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1 **Ultra-sensitive serum interferon- α quantification during SLE remission identifies**
2 **patients at risk for relapse.**

3 Running title: digital IFN- α predicts lupus flare

4 Alexis Mathian, MD, PhD*¹ Suzanne Mouries-Martin, MD, MSc*² Karim Dorgham, PhD*³
5 Hervé Devilliers, MD, PhD⁴ Hans Yssel, PhD³ Laura Garrido Castillo, MSc³ Fleur Cohen-
6 Aubart, MD, PhD¹ Julien Haroche, MD, PhD¹ Miguel Hie, MD, MSc⁵ Marc Pineton de
7 Chambrun, MD, MSc⁵ Makoto Miyara, MD, PhD³ Micheline Pha, MD, MSc⁵ Flore
8 Rozenberg, MD, PhD⁶ Guy Gorochoy, MD, PhD^{†3}, Zahir Amoura, MD, MSc^{†1}

9 *These authors contributed equally to this work.

10 †These authors contributed equally to this work.

11

12 ¹Sorbonne Université, Assistance Publique–Hôpitaux de Paris, Groupement Hospitalier Pitié–
13 Salpêtrière, French National Referral Center for Systemic Lupus Erythematosus,
14 Antiphospholipid Antibody Syndrome and Other Autoimmune Disorders, Service de
15 Médecine Interne 2, Institut E3M, Inserm UMRS, Centre d’Immunologie et des Maladies
16 Infectieuses (CIMI-Paris), Paris, France ;

17 ²Centre Hospitalier Universitaire de Dijon, Hôpital François-Mitterrand, service de médecine
18 interne et maladies systémiques (médecine interne 2), Dijon, France ;

19 ³Sorbonne Université, Inserm UMRS, Centre d’Immunologie et des Maladies Infectieuses
20 (CIMI-Paris), Assistance Publique–Hôpitaux de Paris, Groupement Hospitalier Pitié–
21 Salpêtrière, Département d’Immunologie, Paris, France ;

22 ⁴Centre Hospitalier Universitaire de Dijon, Hôpital François-Mitterrand, service de médecine
23 interne et maladies systémiques (médecine interne 2) et Centre d’Investigation Clinique,
24 Inserm CIC 1432, Dijon, France ;

25 ⁵Assistance Publique–Hôpitaux de Paris, Groupement Hospitalier Pitié–Salpêtrière, French
26 National Referral Center for Systemic Lupus Erythematosus, Antiphospholipid Antibody
27 Syndrome and Other Autoimmune Disorders, Service de Médecine Interne 2, Institut E3M,
28 Paris, France ;

29 ⁶Sorbonne Paris Cité, Assistance Publique–Hôpitaux de Paris, Hôpital Cochin, Service de
30 Virologie, Paris, France

31

32 **Correspondence to** Dr Alexis Mathian, Service de Médecine Interne 2, Institut E3M, Hôpital
33 de la Pitié–Salpêtrière, 47–83, boulevard de l’Hôpital, 75651 Paris Cedex 13, France.

34 Tel: +33 (0)1 42 17 80 25; Fax: +33 (0)1 42 16 58 04; Email: alexis.mathian@aphp.fr

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3

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8

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10

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12

1 **ABSTRACT**

2 **Objectives** Maintenance of remission has become central in the management of systemic
3 lupus erythematosus (SLE). The importance of interferon-alpha (IFN- α) in the pathogenesis
4 of SLE notwithstanding, its expression in remission has been poorly studied as yet. To study
5 its expression in remission and its prognostic value in the prediction of a disease relapse,
6 serum IFN- α levels were determined using an ultrasensitive single-molecule array (Simoa)
7 digital immunoassay which enables the measurement of cytokines at physiological
8 concentrations.

9 **Methods** A total of 254 SLE patients in remission, according to the Definition of Remission
10 in SLE (DORIS) classification, were included in the study. Serum IFN- α concentrations were
11 determined at baseline and patients were followed-up for one year. Lupus flares were defined
12 according to the SELENA-SLEDAI Flare Index, whereas the Kaplan-Meier analysis and Cox
13 regression analysis were used to estimate the time to relapse and to identify baseline factors
14 associated with time to relapse, respectively.

15 **Results** Of all patients in remission, 26% displayed abnormally high IFN- α serum levels that
16 were associated with the presence of antibodies specific for RNP, double stranded (ds)DNA
17 and Ro/SSA60, as well as young age. Importantly, elevated-baseline IFN- α serum levels and
18 remission duration were associated in an independent fashion, with shorter time to relapse,
19 while low serum levels of C3 and anti-dsDNA Abs were not.

20 **Conclusion** Direct serum IFN- α assessment with highly sensitive digital immunoassay permit
21 clinicians to identify a subgroup of SLE patients clinically in remission, but at higher risk of
22 relapse.

23

1 **Key messages**

- 2 - serum IFN- α levels have been poorly studied in SLE patients in remission so far;
- 3 - the single-molecule array (Simoa) assay is an ultrasensitive assay that quantifies directly
- 4 IFN- α at attomolar concentrations;
- 5 - a quarter of SLE patients in remission display elevated serum IFN- α levels;
- 6 - elevated serum IFN- α levels constitute an independent predictive biomarker of lupus flare;
- 7 - adding serum IFN- α to the routine laboratory assessments in patient in remission could help
- 8 clinicians to identify a subgroup of SLE patients at higher risk of relapse.

9

1 INTRODUCTION

2 Systemic lupus erythematosus (SLE) is a chronic autoimmune disease of unknown
3 incompletely known etiology characterized by the presence of anti-nuclear autoantibodies and
4 inflammation in a wide spectrum of organs.[1] Despite the improvement of SLE prognosis in
5 the last decades, SLE patients are still at high risk of disease-related complications and
6 premature death.[2, 3] Several authors have shown that remission defined as the elimination
7 of disease activity predicted a better disease outcome with a lower burden of damage and a
8 lower risk of relapse.[4, 5] Thus, remission achievement and its maintenance have become
9 central in the management of SLE patients.[6, 7] Many definitions have been used to better
10 characterize remission states, but a single one has recently made consensus, the DORIS
11 (Definitions Of Remission In SLE). For defining remission, the DORIS uses a clinical index,
12 such as the clinical SLE disease activity index (SLEDAI)=0 with routine laboratory
13 assessments including anti-double stranded DNA (anti-dsDNA) antibodies (Abs) and
14 complement.[7] Serologically active patients, i.e. patients with an increase in serum anti-
15 dsDNA Abs and/or a complement consumption, are much more likely to experience
16 subsequent flares, as compared to those who are serologically inactive.[7] However, these two
17 markers fail to reliably predict a lupus flare.[8, 9] Consequently, no consensus has yet been
18 reached by the DORIS working group as to a reliable definition that enables the distinction of
19 patients who are serologically active from those who are serologically inactive.[7]

20 Many authors consider the dysregulation of interferons (IFNs), especially IFN-alpha
21 (IFN- α), to be a central cause of the immunological abnormalities observed in SLE.[10-16]
22 Transcriptome analysis using microarray technology revealed up-regulation of numerous
23 IFN-stimulated genes (ISGs) in SLE patients' peripheral blood mononuclear cells,
24 constituting an overall "IFN signature".[17, 18] Many reports showed patients' elevated

1 serum-IFN- α levels to be associated with SLE activity and severity, suggesting that
2 monitoring this cytokine might help physicians to better evaluate disease activity,[17-35]
3 although, unexpectedly, some clinically inactive patients have persistently elevated serum
4 IFN- α levels.[19-23, 25, 35] Since serum concentrations of IFN- α are usually very low and
5 often not detectable by classic immunoassays, monitoring the expression of ISGs is used to
6 evaluate IFN serological activity. The resulting “IFN scores” are calculated based on the
7 expression of several representative ISGs.[29, 36, 37] However, the low availability and high
8 complexity of transcriptome-microarray technology implies that IFN scores are not
9 standardized and therefore cannot be easily used in clinical practice. At present, IFN- α
10 overexpression has been poorly studied in patients in remission. The single-molecule array
11 (Simoa) assay, or digital immunoassay, is an ultrasensitive assay based on enumeration of
12 individual enzyme-labeled immunocomplexes of proteins captured on beads in single-
13 molecule arrays. It enables direct IFN- α quantification at attomolar concentrations,[35, 38-40]
14 corresponding to a 5,000-fold-increased sensitivity over classic ELISAs. We and others have
15 shown that at physiological concentrations, the digital immunoassay is as sensitive as ISG
16 expression as a means to quantify IFN- α levels and simpler to perform and standardize.[35,
17 40] We thus conducted a study to determine the magnitude of serum IFN- α concentrations in
18 SLE patients in remission using the DORIS-validated criteria.[7] Additionally, in order to
19 improve the definition of remission, we determined the clinical and biological features
20 associated with elevated IFN- α levels in patients experiencing a remission and we
21 investigated whether high serum IFN- α levels at baseline in patients in remission were
22 predictive of a flare in the ensuing year.

23

1 PATIENTS, MATERIALS AND METHODS

2 Study design and patients

3 We conducted this longitudinal study between September, 2014 and September 2017 at the
4 National Referral Center for SLE, Paris, France. Serum samples were obtained at day 0 (=
5 baseline) from consecutive patients diagnosed with SLE according to the 1997 American
6 College of Rheumatology criteria for SLE classification, regardless of the activity of the
7 disease.[41] Exclusion criteria were: 1) known or suspected infection or malignancy on the
8 day blood was drawn; 2) increased Hydroxychloroquine (HCQ), prednisone and/or
9 immunosuppressant (IS) during the 4 weeks preceding day 0. SLE clinical characteristics (see
10 [online supplement](#)), SELENA-SLEDAI,[42-44] SLEDAI-2K,[45] class of lupus nephritis
11 according to ISN/RPS-2003,[46] and therapeutic regimen were recorded on day 0. Lupus
12 flares were defined according to the SELENA-SLEDAI Flare Index.[43, 44] The term
13 "clinical" SLEDAI (cSLEDAI) refers to symptoms, signs and routine laboratory testing and
14 disregards only the points that can be given for the presence of anti-dsDNA Abs and/or low
15 complement.[7] Five exclusive disease activity statuses were defined, according to the
16 DORIS [7] and following Wilhelm et al. [47] and Ugarte-Gil et al. [48] without physician
17 global assessment (PGA) and serum C4 analysis:

- 18 - *Complete remission off treatment*: cSELENA-SLEDAI=0, no corticosteroids, no
19 immunosuppressant (IS), no anti-dsDNA Abs and no C3 decrease;
- 20 - *Clinical remission off treatment*: cSELENA-SLEDAI=0, no corticosteroids, no IS.
21 Anti-dsDNA Abs and/or C3 decrease present;
- 22 - *Complete remission on treatment*: cSELENA-SLEDAI=0, prednisone 1-5mg/day,
23 IS allowed, no anti-dsDNA Abs and no C3 decrease;
- 24 - *Clinical remission on treatment*: cSELENA-SLEDAI=0, prednisone 1-5mg/day, IS

1 allowed. Anti-dsDNA Abs and/or C3 decrease present;
2 - *Not in remission*: cSELENA-SLEDAI>0 and/or prednisone >5mg/day.
3 HCQ was allowed in all groups. Duration of remission was recorded. Over a consecutive 5-
4 year period of remission, the patient was considered in “prolonged remission”, as proposed by
5 Steiman et al. [49] and treated in the statistical analysis as a 5-year remission. Alternatively,
6 patients not in remission but fulfilling Lupus Low Disease Activity State (LLDAS) [48, 50]
7 were included in the “*LLDAS without remission*” subgroup ([see online supplement](#)). Sera
8 from age- and gender-matched healthy donors (n=68) were collected (Établissement Français
9 du Sang, Île-de-France, Pitié–Salpêtrière Hospital, Paris, France) during the same time period.
10 The local Ethics Committee of the Pitié-Salpêtrière Hospital approved this study (file N°
11 30052012) and informed consent was obtained from all participants.

12

13 **Patient follow-up**

14 Patients in remission at day 0 were followed for one year ([see online supplement](#)).

15

16 **IFN- α digital immunoassay**

17 Serum IFN- α concentrations, expressed in fg/mL, were determined at day 0 with digital
18 immunoassay technology (IFN- α Reagent Kit, Quanterix SimoaTM, Lexington, MA, USA),
19 based on a 3-step protocol ([see online supplement](#)) using the HD-1 Analyzer
20 (QuanterixTM).[38] The IFN- α digital ELISA positivity threshold was 136 fg/mL, which is 3
21 SDs above the mean serum IFN- α concentration calculated from the sera samples from the 68
22 healthy blood donors.

23

24 **Statistical analysis**

1 Statistical analyses were performed using GraphPad Prism, v5.0 software (GraphPad
2 Software, San Diego, CA, USA) and SAS 9.4 software (Cary, NC, USA) ([see online
3 supplement](#)).

4

1 RESULTS

2

3 Patient characteristics and distribution according to disease activity

4 A total of 407 patients were included in the study, with 254 patients in remission (i.e.
5 cSELENA-SLEDAI=0 and prednisone equal or less than 5 mg/day) and 153 not in remission.
6 Patients' baseline characteristics are described in **Tables 1 and 2**. From the 254 patients in
7 remission, 86 (33.9%) were in complete remission off treatment, 59 (23.2%) in complete
8 remission on treatment, 47 (18.5%) in clinical remission off treatment and 62 (24.4%) in
9 clinical remission on treatment. Sixty-seven (26.3%) patients were in remission for less than 1
10 year, 101 (39.8%) for 1 to 5 years, and 86 (33.9%) for more than 5 years. The median (range)
11 of remission duration was 5 years (0-5) for patients in complete remission off treatment, 1.6
12 years (0-5) for patients in complete remission on treatment, 4.2 years (0.1-5) for clinical
13 remission off treatment and 1.6 years (0-5) for clinical remission on treatment. Among the
14 153 patients not in remission, 19 (12.4%) fulfilled the LLDAS criteria, defining the "LLDAS
15 without remission" subgroup.

16

17 Serum IFN- α in remission and LLDAS

18 The serum IFN- α levels according to the different states of remission are presented in
19 **Table 2**. A total of 26.0% of patients in remission had still detectable serum IFN- α levels
20 exceeding the positivity threshold. Patients in "clinical remission" had the highest rate of
21 abnormal serum IFN- α : 41.9% for "clinical remission on treatment" and 38.3% for "clinical
22 remission off treatment" [versus 16.3% for "complete remission off treatment", $p = 0.0007$
23 and 0.006, respectively]. Among patients in remission, patients in "clinical remission on
24 treatment" had the highest concentration of IFN- α with a median (quartiles) of 109 fg/mL

1 (12-378) [versus 11 fg/mL (0-81) in patients in “complete remission off treatment”,
2 p=0.0002]. Alternatively, we assessed serum IFN- α in different states of remission according
3 to the definition of Zen et al., a definition of remission in which the role of corticosteroids is
4 highlighted in comparison to immunosuppressive treatment.[5] As shown in **supplemental**
5 **table 2**, we found similar results: patients in “clinical remission on or off corticosteroids” had
6 the highest rate of abnormal serum IFN- α levels, as compared to those of patients in
7 “complete remission”. Thus, a significant number of patients in remission had increased IFN-
8 α levels in their serum. Eight (42.1%) patients of the “LLDAS without remission” subgroup
9 displayed abnormal IFN- α serum concentration (OR 3.7 [95% CI 1.3-11.0] and a mean IFN- α
10 level of 111 fg/mL [0-1,647], p=0.02 and p=0.07, respectively, as compared with patients in
11 “complete remission off treatment”).

12

13 **Parameters SLE characteristics associated with abnormal serum IFN- α levels in** 14 **remission**

15 Elevated IFN- α serum levels in remission were significantly associated, in multivariable
16 analysis, with the presence of serum Abs specific for RNP, dsDNA and Ro/SSA60, young age
17 and lower granulocyte counts (**Table 3**). In contrast, disease and remission duration,
18 prednisone intake, immunosuppressant therapy, HCQ, and low C3 serum levels did not show
19 a significant association with IFN- α serum levels.

20

21 **Elevated IFN- α serum levels in SLE patients in remission predicts a lupus flare**

22 Of the 254 patients in remission at day 0, 250 were followed for one year. Twenty-four
23 (9.6%) patients experienced a flare. The median (range) time of the flare occurrence was 141
24 (25-349) days. The type and severity of the flares are reported in **supplemental table 3**. A total

1 of 37.5% of the flares were severe. The most frequent type of relapses was arthritis (n=15)
2 followed by cutaneous flare (n=7) and serositis (n=4). Unadjusted cox regression showed a
3 significantly higher risk of relapse in patients who displayed at baseline elevated IFN- α (HR
4 5.5 [95% CI 2.4-12.5], $p < 0.0001$) or decreased C3 (HR 3.7 [95% CI 1.6-9.1], $p = 0.003$) serum
5 levels, respectively, but not in patients who had a positive Farr assay (HR 1.5 [95% CI 0.7-
6 3.4], $p = 0.3$) (Figure 1). The highest concentrations of IFN- α at baseline were associated with
7 the greatest frequencies of relapse (supplemental figure 1). Other factors at baseline
8 associated with the risk of relapse were prednisone intake 1-5 mg/day (HR 3.2 [95% CI 1.4-
9 7.5], $p = 0.007$), positive anti-RNP Abs (HR 3.1 [1.4-7.0], $p = 0.006$), age < 40 years (HR 0.4
10 [95% CI 0.2-0.9], $p = 0.02$) and disease duration < 10 years (HR 0.2 [95% CI 0.1-0.6],
11 $p = 0.003$). The remission duration was a protective factor for the probability of flare (HR 0.6
12 [95% CI 0.5-0.8] for each consecutive year of remission completed, $p = 0.0002$).

13 Finally, in multivariable analysis, the factors independently associated with the risk of
14 flare were abnormal serum IFN- α levels at baseline (HR 4.0 [95% CI 1.7-9.6], $p = 0.002$) and
15 remission duration (HR 0.7 [95% CI 0.5-0.9], $p = 0.02$, for each year in remission). Low C3
16 (HR 2.4 [95% CI 0.9-6.2], $p = 0.07$) and prednisone intake (HR 2.4 [95% CI 0.9-5.9], $p = 0.06$)
17 were also kept in the model as-associated with a higher risk of flare but these associations
18 were not statistically significant. We performed a sensitivity analysis using a lower
19 concentration of IFN- α as the threshold of elevated IFN- α levels and found similar results
20 (data not shown).

21 Additionally, the risks of relapse according to combined SLE biomarkers (low C3
22 complement, positive Farr assay and positive IFN- α) were independently analysed with a
23 proportional risk cox model (supplemental table 4). In this model, isolated positive IFN- α was

1 a predictive factor of lupus flare (HR of 5.5 [95% CI 1.7-18.1], $p = 0.005$). In contrast,
2 isolated positive Farr assay and isolated low C3 were not predictive of lupus flare.

3

4 **Digital immunoassay, complement C3 and Farr assay performances to predict a flare**

5 The time-dependent ROC area under the curve (AUC) (**Figure 2**) estimates for the IFN- α
6 digital immunoassay to predict a flare was 0.73, better than that of anti-dsDNA Abs (0.60,
7 $p=0.055$), and complement C3 serum levels (0.56, $p=0.2$). The time-dependent ROC AUCs
8 for the three biomarkers were highest around day 90.

1 DISCUSSION

2 The significance of monitoring IFN- α in SLE for the assessment of remission and the risk to
3 develop flares has been poorly documented as yet. In the present study, we show that a
4 significant proportion of patients despite being in remission have high serum IFN- α levels that
5 were found to independently predict the risk of subsequent SLE flares, thereby emphasizing
6 the interest of measuring the production of this cytokine to monitor the course of disease.

7 Earlier publications have reported that between 3 and 32% of SLE patients without active
8 disease present elevated IFN- α serum levels depending on the type of assay used to determine
9 the presence of this cytokine.[19-23, 25, 35] However, because different definitions of disease
10 inactivity were used in these studies, the results cannot directly be compared with our study
11 that takes into account the recently formulated consensual definition of disease remission.[5,
12 7, 47] Contrary to previous report,[51] we noted that serum concentrations of IFN- α above the
13 positive threshold values varied between different remission subgroups. Patients in clinical
14 remission, i.e. with serological activity, presented more frequently elevated serum levels of
15 IFN- α than patients in complete remission. The presence of serum Abs specific for RNP,
16 dsDNA and Ro/SSA60 were found to be independently associated with the magnitude of
17 serum IFN- α levels. These associations have previously been shown in SLE [29-31, 52-57]
18 but, to the best of our knowledge, never been reported for patients in remission. These data
19 are in accordance with those from previous studies that showed a key role of DNA/RNA-
20 associated immune complexes through the activation of TLR7 and TLR9 for the induction of
21 type I IFN production.[58, 59]

22 The DORIS group has recently agreed that the subsequent occurrence of flares was
23 among the most appropriate outcome variables for defining the prognostic value of remission
24 and furthermore suggested the inclusion of serological criteria for this definition.[7] Results

1 from previous reports on smaller SLE patient cohorts, based on the analysis of expression of
2 ISGs in peripheral blood cells by microarray, failed to demonstrate an association between
3 IFN scores and the longitudinal risk of relapse.[30, 31] It is to be noted that type I and type II
4 IFNs largely overlap in the genes that they control, making it difficult to distinguish the
5 signatures of IFN-gamma (IFN- γ) from IFN- α and IFN-beta (IFN- β). Indeed, the results from
6 a modular repertoire analysis have emphasized that IFN signatures in SLE are not restricted to
7 IFN- α , but also involve IFN- β and IFN- γ , thus underscoring the nonspecific nature of IFN
8 scores.[60] Therefore, in a disease like SLE in which IFN- α appears to be central in the
9 pathogenesis, it is likely that clinical activity better correlates with the values of IFN- α serum
10 levels, directly measured by a highly specific digital ELISA, than with the less specific
11 expression of IGSs. Moreover, the IFN scores are likely to be rather poorly sensitive to
12 changes in IFN- α serum concentrations and therefore of little use in monitoring the risk of
13 relapse of disease. The results of our study suggest that the measurement of additional
14 interferons, including IFN- β , - γ and - λ with ultrasensitive digital immunoassay, might help to
15 determine whether the expression of other IFN members also correlates with disease activity
16 or specific characteristics of the disease.[34, 61, 62] Nonetheless, our results corroborate
17 those from an earlier study showing the usefulness of monitoring expression levels of certain
18 IFN-regulated chemokines to predict future flares: serum levels of CXCL10, CCL2 and
19 CCL19 chemokines were found to be linked with SLE activity, and performed better than
20 other laboratory tests to predict a flare over the following year.[63] Yet, these chemokines can
21 be induced by other SLE-associated cytokines beyond IFN- α , such as IFN- γ and their
22 monitoring remains difficult in routine practice. As reported by others, we did not find an
23 association between the presence of anti-dsDNA Abs and the risk of flare. Not surprisingly,
24 the duration of remission was significantly associated with a poor risk of flare in the

1 following year, thus identifying a subgroup of patients in prolonged remission with less
2 relapsing risk and for whom clinical monitoring can probably be lightened.

3 The sustained presence of serum IFN- α in serum of SLE patients in remission may have
4 pathological consequences by itself. IFN- α overexpression could be an explanation of chronic
5 fatigue, depression and reduction of sleep secondary to the stimulation of dopamine
6 metabolism by IFN- α in the central neurological system.[64-66] IFN- α plays also a prominent
7 role in endothelial cell damage and up-regulation of the expression of scavenger receptors in
8 monocyte and macrophages, leading to increased lipid uptake and foam cell formation, a
9 process that may be at the origin of the accelerated atherosclerosis observed in SLE.[67-69]
10 These data suggest that for patients in clinical remission, return of IFN- α to normal values
11 could become one of the objectives of the treatment.

12 Our study has some limitations. It unfortunately lacks data on patients' ethnicity, which is
13 known to bias serum IFN- α levels.[55] We also used a definition of remission and LLDAS in
14 the absence of PGA that has not been assessed in our cohort. However, other teams have
15 already adapted the definition of remission and LLDAS arguing that in the SLE Response
16 Index, the SLEDAI is the variable with the highest impact on the definition of response and
17 that even without PGA, a modified definition of remission or LLDAS is still entirely
18 valid.[48, 70] **Finally, the serum IFN- α concentrations were not assessed during the
19 longitudinal part of the study. Therefore information regarding the intra-individual variability
20 over time of serum IFN- α levels is lacking. These important data will have to be studied in
21 new studies.**

22 In conclusion, our data confirm that a large number of SLE patients in remission display
23 an elevated serum IFN- α concentrations, especially in the presence of anti-dsDNA, anti-

1 ribonucleoprotein Abs (i.e. anti-Ro/SSA 60, anti-RNP), as well as young age. This
2 overexpression is an independent predictive biomarker of lupus flare in the following year.
3 Adding serum IFN- α measurement to the routine laboratory assessments in patient in
4 remission could help clinicians to identify a subgroup of SLE patients clinically in remission
5 but who still overexpress IFN- α and are at higher risk of relapse. These data suggest that the
6 return of normalcy of serum IFN- α could become one of the objectives of the treatment. Our
7 results must be validated in other independent cohorts.

8

9

10

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10

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1 **Figures legends**

2 **Figure 1. Baseline positive serum IFN- α levels and low complement C3 identify patients**

3 **with elevated risk for future SLE flares.** Kaplan-Meier curves. IFN- α = interferon alpha.

4 Serum IFN- α (positivity threshold of 136 fg/mL), anti-dsDNA Abs by Farr assay (cut-off

5 value: 9.0 IU/mL) and complement C3 levels (cut-off value: 0.78g/L) were assessed at day 0.

6 Kaplan-Meier plots show the percentage of patients who flared in any organ system. Vertical

7 tick marks along each curve represent patients who remained flare-free but did not have a full

8 year of clinical follow-up (censored data). Curves were compared using Log-Rank tests.

9 Crude Hazard Ratios (HR) were calculated using proportional risk cox model **A.** The red

10 dashed line represents the 64 patients with positive IFN- α serum level at day 0. The

11 continuous blue line represents the 186 patients with negative IFN- α serum level at day 0. **B.**

12 The red dashed line represents the 99 patients with positive Farr assay at day 0. The

13 continuous blue line represents the 151 patients with negative Farr assay at day 0. **C.** The red

14 dashed line represents the 28 patients with low C3 level at day 0. The continuous blue line

15 represents the 216 patients with normal C3 level at day 0.

16

17 **Figure 2. Cumulative time-dependent receiver operating characteristics (ROC) curves.**

18 IFN- α = interferon alpha; C3 = complement C3. Serum IFN- α digital immunoassay, anti-

19 dsDNA Abs by Farr assay and complement C3 levels were assessed at baseline (day 0). SLE

20 patients in remission were followed for one year. Lupus flares were defined using the

21 SELENA-SLEDAI Flare Index. Patients who had a flare in any organ system were recorded.

22 The diagnostic performances of the baseline serum-IFN- α digital immunoassay (red line),

23 Farr assay (blue line) and complement C3 (green line) to predict SLE flare were investigated

24 by computing cumulative time-dependent ROC curves, with lupus flares serving as the gold

1 standard for those analyses. The dashed line represents the upper and the lower 95%
2 confidence interval (CI). The cumulative time-dependent ROC area under the curve (AUC)
3 estimated to predict a flare are given and were compared based on Uno *et al.*[71]

1 **Table 1. Disease parameters–characteristics in SLE patients at baseline**

	Patients (N=407)
Women	365 (90)
Age, years, mean±SD	40.1±12.9
Disease duration, years, mean±SD	11.2±9.4
SELENA–SLEDAI score, median (range)	2 (0–41)
SELENA–SLEDAI score ≥4	131 (32)
Mild/moderate flare*	31 (8)
Severe flare*	76 (19)
Clinical involvement	
Fever	37 (9)
Weight loss or anorexia	21 (5)
Lymphadenopathy	25 (6)
Any constitutional signs	47 (12)
Active cutaneous lupus	48 (12)
Active lupus serositis	23 (6)
Active lupus arthritis	67 (16)
Active lupus nephropathy	31 (8)
proliferative nephropathy	15 (4)
membranous nephropathy	16 (4)
Active neuropsychiatric lupus	8 (2)
Cytopenia	34 (8)
Treatment regimen	
Hydroxychloroquine use	347 (85)
Prednisone use	205 (50)
Prednisone ≥10 mg/j	56 (14)
Immunosuppressive agent use†	96 (24)
Biological tests	

Positive Farr test	198 (49)
Positive anti-RNP Abs	118 (29)
Positive anti-Sm Abs	51 (13)
Positive anti-Ro/SSA 52 Abs	97 (24)
Positive anti-Ro/SSA 60 Abs	151 (37)
Positive anti-La/SSB Abs	42 (10)
Low C3	98/398‡ (25)
Positive IFN- α	163 (40)

1 Values are expressed as n (%), unless stated otherwise.

2 * Defined using SELENA flare index [43, 44]

3 † Excluding antimalarials and prednisone. Immunosuppressant therapy was mycophenolate mofetil (MMF) for 43 (44%)
4 patients, methotrexate (MTX) for 39 (41%), azathioprine for 13 (14%) and cyclophosphamide for 1 (1%). Two patients were
5 receiving calcineurin inhibitor in addition to MMF and three patients were receiving belimumab in addition to MTX.

6 ‡ Positive assay/number of patients assessed.

7 SD, standard deviation; SELENA-SLEDAI, Safety of Estrogens in Lupus Erythematosus: National Assessment version of the
8 Systemic Lupus Erythematosus Disease Activity Index.

9

10

Table 2. Serum IFN- α levels at baseline in SLE patients in remission or not in remission (according to Wilhelm et al. [47] modified).

	Abnormal serum IFN-α level	Odds Ratio [95% CI]	p value^a	Serum IFN-α fg/mL, median (Q1, Q3)	p value^b
Remission (n=254)	66 (26.0)				
complete remission off treatment (n=86)*	14 (16.3)	1 [Ref.]	Ref.	11 (0-81)	Ref.
complete remission on treatment (n=59)	8 (13.6)	0.8 [0.3-2.1]	NS	15 (0-51)	NS
clinical remission off treatment (n=47)	18 (38.3)	3.2 [1.4-7.2]	0.006	18 (0-314)	NS
clinical remission on treatment (n=62)	26 (41.9)	3.7 [1.7-8.0]	0.0007	109 (12-378)	0.0002
Not in remission (n=153)	97 (63.4)				
prednisone >5 mg/day (n=33)	10 (30.3)	2.2 [0.9-5.7]	NS	78 (0-296)	0.04
cSLEDAI >0 (n=74)	51 (68.9)	11.4 [5.4-24.3]	<0.0001	519 (68-3,087)	<0.0001
prednisone >5 mg/day and cSLEDAI >0 (n=46)	36 (78.3)	18.5 [7.5-45.8]	<0.0001	2,054 (455-8,751)	<0.0001

Values are expressed as n (%), unless stated otherwise.

* reference group for statistical analysis

^a bivariable analysis using Mann–Whitney U-test compared to “complete remission off treatment” as the reference.

^b bivariable comparison using Fisher’s exact test compared with patients in “complete remission off treatment” as the reference.

NS, Non Significant; IFN- α , interferon alpha; fg/mL, femtogram/milliliter; cSLEDAI, clinicalSLEDAI; SD, standard deviation.

Table 3. Parameters-Disease characteristics associated with serum IFN- α levels at baseline in SLE patients in remission

	IFN- α		Bivariable p value ^a	Multivariable p value ^b	Adjusted odds ratio ^b [95% CI]
	Normal	elevated			
	N=188	N=66			
Women	165 (88)	61 (92)	0.30	ND	
Age, years, mean \pm SD	44.0 \pm 13.3	37.5 \pm 10.8	<0.001	0.002	0.96 [0.93-0.98]
Disease duration, years, mean \pm SD	13.4 \pm 10.0	10.4 \pm 8.3	0.029	NS	
Remission duration, years, median (Q1-Q3)	3.4 (1.1-5.0)	1.8 (0.5-5.0)	0.024	NS	
Hydroxychloroquine use	165 (88)	63 (95)	0.08	ND	
Prednisone use	72 (38)	28 (42)	0.55	ND	
Prednisone use, mg/d, median (Q1-Q3)	0 (0-5)	0 (0-5)	0.65	ND	
Immunosuppressive agent use [†]	31 (16)	15 (23)	0.26	ND	
Positive Farr test	59 (31)	42 (64)	<0.001	0.02	2.3 [1.1-4.6]
Positive anti-RNP Abs	21 (11)	29 (44)	<0.001	0.0002	4.6 [2.1-10.2]
Positive anti-Ro/SSA 52 Abs	34 (18)	22 (33)	0.01	NS	
Positive anti-Ro/SSA 60 Abs	52 (28)	31 (47)	0.004	0.01	2.6 [1.3-5.4]
Positive anti-La/SSB Abs	14 (7)	8 (12)	0.25	ND	
Positive anti-Sm Abs	5 (3)	13 (20)	<0.001	NS	
Low C3	14/184 [‡] (8)	16 (25)	<0.001	NS	
Lymphocytes, G/L, median (Q1-Q3)	1.5 (1.2-2.1)	1.2 (0.9-1.5)	0.0001	NS	
Granulocytes, G/L, median (Q1-Q3)	3.9 (2.9-5.5)	3.1 (2.4-3.7)	<0.001	0.001	0.7 [0.5-0.8]
Thrombocytes, G/L, median (Q1-Q3)	249 (215-284)	232 (205-282)	0.5	ND	
Haemoglobin, g/dL, median (Q1-Q3)	13.3 (12.6-14.1)	12.6 (11.8-13.7)	<0.001	NS	

Values are expressed as n (%), unless stated otherwise.

† Excluding antimalarials and prednisone

‡Positive assay/number of patients assessed

^a estimated by Mann-Whitney test, Fisher's exact or χ^2 test

^b evaluated by multivariable logistic regression.

SD, standard deviation; ND, Not Done; NS: Non Significant.

Figure 1

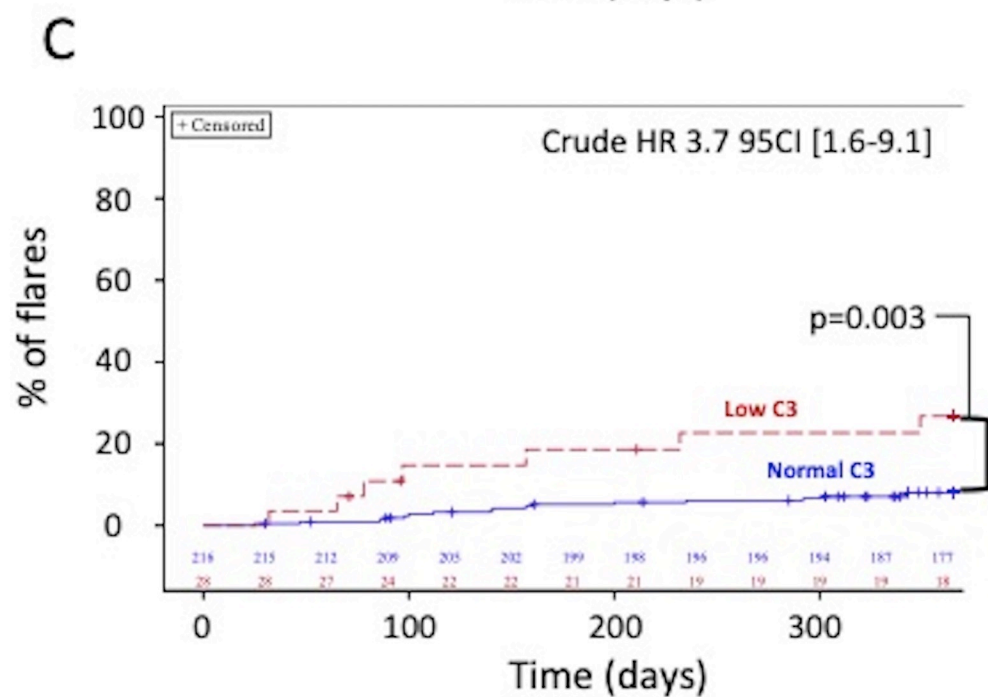
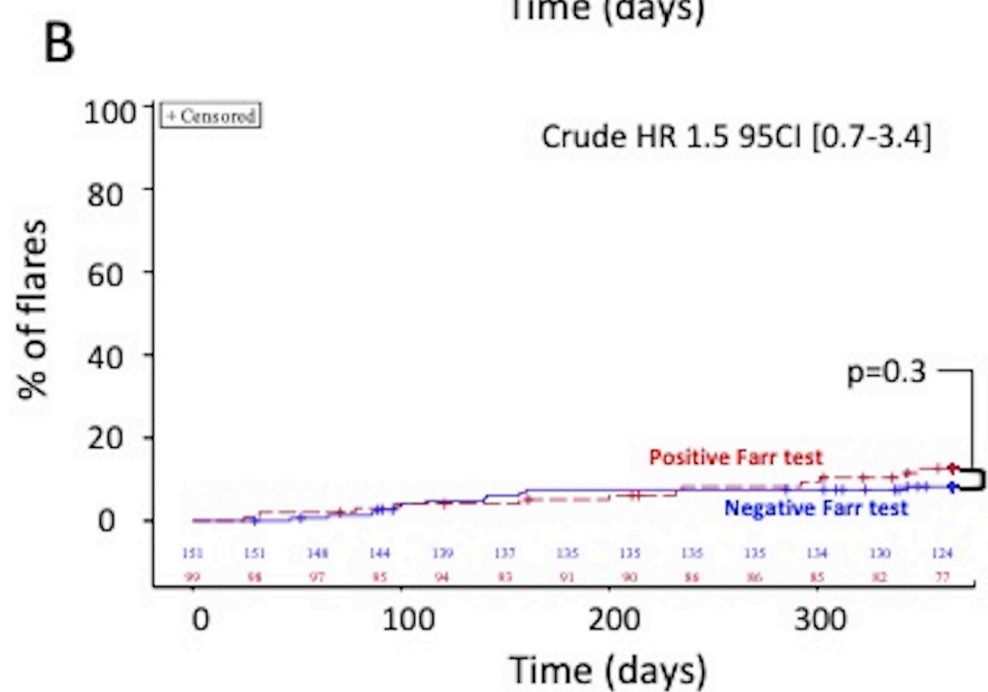
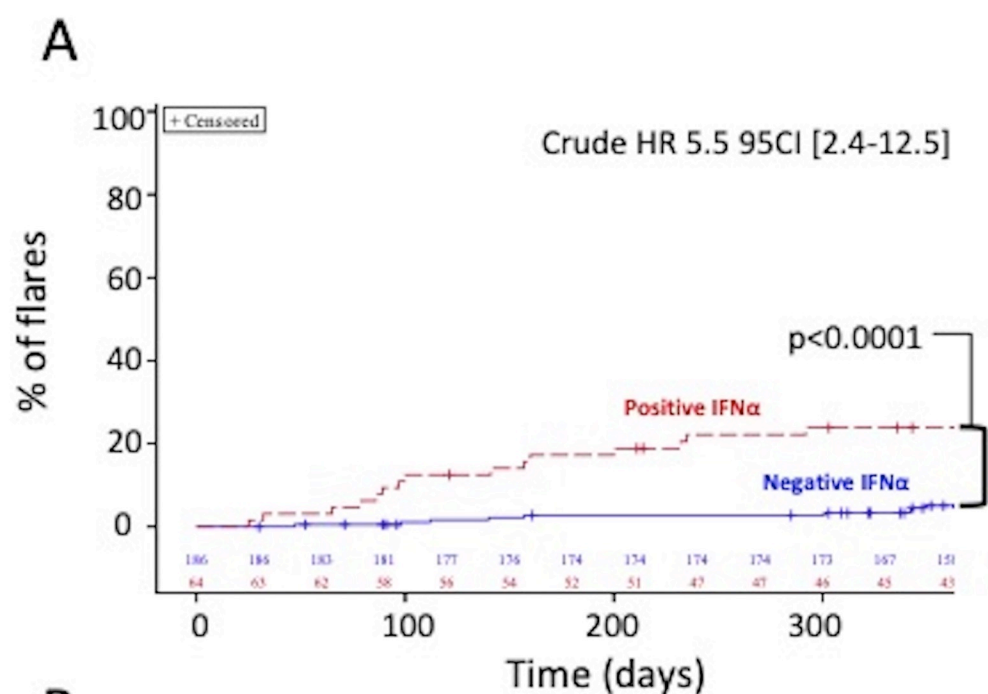


Figure 2

