

RHEUMATOLOGY

# **Clinical science**

# DAS28-γGT for the prediction of major cardiovascular events in rheumatoid arthritis: results from the ESPOIR cohort

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# Abstract

**Objective:** To validate the predictive value of the DAS28 γ-glutamyl transferase (DAS28-γGT) for the occurrence of major cardiovascular (CV) events (MACE) in the 'Etude et Suivi des Polyarthrites Indifférenciées Récentes' ESPOIR cohort.

**Methods:** Analysis of 13-year outcome from the ESPOIR cohort. RA patients with missing data for baseline  $\gamma$ GT activity and those not followed-up to 1 year were excluded. Baseline DAS28- $\gamma$ GT was calculated using the following formula: 0.56\* $\sqrt{TJ-28}$ + 0.28\* $\sqrt{SJ-28}$ + 2\*In( $\gamma$ GT) + 0.014 \* GH. Our primary outcome was the merit of the DAS28- $\gamma$ GT in predicting the occurrence of MACE.

**Results:** Among the 696 patients [536 women, mean (s.b.) age of 49 (12) years], 34 MACE were recorded, with a mean time to event of 71 (44) months. Receiver operating characteristic curve analysis indicated that a DAS28- $\gamma$ GT >9.4 had the best sensitivity and specificity for the diagnosis of MACE during the observation period. DAS28- $\gamma$ GT >9.4 was predictive of the occurrence of MACE, with a hazard ratio (HR) of 3.11 (95% CI 1.41, 5.43). Multivariate Cox analyses confirmed higher DAS28- $\gamma$ GT (HR 2.44, 95% CI 1.05, 5.64) together with age (HR 1.04, 95% CI 1.01, 1.07) and diabetes mellitus (HR 4.12, 95% CI 1.55, 10.95) as independent predictors of MACE. There was a dose effect of the DAS28- $\gamma$ GT for MACE-risk prediction, which was in line with the application of the Framingham risk score.

**Conclusion:** The DAS28- $\gamma$ GT was identified in this large prospective cohort as an independent predictor of MACE in patients with RA. The DAS28- $\gamma$ GT is a simple and useful tool to evaluate CV risk in routine and warn the clinician about the CV risk burden in patients with RA. **Keywords:** RA, cardiovascular risk, composite index

#### Rheumatology key messages

- The DAS28-γGT correlated with the Framingham risk score and steadily increased according to cardiovascular risk.
- Higher baseline DAS28-yGT was identified as an independent predictor of MACE in the ESPOIR cohort.
- The DAS28-γGT is a simple tool to evaluate the cardiovascular risk in routine in RA.

# Introduction

It is now well accepted that patients with RA have an increased cardiovascular (CV) risk. This risk is related to both the burden of traditional CV risk factors and additional disease-related factors, particularly chronic inflammation. Compared with the general population, the CV risk of RA patients is 1.5- to 2-fold higher than age- and sex-matched individuals [1–3], and even higher when traditional CV risk factors are associated. Several lines of evidence have shown that an aggressive management of joint and systemic inflammation, in particular with targeted biologic therapies like TNF- $\alpha$  inhibitors, could significantly reduce the number of CV events [4, 5]. On the other hand, the recent publication of ORAL SURVEILLANCE study has raised concerns around a possible increased risk of major CV events (MACE) in tofacitinib-treated patients in comparison with TNF- $\alpha$ inhibitors [5].

Thus, optimizing the CV risk management is crucial, as stated by the recent EULAR recommendations [6]. However, the guidelines recognize the suboptimal performance of risk scores used in the general population when applied to patients with chronic inflammatory rheumatic disorders, leading to

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underestimation of CV risk in these populations. Thus, given the important gaps in knowledge still existing regarding the approach to CV risk stratification in clinical practice and the need to use disease-specific risk prediction models [7], there is clearly a need for new relevant and feasible methods to predict CV risk in RA.

 $\gamma$ -Glutamyl transferase ( $\gamma$ GT) is a plasma membrane enzyme that is expressed in kidney, liver and pancreatic cells. Their activity is increased in many systemic syndromes, including inflammatory or autoimmune diseases. Ample evidence suggests that elevated  $\gamma$ GT is associated with increased risk of CV diseases [8]. We have previously showed that replacing ESR by  $\gamma$ GT in DAS28 calculation (DAS28- $\gamma$ GT) allowed a combined evaluation of CV risk in patients with RA, in addition to joint disease activity [8]. However, this study was limited by its observational design, the relatively small number of patients included in some analyses and the use of surrogates for CV risk. This justifies the setting of a prospective study to determine the validity of DAS28- $\gamma$ GT levels and its predictive value for the occurrence of CV events in RA populations.

Our objective was to validate with a prospective, longitudinal, multicentre observational cohort the predictive value of the DAS28- $\gamma$ GT for the occurrence of MACE in the 'Etude et Suivi des Polyarthrites Indifférenciées Récentes' (ESPOIR) cohort.

### **Patients and methods**

#### Study design and setting

We took over the ESPOIR cohort, a prospective multicentre observational cohort that included patients with early diagnosis of arthritis from 14 French rheumatology centres [9]. Patients had to have inflammatory arthritis in at least two swollen joints lasting from 6 weeks to 6 months, with the potential to develop into RA, and be naïve to DMARDs and CS therapy. This cohort included 813 patients between 2002 and 2005. They were followed up to 13 years. The objective, design and characteristics of the cohort were previously described [10, 11]. All the patients gave their signed informed consent before the inclusion. The protocol of the ESPOIR cohort and of this study was endorsed by the ethics committee of Montpellier, France (No. 020307). All the data of the ESPOIR cohort were available after being approved by the scientific committee (www.lacohorteespoir.fr).

#### Inclusion and exclusion criteria

Among the 813 patients included in the ESPOIR cohort, we selected patients who fulfilled the 2010 ACR/EULAR criteria for RA over the 13 years of follow-up. We excluded patients with missing data for  $\gamma$ GT activity measurement at inclusion and those not followed up to 1 year.

#### DAS28-yGT calculation and outcome

 $\gamma$ GT was a part of the routine initial workup measured at inclusion in the ESPOIR cohort. The DAS28- $\gamma$ GT is a screening tool providing to the rheumatologist rapid information related to both joint disease activity and CV risk. It was obtained by replacing ESR by baseline  $\gamma$ GT levels with the following formula 0.56 \*  $\sqrt{T}J$ -28 + 0.28 \*  $\sqrt{S}J$ -28 + 2 \* ln( $\gamma$ GT) + 0.14 \* GH, as previously reported [8].

We first examined the association between the DAS28- $\gamma$ GT and CV risk factors at baseline in the entire cohort. We considered smoking, BMI >30 kg/m<sup>2</sup>, diabetes mellitus, high blood pressure and hypercholesterolemia, as previously defined in the cohort [8]. Ten-year risk prediction of CV disease was estimated by the algorithm developed by the Framingham Heart Study [12]. Regular alcohol consumption was defined by up to one drink a day for women and up to two drinks a day for men. Excessive alcohol consumption was defined by a consumption >14 alcohol units a week. We used the Fibrosis-4 (Fib-4) index to evaluate underlying hepatic fibrosis, as previously described [13].

We next assessed the merit of the DAS28- $\gamma$ GT to predict the occurrence of a MACE. To that end, we considered at each study visit the occurrence of MACE during the exposition period, defined as death from CV causes, non-fatal myocardial infarction or non-fatal stroke [5]. For this specific analysis, we excluded the patients with a history of coronary heart disease or ischaemic stroke at baseline.

#### Statistical analysis

All data are presented as mean values (s.D.) or number and percentage (%) for continuous and categorical variables, accordingly. Statistical analysis was performed using GraphPad Prism (v10). We used the unpaired *t*-test for two-group comparisons (continuous variables) and the  $\chi^2$ -test for differences in frequency (binary variables). Three group comparisons were analysed by one-way analysis of variance (ANOVA) with Dunnett's multiple comparisons test. Correlations between the DAS28- $\gamma$ GT and numeric variables were assessed using Spearman's rank correlation coefficient (rS). The diagnostic value of the DAS28- $\gamma$ GT and the DAS28 was assessed by receiver operating characteristic (ROC) curve analysis.

MACE-free survival according to the DAS28- $\gamma$ GT was estimated by Kaplan–Meier survival curves. To identify predictive factors of MACE, we used Cox proportional-hazard regression. This analysis included MACE as the dependent variables and all relevant identified covariates were then entered in one single step. A *P*-value <0.05 was considered statistically significant.

#### Results

#### Study population

A total of 696 patients with RA (536 women, 77%) were included, with a mean (S.D.) age of 49 (12) years and a mean (S.D.) disease duration of 97 (190) days. Positive RF and anti-CCP antibodies were detected in 327 (47%) and 282 (41%) patients, respectively, and bone erosions were present in 258/ 676 (38%) patients. These patients had high disease activity, with a mean (s.D.) DAS28 of 5.24 (1.27). At baseline, 9 patients (1%) received CS and 44 (6%) received conventional synthetic DMARDs (csDMARDs), including 32 (4%) with MTX alone or in combination. During the observation period, 92 patients (13%) were treated by CS, 618 (89%) by csDMARDs and 405 (58%) by MTX. The mean (s.D.) Framingham risk score of this population was 9.4 (8.1) (median 6.0%, 95% CI 5.40, 7.0), 238 (34%) patients had more than two CV risk factors and 9 had a history of coronary heart disease or ischaemic stroke. Detailed patients' characteristics are presented in Table 1.

#### DAS28-yGT for MACE prediction

#### Table 1. Patients' characteristics

|   | Total population $(N = 696)$ | DAS28- $\gamma$ GT $\leq$ 9.4 ( $n = 334$ ) | DAS28- $\gamma$ GT >9.4<br>( <i>n</i> = 362) | P-value |
|---|------------------------------|---|--|---------|
| Baseline demographics                                   |                              |   |  |         |
| Age (years), mean (s.D.)                                | 49 (12)                      | 46 (12.5)                                   | 52 (11)                                      | < 0.001 |
| Women, $n$ (%)  | 536 (77)                     | 285 (85)                                    | 251 (69)                                     | < 0.001 |
| Baseline disease characteristics                        |                              |   |  |         |
| Disease duration (days), mean (s.D.)                    | 97 (190)                     | 93 (192)                                    | 103 (191)                                    | 0.49    |
| Positive RF, n (%)                                      | 327 (47)                     | 165 (49)                                    | 162 (45)                                     | 0.29    |
| Positive anti-CCP2 antibodies, $n$ (%)                  | 282 (41)                     | 136 (41)                                    | 146 (40)                                     | 0.79    |
| Erosions on hand/foot X-rays, $n/N$ (%)                 | 258/676 (38)                 | 109/325 (34)                                | 149/351 (42)                                 | 0.032   |
| Baseline disease activity                               |                              |   |  |         |
| Tender joints, mean (s.D.)                              | 9 (7)                        | 6 (5)                                       | 12 (8)                                       | < 0.001 |
| Swollen joints, mean (s.D.)                             | 8 (5)                        | 6 (4)                                       | 9 (6)  | < 0.001 |
| DAS28, mean (s.D.) <sup>a</sup>                         | 5.24 (1.27)                  | 4.59 (1.05)                                 | 5.83 (1.16)                                  | < 0.001 |
| DAS28 > 3.2, n/N(%)                                     | 645/685 (94)                 | 295/329 (90)                                | 350/356 (98)                                 | < 0.001 |
| DAS28 > 5.1, n/N(%)                                     | 361/685 (53)                 | 102/329 (31)                                | 259/356 (73)                                 | < 0.001 |
| ESR (mm/h), mean $(S.D.)^{b}$                           | 30 (25)                      | 24 (20)                                     | 36 (28)                                      | < 0.001 |
| $CRP (mg/l), mean (S.D.)^{c}$                           | 23 (35)                      | 16 (25)                                     | 29 (40)                                      | < 0.001 |
| γGT activity, mean (s.D.)                               | 38 (44)                      | 18 (9)                                      | 57 (54)                                      | < 0.001 |
| Baseline function                                       |                              |   |  |         |
| HAQ, mean (s.D.)  | 1.0 (0.68)                   | 0.77 (0.58)                                 | 1.22 (0.71)                                  | < 0.001 |
| Baseline cardiovascular risk factors                    |                              |   |  |         |
| Modifiable CV risk factors                              |                              |   |  |         |
| Smokers, $n$ (%)  | 335 (48)                     | 148 (44)                                    | 187 (52)                                     | 0.035   |
| High blood pressure, $n$ (%)                            | 127 (18)                     | 42 (13)                                     | 85 (23)                                      | < 0.001 |
| Diabetes mellitus, $n$ (%)                              | 28 (4)                       | 6 (2)                                       | 22 (6)                                       | 0.008   |
| Hypercholesterolemia, $n$ (%)                           | 104 (15)                     | 36 (11)                                     | 68 (19)                                      | 0.003   |
| $BMI (kg/m^2)$ , mean (s.d.)                            | 25 (5)                       | 24 (4)                                      | 26 (5)                                       | < 0.001 |
| BMI > 30 kg/m <sup>2</sup> , $n$ (%)                    | 296 (43)                     | 112 (35)                                    | 184 (51)                                     | < 0.001 |
| Patients with $\geq 2$ CV risk factors                  | 238 (34)                     | 74 (22)                                     | 164 (45)                                     | < 0.001 |
| Mean Framingham risk score, %, mean (s.D.) <sup>d</sup> | 9.4 (8.1)                    | 6.9 (6.7)                                   | 12.1 (8.6)                                   | < 0.001 |
| Baseline regular alcohol intake                         | 125 (18)                     | 51 (15)                                     | 74 (20)                                      | 0.083   |
| Baseline excessive alcohol intake <sup>e</sup>          | 25 (4)                       | 3 (1)                                       | 22 (6)                                       | < 0.001 |
| Baseline Fib-4 index, mean (s.D.) <sup>f</sup>          | 0.81 (0.48)                  | 0.79 (0.43)                                 | 0.83 (0.62)                                  | 0.25    |
| Treatment received during the observation period        |                              |   |  |         |
| Current CS use, $n$ (%)                                 | 92 (13)                      | 34 (10)                                     | 58 (16)                                      | 0.019   |
| Current use of NSAIDs                                   | 631 (91)                     | 298 (89)                                    | 333 (92)                                     | 0.18    |
| Current conventional DMARD use, n (%)                   | 618 (89)                     | 289 (87)                                    | 329 (91)                                     | 0.091   |
| Current MTX use, <i>n</i> (%)                           | 405 (58)                     | 184 (55)                                    | 221 (61)                                     | 0.11    |

<sup>a</sup> Calculated on 685 patients with available data.

<sup>b</sup> Calculated on 688 patients with available data.

<sup>c</sup> Calculated on 686 patients with available data.

<sup>a</sup> Calculated on 334 patients with available data.

<sup>e</sup> Defined by a consumption >14 alcohol units a week.

<sup>t</sup> Calculated on 694 patients with available data.

γGT: γ-glutamyl transferase; CV: cardiovascular; Fib-4 index: Fibrosis-4 index.

#### Evaluation of DAS28-yGT and CV risk at baseline

For this analysis, the entire cohort of 696 patients was considered. At baseline, the DAS28- $\gamma$ GT correlated with age (rS = 0.27, P < 0.001). It was significantly higher in men [10.39] (1.79) vs 9.50 (1.83), P < 0.001 and in patients presenting the following conditions: excessive alcohol consumption [11.34 (1.87) vs 9.64 (1.83), P < 0.001], active smoking [9.86] (1.86) vs 9.56 (1.84), P = 0.034], high blood pressure [10.40] (1.95) vs 9.55 (1.80), P < 0.001], hypercholesterolemia [10.18 (1.92) vs 9.62 (1.83), P < 0.001], diabetes mellitus [11.03 (2.99) vs 9.65 (1.83), P < 0.001] and obesity (BMI  $>30 \text{ kg/m}^2$  [10.13 (1.85) vs 9.39 (1.85), P < 0.001]. No association was observed between DAS28-yGT and treatment with baseline NSAIDs or CS or with the Fib-4 index. The DAS28-yGT correlated with the Framingham risk score (rS = 0.35, P < 0.001)—it steadily increased according to CV risk (Fig. 1A) and was significantly higher in patients with at least two CV risk factors (Fig. 1B), as well as in the nine patients with a history of coronary heart disease or ischaemic stroke [10.88 (1.47) *vs* 9.64 (1.85), P = 0.046]. The DAS28yGT had a diagnostic value for the presence of at least two CV risk factors characterized by an area under the curve (AUC) of 0.72 (95% CI 0.68, 0.75, P < 0.001) (Supplementary Fig. S1A, available at *Rheumatology* online). Conversely, the DAS28 and the DAS28-CRP did not correlate with the Framingham risk score and their diagnostic value for the presence of at least two CV risk factors was characterized by an AUC of 0.53 (95% CI 0.49, 0.56, P = 0.29) and 0.51 (95% CI 0.48, 0.56), respectively (Supplementary Fig. S1B and C, available at *Rheumatology* online).

# Predictive value of the DAS28-yGT for the occurrence of MACE

For this analysis, the 9 patients a history of coronary heart disease or ischemic stroke were excluded, and 687 patients were thus considered. During the observation period of 130 (34) months, a total of 34 MACE were recorded, with a mean time to event of 71 (44) months. ROC curve analysis indicated



**Figure 1.** DAS28- $\gamma$ GT levels and cardiovascular risk factors. (**A**) DAS28- $\gamma$ GT levels according to cardiovascular risk evaluated by the Framingham risk score (<10%, 10–20% and >20%) (\*\*P<0.01 and \*\*\*P<0.001 by analysis of variance followed by Dunnett's multiple comparisons test); (**B**) DAS28- $\gamma$ GT levels according to the number of cardiovascular risk factors ( $\leq 2$  or >2 risk factors) (\*\*\*\*P<0.0001 by Student's *t*-test).  $\gamma$ GT:  $\gamma$ -glutamyl transferase

that a DAS28- $\gamma$ GT >9.4 had the best sensitivity (74%) and specificity (62%) for the diagnosis of MACE during the observation period. At baseline, patients with a DAS28- $\gamma$ GT >9.4 were more likely to be men, older, with a more active and severe disease, and they were at higher CV risk (Table 1).

DAS28- $\gamma$ GT >9.4 was predictive of the occurrence of MACE during the observation period, with a hazard ratio (HR) of 3.11 (95% CI 1.41, 5.43) (Fig. 2). Conversely, a DAS28 >5.1 or a DAS28-CRP >5.1 were not predictive of MACE (HR 1.12, 95% CI 0.56, 2.20 and 1.15, 95% CI 0.46, 2.93, respectively). The diagnostic value of DAS28-yGT for the occurrence of MACE was characterized by AUC of 0.70 (95% CI 0.61, 0.73) compared with 0.55 (95% CI 0.51, 0.59) for the DAS28 and 0.55 (95% CI 0.51, 0.58) for the DAS28-CRP (Supplementary Fig. S2A-C, available at Rheumatology online). Pairwise comparisons of ROC curves have confirmed an increased predictive value of the DAS28- $\gamma$ GT over the DAS28 (P=0.026) and the DAS28-CRP (P = 0.026) and no significant difference between the DAS28- $\gamma$ GT and the Framingham risk score (P = 0.38).  $\gamma$ GT activity was also predictive of MACE on its own but to a lesser extent than the DAS28-yGT (HR 3.02, 95% CI 1.58, 6.33 and AUC of 0.63, 95% CI 0.61, 0.68).

Multivariate Cox analyses adjusting for the variables age, sex, CRP levels, HAQ, CS use, excessive alcohol intake, smoking, high blood pressure, diabetes, dyslipidaemia and obesity confirmed higher DAS28- $\gamma$ GT (HR 2.44, 95% CI 1.05, 5.64), together with age (HR 1.04, 95% CI 1.01, 1.07) and diabetes mellitus (HR 4.12, 95% CI 1.55, 10.95) as independent predictors of MACE (Table 2). A second model including only statistically significant variables by univariate analysis identified the same predictors of MACE (Table 2).

To determine whether there was a dose effect of the DAS28- $\gamma$ GT for MACE risk prediction, we assessed the predictive value of three tertiles of DAS28- $\gamma$ GT: low (<8.26, 25th percentile), intermediate (8.26–10.72) and high (>10.72, 75th percentile) DAS28- $\gamma$ GT. Fig. 3 illustrates that the risk of MACE was significantly higher in patients with a DAS28- $\gamma$ GT >10.72, compared with those with intermediate or low DAS28- $\gamma$ GT. In particular, patients with a DAS28- $\gamma$ GT <8.26 had a decreased risk of MACE (HR 0.33, 95% CI 0.20, 0.97, *P* = 0.038).

We next compared by multivariate Cox analyses the three tertiles of DAS28-yGT with the validated cut-off of the Framingham risk score (mild <10%, moderate 10–20%, high >20%) (DAS28-yGT <8.26 vs Framingham risk score <10%; DAS28-yGT 8.26-10.72 vs Framingham risk score 10-20%; and DAS28-yGT >10.72 vs Framingham risk score >20%) for the occurrence of MACE (dependent variable) during the observation period. There was a trend for a decreased risk of MACE for patients with a DAS28- $\gamma$ GT < 8.26 (HR 0.18, 95% CI 0.02, 1.37), which was not observed for patients with a Framingham risk score <10% (Table 3). Patients with intermediate DAS28-yGT had an increased risk of MACE (HR 2.79, 95% CI 1.08, 7.22), compared with a trend for patients with an intermediate Framingham risk score (10–20%) (Table 3). A trend for a higher risk of MACE was observed in patients with a DAS28-yGT >10.72 (HR 1.98, 95% CI 0.75, 5.23), compared with a significant risk in patients with an intermediate Framingham risk score >20%(HR 4.04, 95% CI 1.63, 9.96) (Table 3).



**Figure 2.** Predictive value of the DAS28- $\gamma$ GT for the occurrence of MACE during the observation period. Time to MACE according to circulating the DAS28- $\gamma$ GT ( $\leq$  or >9.4). This analysis was performed on the 687 patients with no history of MACE during a mean (s.p.) observation period of 130 (34) months, with a total of 34 MACE recorded.  $\gamma$ GT:  $\gamma$ -glutamyl transferase; MACE: major adverse cardiovascular event

 Table 2. Univariate and multivariate Cox analyses to identify independent predictors of MACE (primary endpoint)

| Variable at baseline      | MACE                                   |  |  |
|---------------------------|--|--|--|
|                           | Univariate<br>analysis<br>(HR, 95% CI) | Multivariate<br>analysis<br>(HR, 95% CI) |  |
| First model <sup>a</sup>  |  |  |  |
| DAS28-γGT >9.4            | 3.11 (1.41, 5.43)                      | 2.44 (1.05, 5.64)                        |  |
| Age                       | 1.05 (1.02, 1.10)                      | 1.04 (1.01, 1.07)                        |  |
| Men                       | 1.64 (1.01, 3.37)                      | 1.02 (0.46, 2.29)                        |  |
| Smokers                   | 2.36 (1.69, 3.30)                      | 1.33 (0.65, 2.70)                        |  |
| Excessive alcohol intake  | 1.67 (0.39, 6.87)                      | 1.18 (0.53, 2.65)                        |  |
| High blood pressure       | 3.09 (1.54, 6.17)                      | 1.67 (0.74, 3.76)                        |  |
| Dyslipidemia              | 1.25 (0.51, 3.03)                      | 0.65 (0.25, 1.64)                        |  |
| Diabetes                  | 6.86 (2.83, 16.61)                     | 4.12 (1.55, 10.95)                       |  |
| $BMI > 30 \text{ kg/m}^2$ | 1.11 (0.56, 2.18)                      | 0.63 (0.30, 1.32)                        |  |
| Current CS use            | 5.06 (0.69, 37.01)                     | 5.12 (0.69, 37.69)                       |  |
| CRP levels                | 1.21 (1.01, 3.22)                      | 1.01 (0.99, 1.01)                        |  |
| HAQ                       | 1.53 (0.96, 2.45)                      | 1.21 (0.72, 1.48)                        |  |
| Second model <sup>b</sup> |  |  |  |
| DAS28-γGT >9.4            | 3.11 (1.41, 5.43)                      | 2.80 (1.09, 4.83)                        |  |
| Age                       | 1.05 (1.02, 1.10)                      | 1.03 (0.99, 1.07)                        |  |
| Men                       | 1.64 (1.01, 3.37)                      | 1.07 (0.49, 2.31)                        |  |
| Smokers                   | 2.36 (1.69, 3.30)                      | 1.30 (0.64, 2.64)                        |  |
| High blood pressure       | 3.09 (1.54, 6.17)                      | 1.51 (0.68, 3.33)                        |  |
| Diabetes                  | 6.86 (2.83, 16.61)                     | 3.87 (1.50, 10.88)                       |  |
| CRP levels                | 1.21 (1.01, 3.22)                      | 1.00 (0.99, 1.01)                        |  |

Bold font is used to highlight significant values. A total of 687 patients with no history of MACE were considered in this analysis.

<sup>a</sup> First model including MACE as the dependent variables and all potentially relevant covariates.

<sup>b</sup> Second model including MACE as the dependent variables and only statistically significant covariates identified through the univariate analysis. HR: hazard ratio; MACE: major adverse cardiovascular event; γGT: γ-glutamyl transferase.

# Discussion

Regardless of the selected method to risk stratification, screening on a regular basis for CV risk factors and adapting treatment based on estimated CV risk are critical to all approaches to primary prevention of MACE in RA [6]. This has also been recently highlighted by the Janus kinase inhibitor–related potential increased risk of MACE in comparison with TNF- $\alpha$ , which now requires, as stated by the Food and Drud Administration and the European Medicines Agency, a detailed and personalized CV evaluation before considering this therapeutic class [14].

A central principle of the EULAR recommendations for CV risk management in rheumatic diseases states that 'rheumatologists are responsible for CV risk assessment and management in collaboration with primary care providers, internists or cardiologists' [6]. This assertion patently sets the rheumatologist at the centre of CV preventive care, which may present several limitations when applied in clinical practice, and some adaptation and flexibility of this approach may be required. Considerable gaps in the management of CV risk are still a major issue despite an increased understanding of this problem among rheumatologists [15, 16]. Studies that attempted to identify obstacles to CV risk management among rheumatologists specified as some of the main reasons lack of time, lack of knowledge of current guidelines for CV risk factors and lack of care coordination [17, 18]. Indeed, patients should undertake measurement of blood pressure, glucose and lipid profiles on a regular basis, and assessment of lifestyle factors for input into a risk-prediction calculator to estimate CV risk. These procedures may be considered as too complex and time consuming in clinical practice by the



**Figure 3.** Predictive value of DAS28-yGT tertiles for the occurrence of MACE during the observation period. Time to MACE according to circulating the DAS28-yGT tertile. This analysis was performed on the 687 patients with no history of MACE during a mean (s.b.) observation period of 130 (34) months, with a total of 34 MACE recorded. yGT: y-glutamyl transferase; MACE: major adverse cardiovascular event

Table 3. Comparison of the DAS28- $\gamma GT$  with the Framingham risk score for the occurrence of MACE

| Variable at baseline                          | Multivariate analysis<br>(HR, 95% CI) | P-value |  |
|---|---------------------------------------|---------|--|
| Model 1                                       |                                       |         |  |
| DAS28- $\gamma$ GT <8.26 ( $n = 150$ )        | 0.18 (0.02, 1.37)                     | 0.18    |  |
| Framingham risk score $<10\%$ ( $n = 185$ )   | 0.64 (0.26, 1.58)                     | 0.59    |  |
| Model 2                                       |                                       |         |  |
| DAS28- $\gamma$ GT 8.26–10.72 ( $n = 357$ )   | 2.79 (1.08, 7.22)                     | 0.034   |  |
| Framingham risk score $10-20\%$ ( $n = 107$ ) | 0.29 (0.05,1.26)                      | 0.29    |  |
| Model 3                                       |                                       |         |  |
| DAS28- $\gamma$ GT >10.72 ( $n = 180$ )       | 1.98 (0.75, 5.23)                     | 0.16    |  |
| Framingham risk score $>20\%$ ( $n = 42$ )    | 4.04 (1.63, 9.96)                     | 0.002   |  |

Only patients with no history of MACE were considered in this analysis. All three comparisons were performed during a mean (s.D.) observation period of 130 (34) months, with a total of 34 MACE recorded. Each model was adjusted on age and sex.  $\gamma$ GT:  $\gamma$ -glutamyl transferase; HR: hazard ratio.

rheumatologist. Thus, it is crucial to provide to rheumatologists a simple tool, feasible in clinical practice, to evaluate CV risk. We constructed a new composite index called DAS28- $\gamma$ GT, replacing ESR with  $\gamma$ GT levels; this index has been shown equivalent to the DAS28 for the assessment of disease activity, and also provided added value to identify the presence of CV risk factors [8]. The present study brings a further step of validation of this index in this 13-year analysis of the ESPOIR cohort. As previously observed [8], the DAS28- $\gamma$ GT was significantly higher in patients with CV risk factors and correlated with the Framingham CV risk score. In addition,

higher baseline DAS28- $\gamma$ GT were able to independently predict the risk of MACE after stratification on potential confounders including alcohol consumption, systemic inflammation and the presence of CV risk factors. Interestingly, there was a dose effect of the DAS28- $\gamma$ GT for MACE risk prediction, similar to what is observed with established risk scores, with lower values being associated with a decreased CV risk and higher values being more at risk.

The EULAR recommendations also highlight some topics regarding CV risk prevention that plainly fall within the scope of practice of all rheumatologists. Achieving optimal control of rheumatic disease activity is an essential treatment objective from a CV standpoint, as studies have observed an association between various measures of disease activity and CV risk [19]. This is also an advantage of the DAS28- $\gamma$ GT, which has been shown to be a reliable marker of RA disease activity, equivalent to the DAS28 for the assessment of disease activity evaluated by clinical examination or power Doppler US [8].

The DAS28- $\gamma$ GT is a simple and useful tool, with the advantage of providing information on disease activity and CV risk with a single index, which may warn the clinician about the CV risk burden in patients with RA. It may be used in clinical practice to assess joint disease activity without losing validity compared with the DAS28 and may help rheumatologists to decide whether RA patients require more in-depth CV evaluation, and need to be referred to a cardiologist [7].

The limitations of the present study include those that cannot be ignored in observational cohort studies, with potential confounding factors that could not be considered. Moreover, most data regarding comorbidities were declarative, with potential recall bias. This cohort included early RA patients with high disease activity, and further data are needed to confirm these findings in patients with established disease and/or lower disease activity. The ascertainment of CV death was performed at a clinic visit, and people who had died in the interim might have been undercounted. However, the prolonged follow-up and the multiple annual data review may have decreased the risk to miss death during the interval of two visits. Cause of death is mentioned in the case report form of ESPOIR and CV events were clearly identified.

Since  $\gamma$ GT activity has also been identified as a predictive factor of MACE in the general population [8], it is not known whether  $\gamma$ GT may predict MACE differently in RA compared with the non-RA population. However, adding the  $\gamma$ GT to the DAS28 increased the prediction of MACE compared with  $\gamma$ GT alone, and this index considers RA disease activity.

Our study has a number of strengths. The ESPOIR cohort allows exploration of the time-dependent risk of MACE in early RA in a real-world setting, providing long follow-up time with a low rate of missing data or drop-out.

In summary, the DAS28- $\gamma$ GT was identified in this large prospective cohort as an independent predictor of MACE in patients with RA. In addition to the assessment of disease activity, the DAS28- $\gamma$ GT is a simple and useful tool to evaluate CV risk in routine and warn the clinician about the CV risk burden in patients with RA.

# **Supplementary material**

Supplementary material is available at Rheumatology online.

# **Data availability**

All the data of the ESPOIR cohort were available after being approved by the scientific committee (www.lacohorteespoir. fr). The authors declare that all other data supporting the findings of this study are available within the paper (and its supplementary information files).

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