

Original article

Impact of disease activity and treatments on ovarian reserve in patients with rheumatoid arthritis in the ESPOIR cohort

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Abstract

Objectives. Patients with RA have a higher prevalence of infertility than the general population. This study sought to examine the impact of RA disease activity and treatments on ovarian reserve measured by serum anti-Müllerian hormone (AMH) levels in the ESPOIR cohort. We sought to better define the indications for fertility preservation.

Methods. Patients and serum analysis data were derived from the French national cohort ESPOIR. Enrolled patients ($n=102$; 18–37-year-olds) fulfilled ACR/EULAR 2010 criteria for RA. Serum AMH levels were measured at T0, T6, T12, T24 and T36 months post-diagnosis. The impacts of RA activity (DAS28 and CRP level) and treatments (MTX only or with other medications) were evaluated at each study visit.

Results. A gradual decrease in patients' serum AMH levels was observed over time, in line with the descending curve described for healthy women. Serum AMH levels of RA patients in comparison with the values considered normal for age did not reveal any significant differences ($P > 0.05$). We did not observe any impact of RA treatments. We demonstrated an inverse correlation between AMH variation and disease activity (DAS28: $r = -0.27$, $P = 0.003$; CRP: $r = -0.16$, $P = 0.06$).

Conclusion. This is the first study to determine serum AMH levels of a large cohort of RA patients over 36 months. Rapid disease activity control appears to be required to limit changes in the ovarian reserve. Fertility preservation is not likely to be necessary if inflammation is promptly controlled.

ClinicalTrials.gov Identifier: NCT03666091.

Key words: rheumatoid arthritis, inflammation, fertility, woman, methotrexate, ovarian reserve, AMH/DAS28, disease activity

Rheumatology key messages

- There is an adverse impact of RA activity on serum AMH levels.
- Rapid disease activity limitation appears to be required to limit changes in the ovarian reserve.
- Fertility preservation for RA women seems not to be necessary if inflammation is promptly controlled.

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Introduction

RA may affect the fertility of women suffering from this pathology [1–6]. Recent studies have shown unexplained

Submitted 14 April 2020; accepted 10 June 2020

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infertility [7], with longer mean time to pregnancy [8–11], higher prevalence of dysovulation [12, 13], nulliparous women [14], and reduced ovarian reserve [12, 15] compared with the general population. However, it is not clear if these women's subfertility is due to RA activity and/or their treatments. Ovarian reserve can be evaluated by measuring the serum anti-Müllerian hormone (AMH) level. AMH is a hormone produced by granulosa cells surrounding oocytes in growing follicles that have undergone recruitment in the ovary [16–18]. Serum AMH level is considered to be an accurate biomarker of ovarian reserve, capable of reflecting the size of the ovarian follicular pool at reproductive age [19–21]. The value of this biomarker is not influenced by the period of the menstrual cycle and will reflect an early reduction of the ovarian follicle reserve [22]. To mitigate the decrease in follicle reserve linked to gonadotoxic treatment or disease, fertility preservation is currently possible for these women through the cryopreservation of oocytes and/or ovarian tissue [23]. When needed, the thawed oocytes may be used for assisted reproductive technology by intra-cytoplasmic sperm injection, and ovarian tissue may be grafted if infertility is confirmed. Recent studies reporting the impact of RA and its treatment on fertility were performed with a short time of follow-up [24] or only small patient cohort sizes [24, 25]. Hence, there is no consensus in the literature on indications of fertility preservation for these patients regarding RA activity and the treatments they receive.

The aim of our study was to examine the impact of RA activity and its related treatments on ovarian reserve measured by serum AMH level in a large cohort of patients for 36 months. The information obtained in this analysis may help to better define indications of fertility preservation for these patients.

Methods

Patients

Patient data and blood sample analyses were obtained from the French prospective multi-centred observational cohort ESPOIR [26] (Etude et Suivi des Polyarthrites Indifférenciées Récentes) sponsored by the French Society of Rheumatology (FSR), in December 2002. The ESPOIR cohort is a French database for various investigations (diagnostic, prognostic, economics, etc.). The study included 813 patients aged 18–70 years with early arthritis and attending 14 rheumatology centres in France between December 2002 and April 2005. To be included, patients had to have RA present for <6 months before inclusion or undifferentiated inflammatory arthritis (UA) with potential for developing RA, also for <6 months. UA is early arthritis that does not meet the criteria for another diagnosis, such as erythema nodosum or sacroiliitis, or that has joint characteristics and extra-articular characteristics incompatible with RA.

All the patients recruited in the ESPOIR cohort were naive to any treatment at inclusion. Only anti-

inflammatory medication taken for <15 days before inclusion was allowed. Patients were followed up by the same investigator every 6 months for 2 years and every year for at least 10 years. The protocol of the ESPOIR cohort was approved by the ethics committee of Montpellier, France (no. 020307). All the patients gave their signed informed consent before inclusion. One biological resources centre (Sarah Tubiana, Paris Bichat) was in charge of centralizing and managing biological data collection.

In our study, we enrolled women ($n = 102$) 18–37 years old who fulfilled the 2010 ACR/EULAR criteria for RA. Fifteen women without treatment who were considered to have UA with the potential to become RA were used as a chronic inflammatory disorder control group. This group allowed us to analyse the impact of inflammation itself on ovarian reserve without confounding by any potential effect of treatment. We excluded men, and women under the age of 18 and over 37 years old. We chose the age limit of 37 years because the serum AMH level decreases significantly and physiologically at this age due to follicular atresia [18]. We also excluded patients with ovarian surgery and premature ovarian insufficiency (POI) defined, according to the ESHRE guideline [27] on the management of POI 2015, by loss of ovarian activity before the age of 40 years. POI is characterized by menstrual disturbance (amenorrhoea or oligomenorrhoea) with raised gonadotropins and low oestradiol.

Measurements

The primary outcome was the global evolution of the serum AMH level (ng/ml), adjusted for age, over 36 months of follow-up in patients with RA. The measurement of AMH level was performed using the electrochemiluminescence method with a Cobas® e411 analyser (Roche Diagnostics, Meylan, France). The threshold of <1 ng/ml was considered to be a proof of severe decrease in ovarian reserve. The serum AMH assay was performed on patients from the ESPOIR cohort at baseline (T0), 6, 12, 24 and 36 months after inclusion (T6, T12, T24 and T36, respectively). All the results were expressed as the median value of AMH level (ng/ml). Blood samples were available for 117 patients at T0, 111 at T6, 106 at T12, 99 at T24, and 82 at T36 months of follow-up.

We analysed the impact of RA activity and inflammation on the serum AMH level. Disease activity was scored using the Disease Activity Score in 28 joints (DAS28 CRP) and biological parameters of inflammation: CRP in mg/l and ESR in mm/h. Anti-CCP antibodies were measured from thawed frozen sera using Immunoscan CCplus®, Euro-Diagnostica BV, Arnhem, the Netherlands.

Finally, we evaluated the potential impact of treatments used on AMH level. The treatments used were MTX only or with other treatments. The focus on MTX was justified by the high frequency of its use as the first line of treatment in RA according to the EULAR and

FSR recommendations [28, 29]. To study the potential effects of the treatments, we analysed four groups of patients according to their treatment: MTX only ($n = 27$); MTX and other treatments ($n = 56$) (hydroxychloroquine, leflunomide, sulfasalazine, etanercept, adalimumab, rituximab and infliximab); treatment without MTX ($n = 19$) (leflunomide, sulfasalazine and hydroxychloroquine); and a control group without treatment ($n = 15$). We also studied the impact of corticosteroid on AMH level, with 69 exposed patients and 37 patients without corticosteroid treatment.

Statistical analysis

Statistical analyses were performed using Stata software (version 13, StataCorp, College Station, TX, USA). All tests were two-sided, with a type I error set at 0.05. Continuous data were expressed as median (interquartile range) according to the statistical distribution. The assumption of normality was assessed using the Shapiro-Wilk test.

Concerning baseline data, the comparisons were conducted using (i) χ^2 and Fisher's exact tests, as appropriate, for categorical data, and (ii) analysis of variance (ANOVA), or the Kruskal-Wallis test when the assumptions of ANOVA were not met, for continuous parameters. When appropriate (omnibus P -value < 0.05), a post-hoc test for multiple two-by-two comparisons was performed: Tukey-Kramer after ANOVA and Dunn post Kruskal-Wallis. To study the relationships between continuous parameters, Pearson or Spearman correlation coefficients were estimated. Random-effects models

(more precisely, linear mixed models) were performed to compare the evolution of the AMH level between treatment groups, (i) studying the following fixed effects: group, time-points, and their interaction, and (ii) taking into account the between- and within-patient variability (considering the subjects as random-effect). Sidak's type I error correction was applied to perform multiple comparisons. The normality of residuals was analysed using the Shapiro-Wilk test and a logarithmic transformation of the AMH level was then conducted to assess the normality. Multivariable analyses were then carried out with adjustment for age and DAS28 at inclusion, as the treatment groups differed in these parameters at inclusion. A sensitivity analysis was performed, considering the AMH level as a normal/non-normal dichotomous parameter. The age-related threshold reported in the literature was applied. To compare the evolution of the AMH level between treatment groups, a generalized linear mixed model with a logit link function was used.

Results

Baseline demographics and disease activity

Study population details are shown in Table 1. In all, 117 women from the ESPOIR cohort were included. They were 29.9 (24.8–33.5) years old. Among these enrolled women, 102 (90%) fulfilled the 2010 ACR/EULAR RA classification criteria and were diagnosed as RA patients. They began their treatment on the day of inclusion (T0). Fifteen (10%) women, diagnosed as

TABLE 1 Baseline characteristics of the patient population

Characteristics of population at inclusion (T0)	Total patients ($n = 117$)	MTX group ($n = 27$)	MTX + other group ($n = 56$)	No MTX group ($n = 19$)	Control group ($n = 15$)	P
Age (years)	29.9 (24.8–33.5)	31.5 (27.7–34.4)	30.1 (24.7–33.6)	28.2 (23.9–32.1)	28.2 (25.0–33.8)	0.40
BMI (kg/m²)	22.2 (20.5–25.0)	23.6 (20.6–25.5)	22.2 (20.4–24.5)	21.9 (21.0–25.0)	25.5 (19.4–28.0)	0.31
Smokers, n (%)	67 (51%)	12 (44%)	29 (52%)	8 (42%)	9 (60%)	0.53
RF positivity, n (%)	43 (33%)	11 (40%)	26 (46%)	4 (21%)	0 (0%)	0.002
ACPA positivity, n (%)	45 (35%)	15 (55%)	25 (44%)	2 (10%)	0 (0%)	0.001
CRP (mg/l)	10 (5–24)	11 (6–32)	10 (4–26)	7 (4–18)	6 (4–24)	0.83
ESR (mm/h)	20 (10–35)	34 (16–56)	18 (10–35)	19 (8–25)	18 (10–35)	0.18
DAS28	5.0 (4.3–5.8)	5.5 (4.5–6.2)	5.0 (4.3–5.8)	4.6 (3.8–5.5)	3.9 (2.8–4.8)	0.001
HAQ	0.94 (0.50–1.38)	1.25 (0.88–1.63)	0.88 (0.50–1.50)	0.88 (0.5–1.13)	0.88 (0.13–1.25)	0.02

Continuous variables are presented with median [interquartile range (IQR)] and categorical parameters variable with number of patients and percentages. DAS28, disease activity score in 28 joints.

having UA without treatment, were included in the chronic inflammatory disorder control group. These women were 28.2 (25.0–33.8) years old.

Median BMI was 22.2 (20.5–25.0) kg/m². Sixty-seven patients (51%) were smokers. As shown in Table 1, there were no differences in these characteristics between the evaluated patients and the control group.

Rheumatoid factor (RF) was positive in 43 (33%) patients and ACPAs were positive in 45 (35%) patients. There was a significant difference in both these parameters between the subgroup of patients treated with MTX and the other groups without MTX ($P < 0.05$). There was no difference in serum AMH level between ACPA-positive patients and ACPA-negative patients, with respective values of 1.52 (0.63–3.26) ng/ml and 1.72 (0.97–2.81) ng/ml ($P = 0.64$). The median serum CRP level was 10 (5–24) mg/l and the median ESR level was 20 (10–35) mm/h. There was no significant difference in CRP and ESR levels at T0 between any groups of patients.

The patients' median value of DAS28 was 5.0 (4.3–5.8), corresponding to moderate disease activity. The DAS28 score of the subgroup of patients treated with MTX was higher at 5.5 (4.5–6.2) corresponding to high RA activity compared with the other groups ($P = 0.001$), and a higher level of inflammation and disease activity for these patients. The HAQ score was significantly higher in the MTX group [1.25(0.88–1.63)] compared with the other groups ($P < 0.02$).

Impact of disease activity and inflammation on serum AMH level

The analysis of serum AMH level over the time and as function of age describes a gradual decrease (Fig. 1). There was no significant difference between AMH level at T12, T24 and T36 compared with T0 ($P > 0.05$). We observed a lower AMH level at T0 compared with T6 (1.70 ng/ml vs 2.20 ng/ml; $P = 0.017$) in the group of 102 patients with treatment. Control group serum AMH

levels measured at the different times during the 36-month follow-up period did not differ significantly.

At T0, the serum AMH level was negatively correlated with the DAS28 score ($r = -0.12$; $P = 0.15$) and with the biological parameters of inflammation (CRP, $r = -0.26$; ESR, $r = -0.19$; $P = 0.002$) (Fig. 2).

There was a statistically significant decrease in RA activity (measured by DAS28 score; $r = -0.28$; $P = 0.003$) and a trend for inflammation (measured by serum CRP level; $r = -0.17$; $P = 0.06$), which was correlated with the increase in AMH level (1.70 ng/ml vs 2.20 ng/ml) between T0 and T6 for all the patients evaluated, whatever treatment they received. The control of inflammation was correlated with an improvement in AMH level. These variations were not observed in the control group ($r = 0$; $P = 0.99$).

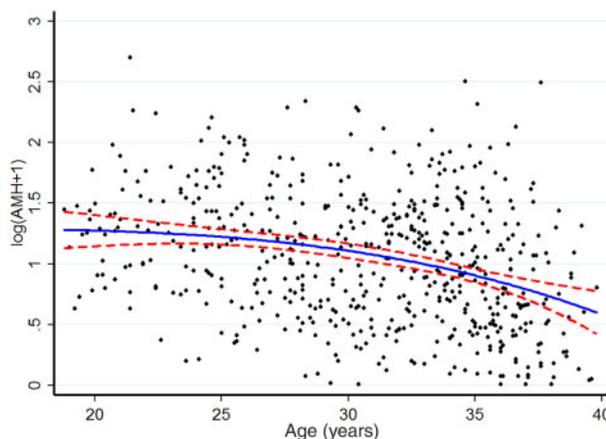
There was no statistical difference between groups regarding the evolution of AMH level between T6, T12 ($P = 0.09$), T24 ($P = 0.35$) and T36 ($P = 0.61$).

Impact of treatments on serum AMH level during 36 months of follow-up

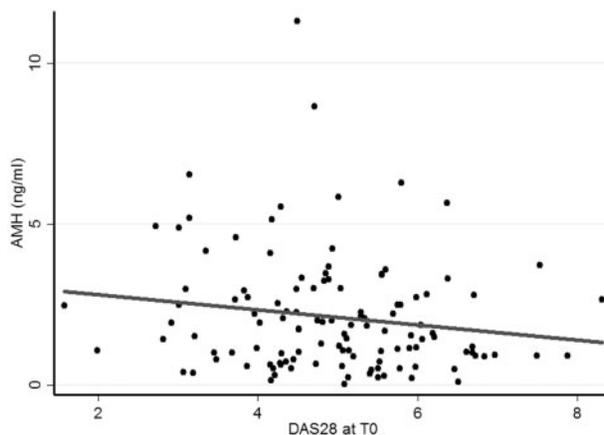
At T0, we found a lower serum AMH level in the MTX-only patient group ($n = 27$) compared with the without-MTX patient group ($n = 19$; $P = 0.002$) and the control group ($n = 15$; $P > 0.05$). The DAS28 score was higher in the with MTX patient group [DAS28 at 5.5 (4.5–6.2)] compared with the no MTX patient group [DAS28 at 4.6 (3.8–5.5)], $P = 0.001$.

Although the mean AMH level tended to decrease during the follow-up, we did not observe a significant difference between the serum AMH levels at T12, T24 and T36 compared with T0 ($P > 0.05$), whatever the treatments (MTX or not) in the patients evaluated, even after adjustment for the DAS28 score. Indeed, the serum AMH level evolved from 1.13 ng/ml to 1.14 ng/ml at T36 in the MTX-only group ($n = 27$), from 1.52–1.34 ng/ml at T36 in the MTX with or without other treatments ($n = 83$) group, and from 1.72 ng/ml to

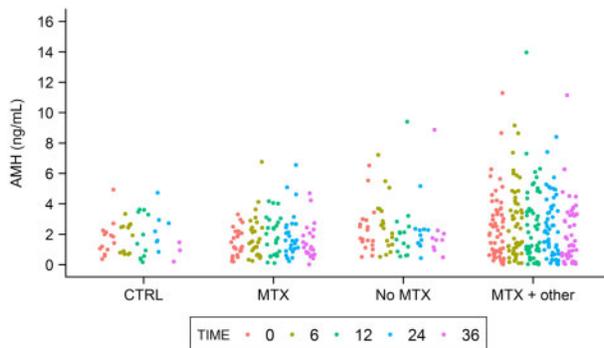
Fig. 1 Evolution of serum AMH level during 36 months of follow-up according to age of patients



The relationship between age of the patients and serum AMH level was analysed using fractional-polynomial prediction (blue line). Red line corresponds to the 95% CI. AMH, anti-Müllerian hormone.

Fig. 2 Correlation between serum AMH level and parameter of inflammation

The serum AMH level at the diagnosis of the disease is negatively correlated with the DAS28 score ($r = -0.12$; $P = 0.15$). AMH, anti-Müllerian hormone; DAS, disease activity score in 28 joints.

Fig. 3 Evolution of serum AMH level for each treatment

Serum AMH level according to the treatments received by the patients [MTX only ($n = 27$); No MTX ($n = 19$); MTX + other treatment group ($n = 56$); CTRL: control group ($n = 15$) at different time of evolution of the RA disease (diagnosis: T0; 6 months: T6; 12 months: T12; 24 months: T24; and 36 months: T36)]. AMH, anti-Müllerian hormone.

1.62 ng/ml at T36 in the without MTX group ($n = 19$), vs from 1.62 ng/ml to 0.94 ng/ml in the control group ($n = 15$) (Fig. 3).

There was no difference in the evolution of the serum AMH level (Fig. 1) over time between RA patients under treatment ($n = 102$) and the control group ($n = 15$; $P = 0.13$) and between groups of patients who fulfilled ($n = 102$) or did not fulfil ($n = 15$) the ACR/EULAR 2010 criteria ($P > 0.05$).

In regard to corticosteroid treatments, we did not find a difference in serum AMH levels between users ($n = 69$) and non-users ($n = 37$) at any time of the comparison, even with doses higher than 7.5 mg per day ($n = 6$).

Discussion

By following this cohort of women of reproductive age with early RA for a period of 36 months, we were able to

clearly show that RA activity was correlated with a decrease in serum AMH level. However, AMH level as a function of age in these patients is similar to the physiological concentration of AMH over time reported by Kelsey *et al.* [30] in women without RA. In addition, we did not observe any impact of treatments used to treat RA on AMH level of these women.

This study enabled us to show clearly the impact of RA activity on serum AMH level early in the disease. This decline in serum AMH level is linked to the decrease in ovarian reserve we observed when diagnosing RA before starting treatment. The strength of our study lies in the size of the ESPOIR cohort and the long follow-up period of 36 months. No previous study has carried out as lengthy a longitudinal follow-up of an equally large number of patients. We were only able to evaluate ovarian reserve on the basis of serum AMH level because ovarian echography had not been performed for the patients included in the study.

The impact of RA on female fertility remains controversial. Henes *et al.* [25] showed that patients with RA had a lower AMH level, a lower pregnancy rate (Del Junco *et al.* [1]), and a longer waiting time to pregnancy [8, 9] compared with the general population. Other authors focussed more particularly on the effect of MTX, since it is a first-line treatment for RA. Two studies pointed to a reduction of ovarian reserve after exposure to MTX [31, 32]. Nonetheless, these observations were not confirmed by Brouwer *et al.* [24] in a follow-up of RA patients over a 6-month period of exposure to MTX. A similar result was observed in a recent study of Eudy *et al.* in 2019 [7], with no impact of MTX on AMH level. In our study, we did not find an impact of MTX during the 36-month follow-up period in the patients exposed only to MTX ($n = 27$) or in those exposed to MTX in addition to other treatments ($n = 56$). The diversity of groups of treatments in our study is explained by the fact that the cohort of patients is multi-centred and by the difficulty of clearly determining the diagnosis of RA in the framework of early arthritis. Such diagnosis involves a variety of recommendations and diverse clinical rheumatologic practices, as shown by Benhamou *et al.* [33]. This likely explains our study's possible lack of power in its evaluation of the effects of treatments, due to the lower number of patients in each group after adjustment. However, it is noteworthy that a large proportion of the treatments used are not known to affect fertility—this reduces the risk of interference in the interpretation of our results with, in particular, leflunomide, hydroxychloroquine, and sulfasalazine. Only a severe impact would justify preserving fertility. Our study therefore confirms results published previously that do not recommend preserving fertility for patients undergoing treatment with immunosuppressive doses of MTX [34]. However, it is important to note that since the formation of this cohort new and effective RA treatments have been developed. It will be interesting to analyse the impact of the new generation of treatments on AMH in a further study.

Regarding corticosteroid therapy, no significant difference in serum AMH level was observed between the exposed and unexposed patients. This observation does not confirm the data in the literature. Brouwer *et al.* [9] highlighted a link between corticosteroid therapy and a reduction in fertility in 66% of the 245 women in the Pregnancy-induced Amelioration of RA (PARA) cohort composed of patients following corticosteroid therapy at a dose of 7.5 mg/day. When patients followed corticosteroid therapy at a dose higher than this daily dose, an increase in the waiting time to pregnancy was observed in comparison with a control group. In our study, the number of patients exposed to this dose was very low, which perhaps explains the observed absence of difference.

The analysis of AMH levels according to ACPA status did not show any significant difference contrary to the previous study in 2019 by Brouwer *et al.* [35], which demonstrated lower AMH levels in ACPA-positive

patients compared with healthy women as a control group. In our study the control group consisted of patients with UA. Moreover, we applied the same assay to measure serum AMH level in patients and control groups. This may explain this lack of difference in our study.

No correlation was found in previous studies between serum inflammatory parameters, disease activity and AMH level. Brouwer *et al.* [35] did not show any association between disease activity and serum AMH level in a recent study. However, their study was based on a population whose disease activity was under control and who had participated in a pregnancy project. Our study clearly shows a correlation between RA activity and AMH level. At inclusion, inflammation linked to RA was not controlled by treatment, with RA activity evaluated as moderate or high as a function of group, according to the DAS28 score. The organization of treatment from the time of diagnosis of RA makes it possible to see its probable beneficial impact on disease evolution at 6 months, with a significant improvement in serum AMH level whatever the treatment group. An increased prevalence of infertility of unknown aetiology was shown by Brouwer *et al.* [12] in these patients. Our data appear to show that it might be an effect of inflammation on the ovarian reserve.

RA activity, which reflects the magnitude of articular inflammation, generally appears to have a significant impact on serum AMH level. Previously, it was only possible to draw a link between increased waiting time to pregnancy and the inflammatory activity of RA [3]. The hypothesis used to explain this association was potential sexuality disorders linked to a fall in libido related to inflammation-induced pain. Pikwer *et al.* showed an association between the occurrence of the menopause before the age of 45 and the development of RA, with an odds ratio (OR) of 2.42, 95% CI: 1.32, 4.45 [36]. The authors hypothesized a link between the inflammation present before the diagnosis and the reduction of ovarian reserve, leading to premature menopause. The same observation was made by Del Junco *et al.* who demonstrated that female fertility was reduced before the diagnosis of RA was made [1]. In our study also, it was at T0 that we found the lowest AMH level. In comparison with other models of chronic systemic diseases such as Crohn's disease, it has been shown that serum AMH values are also clearly reduced in patients with intense disease activity compared with a control group of patients in remission [37]. Inflammation therefore appears to play a crucial role in fertility in RA.

Indeed, proinflammatory cytokines such as IL and TNF α can harm the vascular system, possibly explaining the link between the inflammatory activity of a disease and damage to the ovarian reserve [33, 36]. Paradisi *et al.* [38] studied the potential link between cytokines and the ovarian reserve in patients with Hodgkin and non-Hodgkin lymphoma before the administration of chemotherapy. A negative correlation was found

between AMH level and IL-6 in these patients [38]. In an animal model, IL-1 was implicated in the regulation of the ovarian reserve through an increase of serum AMH level, with a better response of the ovaries to gonadotropins in IL-1 beta-deficient mice [39]. It would therefore be interesting to extend our study by measuring cytokine levels in the serum and/or follicular fluid to analyse their potential influence and better understand how inflammation affects the ovarian reserve of patients with RA. One might imagine that reducing inflammation would remove this shock to the production of AMH by the granulosa cells. Accordingly, it would be interesting to know whether the granulosa cells can be destroyed by long exposure to inflammation (driven by cytokines), since that would clearly link inflammation to a reduced ovarian reserve.

In conclusion, our study highlights the harmful impact of RA activity on the serum AMH level, an indicator of the ovarian reserve. We did not demonstrate the impact of treatments. Our study therefore tends to show that patients with RA do not need to preserve their fertility. Nevertheless, it appears important to thoroughly control the inflammatory activity of RA from the time it is considered as a diagnosis, in order to limit the premature impact of inflammation on the ovarian reserve. In the future, it would be interesting to measure IL levels in follicular fluid to better understand the physiopathology of the damage incurred.

Acknowledgements

We wish to thank Nathalie Rincheval (CHU Montpellier and EA 2415) who did expert monitoring and data management, and all the investigators who recruited and followed the patients (F. Berenbaum, Paris-Saint Antoine; M. C. Boissier, Paris-Bobigny; A. Cantagrel, Toulouse; B. Combe, Montpellier; M. Dougados, Paris-Cochin; P. Fardellone and P. Boumier, Amiens; B. Fautrel, Paris-La Pitié; R. M. Flipo, Lille; Ph. Goupille, Tours; F. Liote, Paris-Lariboisière; O. Vittecoq, Rouen; X. Mariette, Paris Bicêtre; P. Dieude, Paris Bichat; A. Sarau, Brest; T. Schaeverbeke, Bordeaux; and J. Sibilia, Strasbourg). We finally wish to thank A. Vega and C. Lambert [Biostatistics Unit (DRCI)], CHU Clermont-Ferrand for their contributions for the preparation of figures and submission of the manuscript. At last we would like to thank Roche Diagnostics for the generous gift of the AMH assays to support this study.

Funding: An unrestricted grant from Merck Serono was allocated for this study. Two additional grants from INSERM were obtained to support part of the biological database of the ESPOIR cohort. The French Society of Rheumatology, Pfizer, Abbvie and Lilly also supported the ESPOIR cohort study.

Disclosure statement: The authors have declared no conflicts of interest.

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