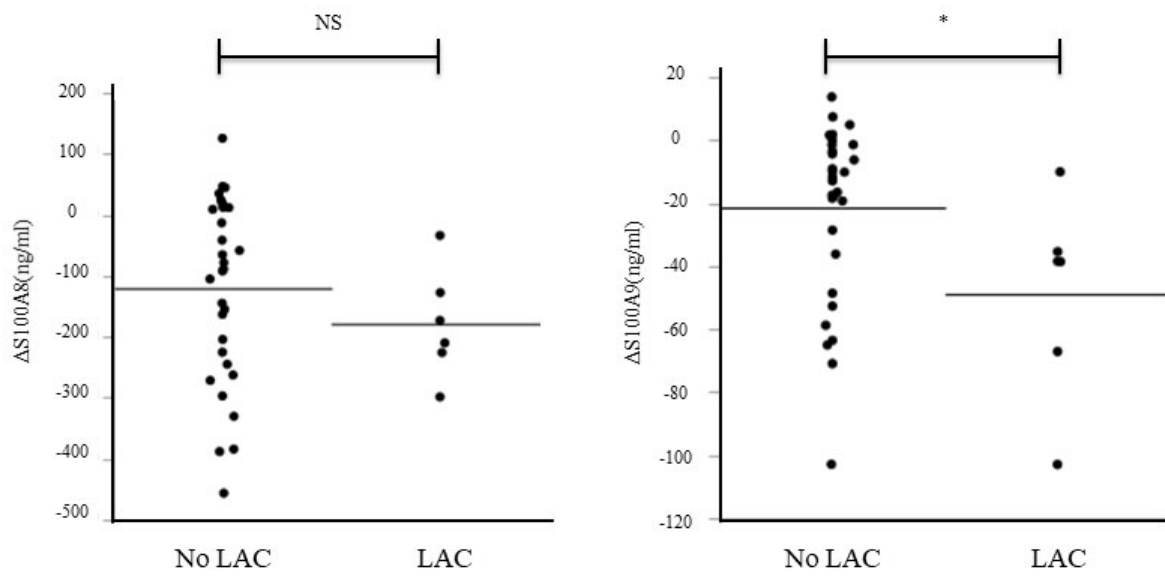
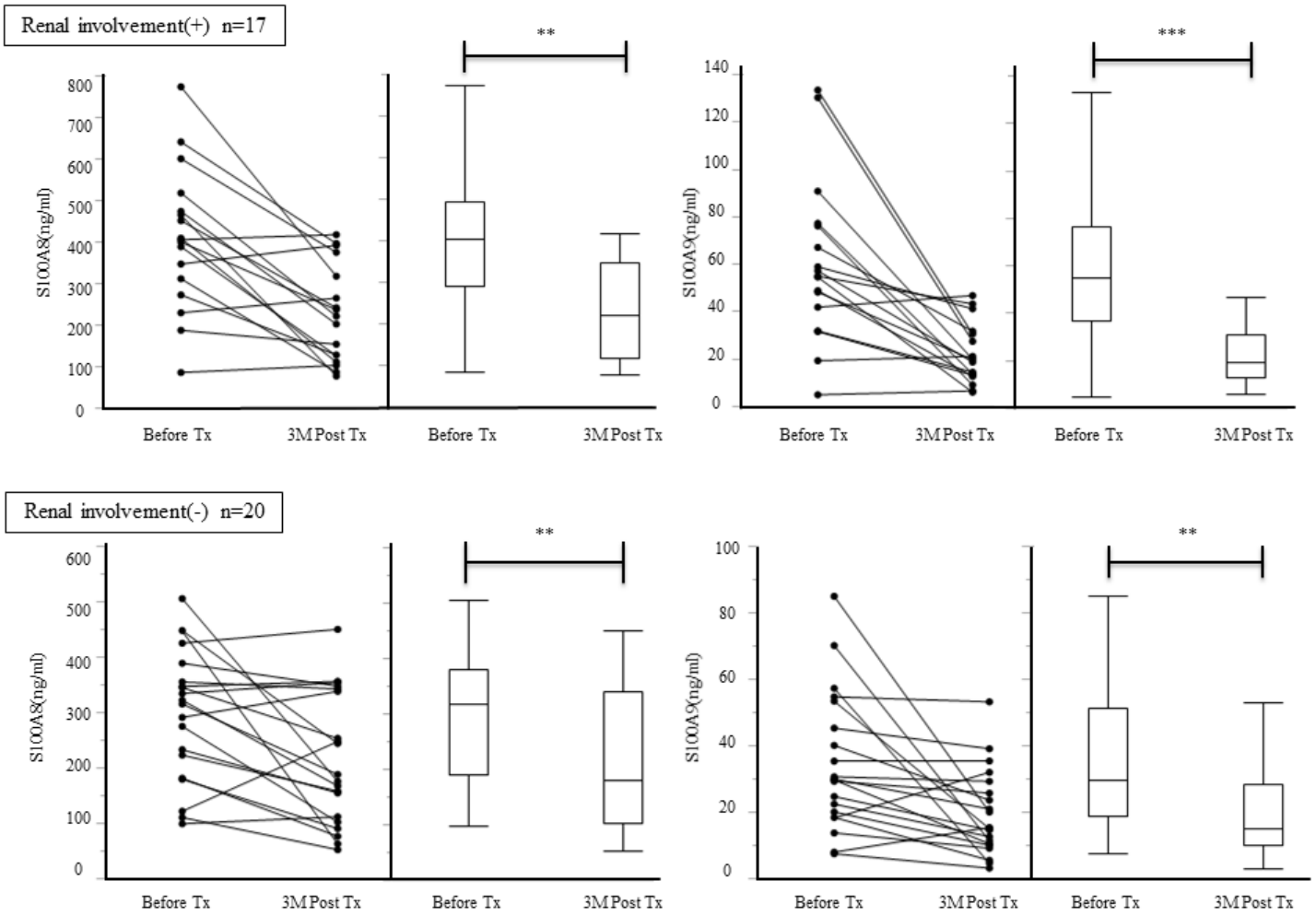
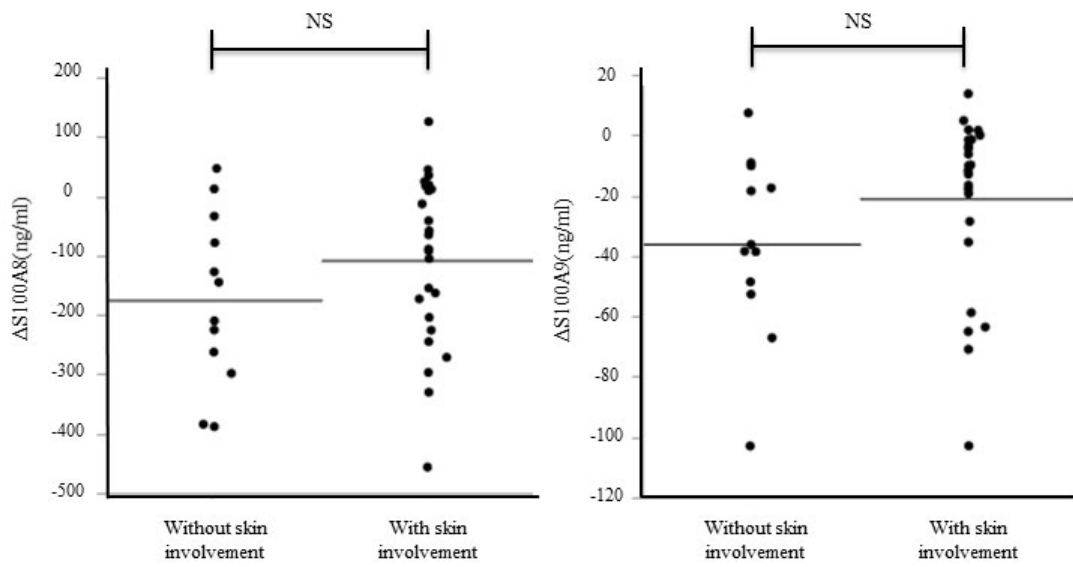


Figure 5

Supplementary Material Figure S1

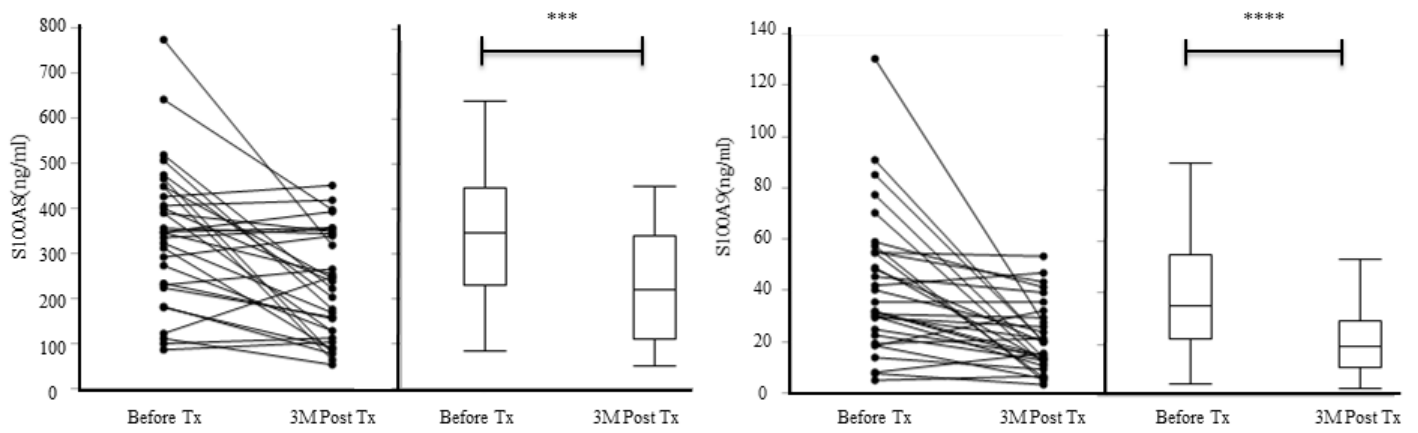


Supplementary Material Figure S2



Supplementary Material Figure S3

LAC -negative (n=33)



LAC -positive (n=6)

