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Association Between Baseline Anti-cyclic Citrullinated Peptide Antibodies and 6-Month Clinical Response Following Abatacept or TNF Inhibitor Treatment: A Real-World Analysis of Biologic-Experienced Patients with RA

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Association Between Baseline Anti-cyclic Citrullinated Peptide Antibodies and 6-Month Clinical Response Following Abatacept or TNF Inhibitor Treatment: A Real-World Analysis of Biologic-Experienced Patients with RA

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ABSTRACT

Introduction: Anti-cyclic citrullinated peptide (anti-CCP) antibodies are associated with poor prognosis in patients with rheumatoid arthritis (RA). Previous data from randomized controlled trials and clinical practice have shown anti-CCP-positive (+) patients had a better response to treatment with abatacept or tumor necrosis factor inhibitor (TNFi) treatment than those who were anti-CCP negative. This study assessed the association between baseline anti-CCP2 [a surrogate for anti-citrullinated protein antibody (ACPA)] concentration and 6-month treatment responses to abatacept or TNFi in patients with RA.

Supplementary Information The online version contains supplementary material available at <https://doi.org/10.1007/s40744-021-00310-2>.

Sabrina Rebello: CorEvitas employee at the time of the work that was conducted on this project.

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Methods: This real-world analysis included biologic-experienced patients from CERTAIN (Comparative Effectiveness Registry to study Therapies for Arthritis and Inflammatory CoNditions) who initiated abatacept or TNFi, had prior biologic disease-modifying drug exposure and baseline anti-CCP2 concentration/serostatus and serum samples (baseline and 6 months). Baseline demographics and disease characteristics were compared. Change from baseline at 6 months in Clinical Disease Activity Index (CDAI) score and patient-reported outcomes [PROs: pain, fatigue, patient global assessment (PtGA), modified Health Assessment Questionnaire (mHAQ) score], by baseline anti-CCP2 quartile and binary cut-off (> 10–250 and > 250 U/ml), were evaluated separately in the abatacept and TNFi groups using a linear regression model adjusted for age, sex, CDAI/PROs, comorbidity index, and methotrexate use.

Results: Included were 138 abatacept and 137 TNFi initiators who were anti-CCP2+. At

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baseline, there were significant differences between anti-CCP2 quartiles and mean CDAI, swollen joint count 28, C-reactive protein (CRP), Disease Activity Score 28 (CRP), rheumatoid factor (RF), mHAQ and physician global assessment among abatacept initiators, and in mean RF, mHAQ, and PtGA among TNFi initiators. Among abatacept (but not TNFi) initiators, CDAI numerically improved ($p = 0.208$) and PROs significantly improved ($p < 0.05$) with increasing baseline anti-CCP2.

Conclusions: In patients treated with abatacept, not TNFi, higher anti-CCP2 concentrations at baseline were associated with numerically greater improvements in CDAI and significant improvements in PROs after 6 months.

Clinical Trial Number: NCT01625650.

PLAIN LANGUAGE SUMMARY

Rheumatoid arthritis (RA) is an autoimmune disease – a disease that causes the immune system to attack an individual's own body. A key feature of RA is the presence of proteins called autoantibodies in the blood. While antibodies help protect against external threats such as viruses, autoantibodies mistakenly target an individual's own tissues and organs. One type of autoantibody often found in patients with RA is called anti-cyclic citrullinated peptide (anti-CCP). Studies have shown that patients with RA with anti-CCP antibodies may experience worse physical symptoms, function, disease activity, and outcomes than patients with RA without anti-CCP antibodies. Clinical trials suggest that some drugs may be more effective than others at managing symptoms of RA in patients who have anti-CCP in their blood. It is important to study this further to give doctors a sense of how patients respond to drug therapy in the 'real world', without clinical trial constraints. This study examined real-world patient data to see whether the presence of anti-CCP in patients' blood impacted how their RA symptoms responded to treatment with two different drugs: abatacept or a tumor necrosis factor inhibitor (TNFi). This study found that patients with higher levels of anti-CCP at the start of the study, compared with patients with lower levels of anti-CCP, experienced less disease activity and greater improvement in

physical function after 6 months of treatment with abatacept. The study found no relationship between anti-CCP and treatment response after 6 months of treatment with a TNFi.

Keywords: Abatacept; Patient-reported outcome measures; Rheumatoid arthritis; Tumor necrosis factor inhibitors

Key Summary Points

Why carry out this study?

Anti-cyclic citrullinated peptide (anti-CCP) antibodies are highly specific serological biomarkers associated with poor clinical prognosis in patients with rheumatoid arthritis (RA), including higher disease activity, more rapid radiographic disease progression, and greater impairment of physical function.

An important consideration for physicians treating patients with RA is that treatment responses can vary depending on the clinical characteristics of the patients; therefore, an individualized treatment approach may be needed.

Previous data from clinical trials and clinical practice suggest that bDMARDs with different mechanisms of action may result in differential treatment responses, depending on a patient's anti-CCP serostatus.

What was learned from the study?

In this post hoc analysis of the prospective CERTAIN cohort study, biologic-experienced patients with RA and higher (versus lower) anti-CCP2 concentrations at baseline had numerically, but not statistically, greater improvements in CDAI score and significantly greater improvements in patient-reported outcomes after 6 months of treatment with abatacept.

No association between baseline anti-CCP2 concentration and treatment response was observed in patients treated with a tumor necrosis factor inhibitor.

DIGITAL FEATURES

This article is published with digital features, including a summary slide and plain language summary to facilitate understanding of the article. To view digital features for this article go to <https://doi.org/10.6084/m9.figshare.14438786>.

INTRODUCTION

Rheumatoid arthritis (RA) is characterized by the production of autoantibodies, including rheumatoid factor (RF) and anti-citrullinated protein antibodies (ACPAs) [1, 2]. Anti-cyclic citrullinated peptide (anti-CCP, a surrogate for ACPA), is a highly specific serological biomarker that physicians can use in the clinic to guide therapy; anti-CCP positivity is associated with poor prognosis in patients with RA [3–5]. Patients with RA who are ACPA/anti-CCP positive (+), compared with those who are ACPA/anti-CCP negative (–), have been shown to have worse clinical outcomes, including higher disease activity, more rapid radiographic disease progression, greater impairment of physical function, and increased mortality [3–7].

Current European League Against Rheumatism (EULAR) and American College of Rheumatology (ACR) treatment guidelines recommend a treat-to-target approach in order to achieve and maintain a state of sustained remission or low disease activity [4, 8]. However, there are few data available to guide sequencing of biologic disease-modifying antirheumatic drugs (bDMARDs) in the event of treatment failure with first-line therapy with conventional synthetic DMARDs, such as methotrexate (MTX). An important consideration for physicians treating patients with RA is that treatment responses can vary depending on the clinical characteristics of the patients. Thus, an individualized treatment approach may be needed [9, 10]. Specific serological biomarkers, such as ACPAs, that are indicative of a poor prognosis may be useful as part of such an approach. ACPAs are measured using anti-CCP antibody assays; the second and third generations of anti-CCP assays (CCP2 and CCP3) have

been shown to be good measures of CCP [11, 12].

Clinical trial data suggest that bDMARDs with different mechanisms of action may result in differential treatment responses depending on a patient's ACPA serostatus [13–15]. A post hoc analysis of the AMPLE (Abatacept versus adaliMumab comParison in bioLogic-naivE RA subjects with background MTX) trial demonstrated that patients treated with abatacept or a tumor necrosis factor inhibitor (TNFi; adalimumab) who were anti-CCP2+, versus those who were anti-CCP2–, had a better response to treatment at 12 months [13]. Additionally, patients with RA who had higher baseline anti-CCP2 concentrations showed a better response to treatment with abatacept than those with lower concentrations, an association that was not observed with adalimumab treatment [13]. Real-world data from the large, US-based, Corrona RA Registry have demonstrated similar findings. An analysis from the Corrona RA Registry, which included patients with RA initiating treatment with abatacept ($n = 566$) or a TNFi ($n = 1715$), showed that anti-CCP status was associated with a differential treatment response to abatacept but not to TNFi [14].

CERTAIN (Comparative Effectiveness Registry to study Therapies for Arthritis and Inflammatory CoNditions) was a prospective cohort study of adult patients with RA (ClinicalTrials.gov: NCT01625650) [16]. Patients included in CERTAIN were recruited from the Corrona RA Registry network and had at least moderate disease activity [Clinical Disease Activity Index (CDAI) score ≥ 10], had either started therapy with or switched to a TNFi or non-TNFi bDMARD and were followed for 12 months or until they switched/discontinued the bDMARD [16, 17]. Prior analyses of patients who were anti-CCP+ (bDMARD-naive or -experienced) from the Corrona RA Registry network have shown that patients had similar outcomes at 6 months when treated with abatacept versus a TNFi: there was no significant difference in response [17]. However, a subgroup analysis found that there was a significant difference in reduction of disease activity at 6 months in favor of abatacept, compared with TNFi treatment, among patients with anti-

CCP+RA who were biologic-experienced [17]. Therefore, the purpose of the present study was to use data from CERTAIN to evaluate the effects of different baseline anti-CCP2 concentrations on the change in clinical and patient-reported outcome (PRO) response from baseline to 6-month follow-up among patients separately initiating abatacept or a TNFi.

METHODS

Study Sample

Overall CERTAIN Population

The design of the CERTAIN study has been published previously [16]. Patients in CERTAIN, recruited from the Corrona RA Registry network [16, 17], were enrolled from 43 private practices and academic sites with over 100 participating rheumatologists. CERTAIN included information on 2350 patients, with 10,077 patient visits and a mean patient follow-up time of 10.3 months (median 11.7 months). Patients were included in CERTAIN if they were at least 18 years of age, had at least moderate disease activity (CDAI score > 10) at enrollment, either started therapy (biologic-naïve) with or switched from a prior bDMARD to a TNFi or non-TNFi bDMARD (biologic-experienced) and were followed for 12 months or until they switched/discontinued bDMARD therapy [16]. Regular data collection at 3-month intervals was mandated in CERTAIN, with uniform collection of anti-CCP status for all patients initiating treatment with a bDMARD.

Analysis Population

Patients from CERTAIN were included in this analysis if they initiated abatacept or any TNFi (adalimumab, etanercept, certolizumab, golimumab, infliximab), had visits and serum samples available at baseline and 6-month follow-up, had prior bDMARD exposure, and were anti-CCP3+(> 20 U/ml) at baseline.

Propensity Score Matching A propensity score model was used to identify patients with similar disease activity and similar number of prior biologics at baseline in order to determine

which serum samples to send for anti-CCP2 testing. Using a logistic regression model with abatacept initiation as the outcome, a propensity score was estimated using baseline CDAI as the only covariate stratified by prior number of biologics: 1 prior biologic and > 1 prior biologic. Using the propensity score, patients treated with TNFi were matched to patients treated with abatacept, using 1:1 matching based on the propensity score (log odds) with a caliper of 0.025 within each prior biologic strata. Serum samples for 186 abatacept patients and 186 TNFi patients were sent for anti-CCP2 testing. Anti-CCP2 levels ≤ 10 U/ml were defined as negative and those > 10 U/ml as positive.

Anti-CCP Analysis

In CERTAIN, there was a uniform collection of specimens for biomarker analysis, thus allowing for simultaneous testing of anti-CCP status at the same time and by a single laboratory for the entire cohort. For this analysis, patients with anti-CCP3 positivity were selected from CERTAIN. Upon enrolment in the CERTAIN database from 2010 to 2014, anti-CCP3 testing was centrally performed on a rolling basis by ICON Laboratories (Farmingdale, NY, USA) who were blinded to the clinical outcomes. Anti-CCP2 is more commonly used in clinical practice than anti-CCP3 and is considered the gold standard. As such, all anti-CCP2 testing was performed on the samples from patients in CERTAIN in a single batch by Exagen Diagnostics (Vista, CA, USA). The patient samples were evaluated in a single batch to minimize variability in processing or reagents that could have occurred over time. Patients who were anti-CCP2+ at baseline (> 10 U/ml) were further divided into equal quartiles: quartile 1, 11–94 U/ml; quartile 2, 95–296 U/ml; quartile 3, 297–876 U/ml; and quartile 4, > 876 U/ml.

Outcome Measures

Patient data from baseline and 6-month follow-up visit were included. All baseline visits and sampling occurred within 36 h prior to the first dose of the new biologic agent. Variables

collected at baseline included patient demographics, lifestyle characteristics, disease characteristics, comorbidities, prior medication use, RA disease activity, and PROs. Treatment response by baseline anti-CCP2 quartile was assessed in patients initiating treatment with abatacept or a TNFi.

Primary Outcome

The primary outcome was change from baseline at 6 months in CDAI score.

Secondary Outcomes

Secondary outcomes were change from baseline at 6 months in PROs: patient-reported pain and fatigue as measured by visual analog scale (VAS; 0–100), patient global assessment (PtGA; VAS 0–100), and modified Health Assessment Questionnaire (mHAQ; modification of the standard HAQ where the number of activities of daily living was reduced from 20 to 8) score.

Statistical Analysis

Baseline patient demographics and clinical characteristics were evaluated for abatacept and TNFi groups separately using descriptive statistics across anti-CCP2+ quartiles. Continuous characteristics were evaluated using one-way analysis of variance or Kruskal–Wallis test for characteristics that were not normally distributed. Categorical characteristics were evaluated using the Fisher's exact test. Outcomes were evaluated separately in the abatacept and TNFi groups and were compared across the anti-CCP2+ quartiles with a test of trend using anti-CCP2+ quartiles as the ordinal predictor in both unadjusted and adjusted linear regression models. Missing data were not imputed. Patient numbers < 5 (or < 5 derived through subtraction from the total) or a percentage of 0% or 100% (indicative of certainty) were suppressed due to potential loss of confidentiality. Covariates selected a priori for multivariate adjustment of outcomes included age, sex, CDAI, or PRO score at initiation, modified Charlson Comorbidity Index and current MTX use. Additionally, a sensitivity analysis was conducted using the typical cut-off for anti-CCP

(> 250 U/ml) used by commercial laboratories: binary anti-CCP2 cut-offs of > 10–250 and > 250 U/ml were used. Similarly, outcomes were evaluated separately in the abatacept and TNFi groups and compared between the binary anti-CCP2 cut-offs using anti-CCP2 binary cut-off as the predictor in both unadjusted and adjusted linear regression models. A *P* value < 0.05 was considered statistically significant. All statistical analyses were performed using Stata version 15 (StataCorp LLC, College Station, TX, USA).

Patient and Public Involvement

Study participants were required to provide informed consent for involvement in the study, which included demonstrating full understanding of the study design and outcomes. The research questions and outcome measures were investigated using standard, established physician and patient-derived clinical metrics of disease activity. In addition to implementation of the treatment (which was physician-led as part of routine clinical care), study participants were required to provide the PROs that were key components of the study clinical efficacy evaluation. Patients were not involved in the study design, interpretation, writing or editing of this manuscript but will be informed of the outcomes of the study.

Compliance with Ethics Guidelines

The study was conducted in accordance with the Declaration of Helsinki and International Conference on Harmonization Good Clinical Practice. The protocol and patients' informed consent received institutional review board/independent ethics committee approval prior to initiation of the study. All participating investigators were required to obtain full board approval for conducting research involving human subjects. Sponsor approval and continuing review was obtained through a central IRB (New England Independent Review Board, NEIRB No. 120160610). For academic investigative sites that did not receive a waiver to use the central IRB, approval was obtained from the respective governing IRBs and documentation

of approval was submitted to the Sponsor prior to initiating any study procedures. All registry subjects were required to provide written informed consent prior to participating.

RESULTS

Study Sample

From CERTAIN, 186 patients initiating abatacept and 557 patients initiating a TNFi met the criteria for inclusion in this post hoc analysis (Fig. 1). In total, 151 biologic-experienced patients initiating treatment with abatacept or a TNFi were included in the analysis. Of these, 13 abatacept- and 14 TNFi-treated patients were anti-CCP2– and 138 and 137 patients were anti-CCP2+, respectively.

At baseline, for patients treated with abatacept who were anti-CCP2+, the majority of patient demographics and disease characteristics were similar across anti-CCP2 quartiles (Table 1). Mean patient age was 56–59 years, 63–80% were female, the mean duration of RA was 11.5–13.6 years, and the mean body mass index (BMI) was 29.2–30.1 kg/m². There were significant differences between anti-CCP2 quartiles at baseline for several patient demographics and disease characteristics: mean CDAI score, swollen joint count in 28 joints (SJC28), C-reactive protein (CRP), Disease Activity Score 28 (DAS28) (CRP), RF, mHAQ score, and physician global assessment (all $p < 0.05$).

At baseline, for patients initiating treatment with a TNFi who were anti-CCP2+, the majority of demographic and disease characteristics were similar across anti-CCP2 quartiles (Table 2). Mean patient age was 55–61 years, 71–78% were female, the mean duration of RA was 12.1–14.7 years and the mean BMI was 27.4–30.7 kg/m². There were significant differences between anti-CCP2 quartiles at baseline in mean RF, mHAQ score, and PtGA (all $p < 0.05$).

There were no statistically significant differences in medication history and current MTX use at time of initiation across the anti-CCP2 quartiles in either the abatacept or the TNFi treatment groups.

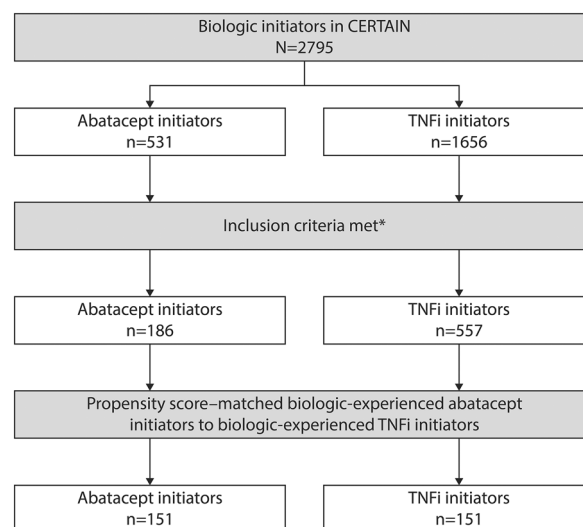


Fig. 1 Patient disposition. *Serum available at baseline and 6-month visit, has 6-month follow-up visit in CERTAIN, moderate or severe CDAI score at baseline visit, CCP3+ at baseline visit. CCP3 cyclic citrullinated peptide 3, CERTAIN Comparative Effectiveness Registry to study Therapies for Arthritis and Inflammatory CoNditions, TNFi tumor necrosis factor inhibitor/s

Clinical and Patient-Reported Outcomes by Anti-CCP2 Quartile

In the unadjusted model among patients initiating treatment with abatacept, there was a statistically significant improvement in CDAI score and in all PROs (patient-reported pain, fatigue, PtGA, and mHAQ score) across increasing anti-CCP2 quartiles (all $p < 0.05$; Supplementary Tables S1 and S2). No relationship was observed between anti-CCP concentration and clinical outcomes or PROs among patients initiating treatment with a TNFi. In the adjusted model, patients initiating treatment with abatacept (Fig. 2), but not those initiating treatment with a TNFi (Fig. 3), demonstrated increasing improvement in CDAI score at 6 months with increasing anti-CCP2 quartile; however, this trend was not statistically significant. In the adjusted model, patients initiating treatment with abatacept, but not those initiating treatment with a TNFi, demonstrated a significantly greater improvement in all PROs at 6 months with increasing anti-CCP2 quartile (Figs. 2 and 3).

Table 1 Baseline characteristics among abatacept-treated patients by anti-CCP2 quartile

Characteristic	Anti-CCP2+, U/ml ^a				<i>p</i> value ^b
	Quartile 1 (<i>n</i> = 30)	Quartile 2 (<i>n</i> = 37)	Quartile 3 (<i>n</i> = 36)	Quartile 4 (<i>n</i> = 35)	
Age, years	57.7 (12.8)	57.2 (15.1)	55.8 (12.0)	59.0 (10.3)	0.754
Female sex, <i>n</i> (%)	24 (80.0)	26 (70.3)	28 (77.8)	22 (62.9)	0.406
White race, <i>n</i> (%) ^c	NP ^d	31 (83.8)	NP ^d	NP ^d	NP ^d
Hispanic ethnicity, <i>n</i> (%) ^e	NP ^d	NP ^d	NP ^d	5 (14.3)	NP ^d
BMI, kg/m ²	30.1 (5.8)	29.2 (6.8)	30.1 (7.1)	30.1 (6.5)	0.879
Smoking history, <i>n</i> (%)					
Never	17 (56.7)	19 (51.4)	14 (38.9)	12 (34.3)	0.223
Former/Current	13 (43.3)	18 (48.6)	22 (61.1)	23 (65.7)	
Comorbidities, <i>n</i> (%)					
Hypertension	10 (33.3)	14 (37.8)	13 (36.1)	18 (51.4)	0.450
Diabetes	5 (16.7)	8 (21.6)	1 (2.8)	5 (14.3)	0.087
Malignancy	1 (3.3)	3 (8.1)	2 (5.6)	5 (14.3)	0.433
Serious infections	4 (13.3)	3 (8.1)	6 (16.7)	4 (11.4)	0.717
Cardiovascular	6 (20.0)	6 (16.2)	7 (19.4)	4 (11.4)	0.780
Duration of RA, years	11.5 (10.3)	13.6 (10.7)	12.7 (10.3)	12.0 (8.3)	0.804
CDAI score	24.2 (9.8)	28.0 (12.2)	29.2 (11.9)	32.3 (12.5)	0.041
TJC28	9.4 (6.0)	10.1 (6.6)	9.4 (6.0)	11.9 (7.4)	0.437
SJC28	5.3 (4.3)	7.9 (4.6)	9.5 (5.9)	9.1 (5.4)	0.005
CRP, mg/l	2.1 (1.8)	9.5 (14.7)	11.1 (13.1)	19.3 (19.2)	< 0.0001
DAS28 (CRP) score	4.3 (0.8)	4.8 (1.1)	4.8 (1.0)	5.3 (1.1)	0.004
RF, IU/ml	110.8 (140.8) ^f	136.4 (163.6)	362.8 (672.9)	320.7 (380.2)	< 0.0001
mHAQ	0.6 (0.5)	0.4 (0.4)	0.6 (0.6)	0.7 (0.6)	0.049
PtGA (VAS 0–100)	54.3 (28.1)	53.0 (23.4)	51.3 (29.5)	58.8 (29.6)	0.720
PGA (VAS 0–100)	40.5 (18.6)	47.0 (20.4)	51.2 (18.3)	54.5 (20.9)	0.031
Patient-reported pain (VAS 0–100)	53.9 (27.8)	51.4 (25.4)	52.8 (28.0)	65.7 (28.9)	0.067
Patient-reported fatigue (VAS 0–100)	59.0 (28.1)	55.7 (24.1)	54.0 (32.1)	58.2 (29.0)	0.844

Table 1 continued

Characteristic	Anti-CCP2+, U/ml ^a				<i>p</i> value ^b
	Quartile 1 (<i>n</i> = 30)	Quartile 2 (<i>n</i> = 37)	Quartile 3 (<i>n</i> = 36)	Quartile 4 (<i>n</i> = 35)	
Prior biologic use, <i>n</i> (%)					
1	11 (36.7)	16 (43.2)	9 (25.0)	14 (40.0)	0.593
2	13 (43.3)	13 (35.1)	14 (38.9)	11 (31.4)	
3+	6 (20.0)	8 (21.6)	13 (36.1)	10 (28.6)	
Prior TNFi use, <i>n</i> (%) ^g	30 (100.0)	35 (94.6)	36 (100.0)	35 (100.0)	0.246
Prior non-TNFi use, <i>n</i> (%) ^g	4 (13.3)	8 (21.6)	6 (16.7)	8 (22.9)	0.756
Current MTX use, <i>n</i> (%) ^h	17 (56.7)	20 (54.1)	21 (58.3)	17 (48.6)	0.860

Data are mean (SD), unless otherwise stated

^a Anti-CCP2 quartiles: quartile 1, 11–94 U/ml; quartile 2, 95–296 U/ml; quartile 3, 297–876 U/ml; and quartile 4, > 876 U/ml

^b Differences between baseline measures of disease activity across anti-CCP2+ quartiles were assessed using one-way analysis of variance for DAS28 (CRP) and PGA and the Kruskal–Wallis test for nonparametric measures (CDAI, TJC28, SJC28, CRP, RF, mHAQ, and PtGA)

^c *n* = 35 and 32 for quartiles 3 and 4, respectively

^d Patient numbers < 5 (or < 5 derived through subtraction from the total) or a percentage of 0% or 100% (indicative of certainty) were suppressed (not presented, NP) due to potential loss of confidentiality

^e *n* = 36 and 35 for quartiles 2 and 3, respectively

^f *n* = 29

^g Prior use reflects all prior biologic history; may not sum to 100%

^h At time of biologic initiation

Anti-CCP2 anti-cyclic citrullinated peptide 2, *BMI* body mass index, *CDAI* Clinical Disease Activity Index, *CRP* C-reactive protein, *DAS28 (CRP)* Disease Activity Score 28 (C-reactive protein), *mHAQ* modified Health Assessment Questionnaire, *NP* not presented, *PGA* physician global assessment, *PtGA* patient global assessment, *RA* rheumatoid arthritis, *RF* rheumatoid factor, *SD* standard deviation, *SJC28* swollen joint count in 28 joints, *TJC28* tender joint count in 28 joints, *TNFi* tumor necrosis factor inhibitor/s, *VAS* visual analog scale

Sensitivity Analyses

Sensitivity analyses using binary anti-CCP2 cut-offs (> 10–250 and > 250 U/ml) showed that baseline characteristics were similar to those from the anti-CCP2+ quartile analysis in patients initiating treatment with abatacept or a TNFi (Supplementary Tables S3 and S4). Among patients treated with abatacept, there was a significant improvement in unadjusted CDAI score at 6 months with the higher cut-off versus the lower cut-off (*p* < 0.05; Supplementary Table S5). However, the change in CDAI score was no longer statistically significant after

adjusting for baseline CDAI score and other covariates (*p* = 0.186; Table 3). In addition, among patients treated with abatacept, patient-reported pain, fatigue, PtGA, and mHAQ score each showed a significant association with binary anti-CCP2+ status in both unadjusted (*p* < 0.05; Supplementary Table S5) and adjusted (*p* < 0.05; Table 3) models. Among patients initiating treatment with a TNFi, there was no significant association with binary anti-CCP2+ status for change in CDAI score or PROs in either the unadjusted (Supplementary Table S6) or adjusted (Table 4) models.

Table 2 Baseline characteristics among TNFi-treated patients by anti-CCP2 quartile

Characteristic	Anti-CCP2+, U/ml ^a				<i>p</i> value ^b
	Quartile 1 (<i>n</i> = 36)	Quartile 2 (<i>n</i> = 34)	Quartile 3 (<i>n</i> = 32)	Quartile 4 (<i>n</i> = 35)	
Age, years	60.1 (11.5)	60.6 (12.7)	55.3 (13.3)	58.9 (12.5)	0.306
Female sex, <i>n</i> (%)	28 (77.8)	26 (76.5)	24 (75.0)	25 (71.4)	0.946
White race, <i>n</i> (%) ^c	NP ^d	NP ^d	24 (82.8)	NP ^d	NP ^d
Hispanic ethnicity, <i>n</i> (%) ^e	NP ^d	NP ^d	NP ^d	5 (15.2)	NP ^d
BMI, kg/m ²	30.7 (9.0)	27.4 (6.0)	28.6 (6.5)	29.3 (7.7)	0.611
Smoking history, <i>n</i> (%) ^f					
Never	21 (58.3)	12 (36.4)	16 (53.3)	14 (40.0)	0.213
Former/Current	15 (41.7)	21 (63.6)	14 (46.7)	21 (60.0)	
Comorbidities, <i>n</i> (%)					
Hypertension	17 (47.2)	10 (29.4)	9 (28.1)	15 (42.9)	0.263
Diabetes	4 (11.1)	3 (8.8)	1 (3.1)	2 (5.7)	0.683
Malignancy	1 (2.8)	2 (5.9)	1 (3.1)	3 (8.6)	0.748
Serious infections	3 (8.3)	4 (11.8)	2 (6.3)	4 (11.4)	0.885
Cardiovascular	4 (11.1)	3 (8.8)	6 (18.8)	4 (11.4)	0.681
Duration of RA, years	14.1 (9.6)	14.7 (13.7)	12.1 (8.2)	13.1 (9.3)	0.897
CDAI score	29.1 (10.8)	32.3 (12.5)	26.2 (11.2)	29.8 (13.7)	0.255
TJC28	11.3 (7.2)	11.9 (6.3)	9.6 (6.1)	11.0 (7.1)	0.537
SJC28	7.6 (4.6)	9.4 (5.9)	7.4 (4.2)	7.6 (6.0)	0.427
CRP, mg/l	15.5 (30.8)	17.1 (24.3)	12.2 (14.2) ^g	13.3 (27.0)	0.966
DAS28 (CRP)	4.9 (1.1)	5.1 (1.0)	4.6 (1.0) ^g	4.9 (1.2)	0.315
RF, IU/ml ^h	111.0 (173.2)	182.9 (353.3)	342.0 (559.1)	617.8 (1042.6)	< 0.0001
mHAQ	0.5 (0.4)	0.6 (0.6)	0.3 (0.3)	0.8 (0.6)	0.003
PtGA (VAS 0–100)	50.4 (23.3)	57.1 (22.5)	40.5 (27.2)	59.4 (24.8)	0.012
PGA (VAS 0–100)	52.1 (14.5)	52.9 (17.4)	51.3 (19.8)	51.9 (19.2)	0.987
Patient-reported pain (VAS 0–100)	54.8 (28.3)	56.8 (26.4)	45.4 (26.3)	60.8 (26.4)	0.136
Patient-reported fatigue (VAS 0–100)	51.6 (29.4)	51.7 (29.7)	47.4 (29.8)	58.2 (28.7)	0.463

Table 2 continued

Characteristic	Anti-CCP2+, U/ml ^a				<i>p</i> value ^b
	Quartile 1 (<i>n</i> = 36)	Quartile 2 (<i>n</i> = 34)	Quartile 3 (<i>n</i> = 32)	Quartile 4 (<i>n</i> = 35)	
Prior biologic use, <i>n</i> (%)					
1	14 (38.9)	15 (44.1)	13 (40.6)	14 (40.0)	0.821
2	13 (36.1)	8 (23.5)	13 (40.6)	12 (34.3)	
3+	9 (25.0)	11 (32.4)	6 (18.8)	9 (25.7)	
Prior TNFi use, <i>n</i> (%) ⁱ	34 (94.4)	34 (100.0)	31 (96.9)	34 (97.1)	0.747
Prior non-TNFi use, <i>n</i> (%) ⁱ	15 (41.7)	14 (41.2)	9 (28.1)	13 (37.1)	0.645
Prior abatacept use, <i>n</i> (%) ⁱ	11 (30.6)	12 (35.3)	6 (18.8)	11 (31.4)	0.485
Current MTX use, <i>n</i> (%) ^j	21 (58.3)	18 (52.9)	20 (62.5)	19 (54.3)	0.868

Data are mean (SD), unless otherwise stated

^a Anti-CCP2 quartiles: quartile 1, 11–94 U/ml; quartile 2, 95–296 U/ml; quartile 3, 297–876 U/ml; and quartile 4, > 876 U/ml

^b Differences between baseline measures of disease activity across anti-CCP2+ quartiles were assessed using one-way analysis of variance for DAS28 (CRP) and PGA and the Kruskal–Wallis test for nonparametric measures (CDAI, TJC28, SJC28, CRP, RF, mHAQ and PtGA)

^c *n* = 29 and 34 for quartiles 3 and 4, respectively

^d Patient numbers < 5 (or < 5 derived through subtraction from the total) or a percentage of 0% or 100% (indicative of certainty) were suppressed (not presented, NP) due to potential loss of confidentiality

^e *n* = 35 and 33 for quartiles 1 and 4, respectively

^f *n* = 33 and 30 for quartiles 2 and 3, respectively

^g *n* = 31

^h *n* = 35, 32, 31, and 34 for quartiles 1 to 4, respectively

ⁱ Prior use reflects all prior biologic history; may not sum to 100%

^j At time of biologic initiation

Anti-CCP2 anti-cyclic citrullinated peptide 2, *BMI* body mass index, *CDAI* Clinical Disease Activity Index, *CRP* C-reactive protein, *DAS28 (CRP)* Disease Activity Score in 28 joints (C-reactive protein), *mHAQ* modified Health Assessment Questionnaire, *MTX* methotrexate, *PGA* physician global assessment; *PtGA* patient global assessment, *RA* rheumatoid arthritis, *RF* rheumatoid factor, *SD* standard deviation, *SJC28* swollen joint count in 28 joints, *TJC28* tender joint count in 28 joints, *TNFi* tumor necrosis factor inhibitor/s, *VAS* visual analog scale

DISCUSSION

In this prospective, adjusted analysis of biologic-experienced patients with RA, higher (versus lower) anti-CCP2 concentrations at baseline were associated with a numerically greater improvement in CDAI score and significantly greater improvements in PROs after 6 months of treatment with abatacept. No association between baseline anti-CCP2 concentration and treatment response was observed in patients treated with a TNFi.

Baseline disease characteristics (CDAI score, SJC28, CRP, DAS28 [CRP], RF, mHAQ score, physician global assessment) were significantly different between anti-CCP2 levels for patients initiating abatacept. For patients initiating a TNFi, RF, mHAQ score and PtGA were significantly different between anti-CCP2 levels. The association of increased baseline disease characteristics with a higher baseline anti-CCP2 titer (compared with a lower baseline anti-CCP2) in patients is consistent with previous research where patients with high levels of

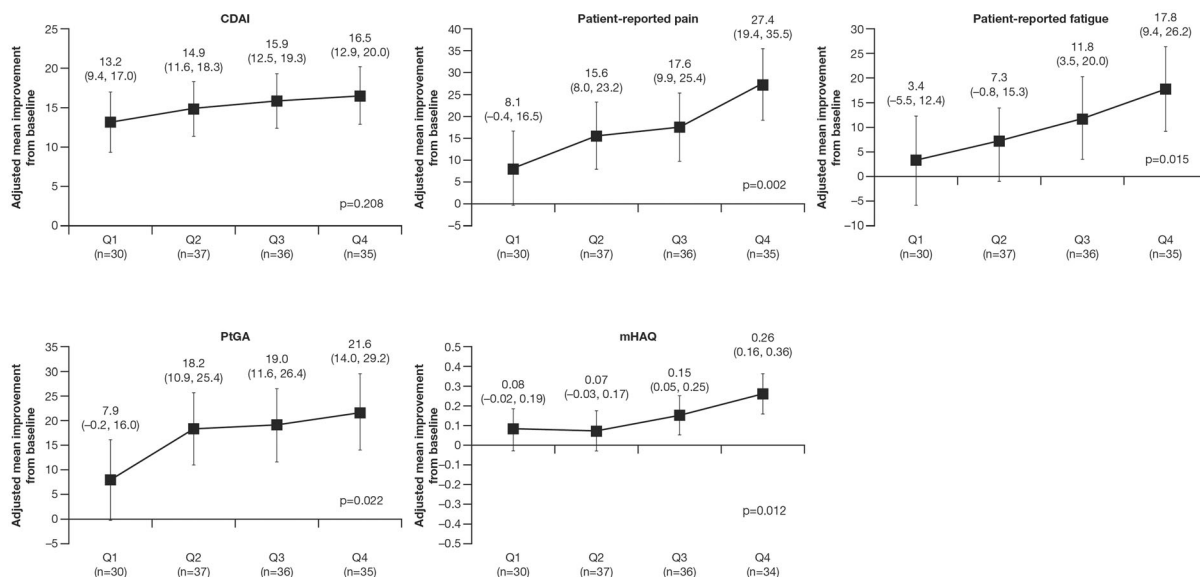


Fig. 2 Adjusted mean improvement from baseline* in CDAI score and PROs for abatacept-treated patients, by anti-CCP2 quartile. Data are mean (95% CI); quartile 1 (n = 30) was used as the reference. *Adjusted for age, sex, baseline CDAI score or PROs, Charlson Comorbidity

Index and current methotrexate use. *Anti-CCP2* anti-cyclic citrullinated peptide 2, *CDAI* Clinical Disease Activity Index, *CI* confidence interval, *mHAQ* modified Health Assessment Questionnaire, *PRO* patient-reported outcome, *PtGA* patient global assessment, *Q* quartile

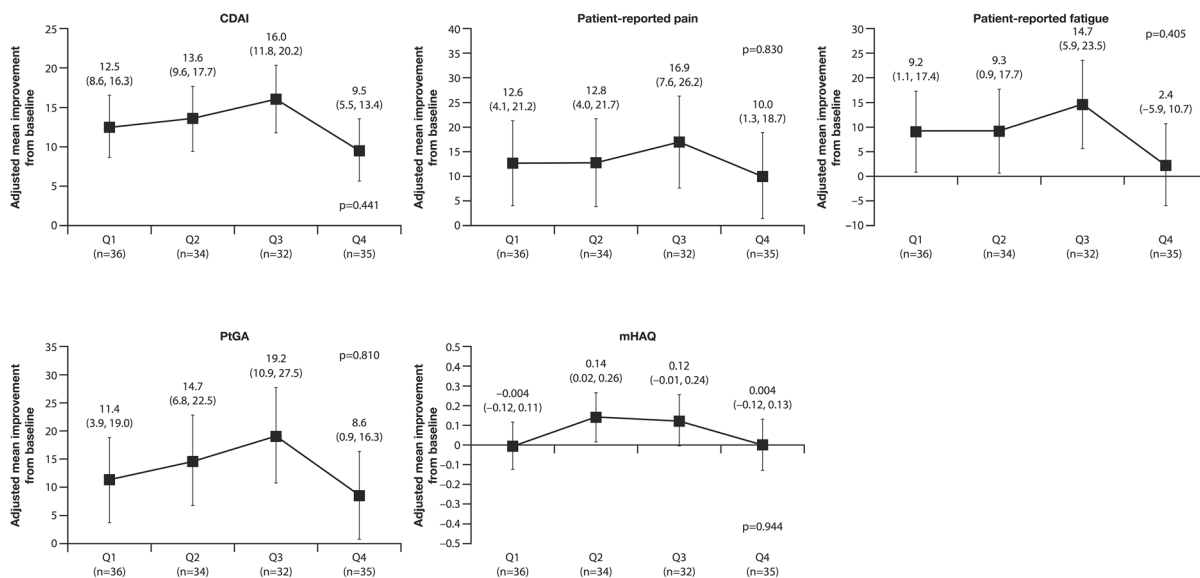


Fig. 3 Adjusted mean improvement from baseline* in CDAI score and PROs for TNFi-treated patients, by anti-CCP2 quartile. Data are mean (95% CI); quartile 1 (n = 36) was used as the reference. *Adjusted for age, sex, baseline CDAI score or PROs, Charlson Comorbidity

Index and current methotrexate use. *Anti-CCP2* anti-cyclic citrullinated peptide 2, *CDAI* Clinical Disease Activity Index, *CI* confidence interval, *mHAQ* modified Health Assessment Questionnaire, *PRO* patient-reported outcome, *PtGA* patient global assessment, *Q* quartile, *TNFi* tumor necrosis factor inhibitor/s

Table 3 Adjusted mean improvement from baseline in CDAI score and PROs for abatacept-treated patients, by binary anti-CCP2 cut-off

Full adjusted outcome	Anti-CCP2+ > 10–250 U/ml (n = 61)	Anti-CCP2+ > 250 U/ml (n = 77)	p value
Δ CDAI, mean (95% CI)	13.9 (11.2, 16.5)	16.3 (13.9, 18.6)	0.186
Δ mHAQ, mean (95% CI)	0.07 (− 0.01, 0.14)	0.20 (0.13, 0.27) ^a	0.014
Δ patient-reported pain, mean (95% CI)	11.1 (5.1, 17.0)	22.6 (17.3, 27.9)	0.005
Δ patient-reported fatigue, mean (95% CI)	4.6 (−1.7, 10.9)	14.8 (9.2, 20.3)	0.018
Δ patient global assessment, mean (95% CI)	12.6 (6.9, 18.3)	20.5 (15.5, 25.6)	0.042

Anti-CCP2 > 10–250 U/ml was used as the reference. Adjusted for age, sex, baseline CDAI score or PROs, comorbidity index and current methotrexate use.

Anti-CCP2 anti-cyclic citrullinated peptide 2, *CDAI* Clinical Disease Activity Index, *CI* confidence interval, *mHAQ* modified Health Assessment Questionnaire, *PRO* patient-reported outcome

^a n = 76 for mHAQ

autoantibodies, including ACPA, have higher disease activity and greater impairment of physical function [3, 4, 6].

In the adjusted model, the CDAI score was numerically improved and PROs were significantly improved after 6 months across both increasing anti-CCP2 quartiles and binary cut-offs for patients treated with abatacept. There were no significant improvements in CDAI score or PROs across anti-CCP2 quartiles or binary cut-offs for patients initiating a TNFi. Data from previous clinical trials have shown an

Table 4 Adjusted mean improvement from baseline in CDAI score and PROs for TNFi-treated patients, by binary anti-CCP2 cut-off

Full adjusted outcome	Anti-CCP2+ > 10–250 U/ml (n = 63)	Anti-CCP2+ > 250 U/ml (n = 74)	p value
Δ CDAI, mean (95% CI)	13.1 (10.2, 16.1)	12.6 (9.8, 15.3)	0.776
Δ mHAQ, mean (95% CI)	0.08 (− 0.01, 0.17)	0.04 (− 0.04, 0.13)	0.563
Δ patient-reported pain, mean (95% CI)	13.0 (6.5, 19.4)	13.1 (7.1, 19.0)	0.985
Δ patient-reported fatigue, mean (95% CI)	10.7 (4.5, 16.9)	7.1 (1.4, 12.8)	0.404
Δ patient global assessment, mean (95% CI)	13.4 (7.7, 19.2)	13.2 (7.9, 18.5)	0.956

Anti-CCP2 > 10–250 U/ml was used as the reference. Adjusted for age, sex, baseline CDAI score or PROs, comorbidity index and current methotrexate use.

Anti-CCP2 anti-cyclic citrullinated peptide 2, *CDAI* Clinical Disease Activity Index, *CI* confidence interval, *mHAQ* modified Health Assessment Questionnaire, *PRO* patient-reported outcome, *TNFi* tumor necrosis factor inhibitor/s

association between ACPA+ status and clinical response to abatacept, in combination with MTX, in both bDMARD-naïve patients (AMPLE) [13] and MTX-naïve patients with very early RA (AVERT; Assessing Very Early Rheumatoid arthritis Treatment) [18, 19]. In a post hoc analysis of the AMPLE trial, a more pronounced treatment effect was observed in patients with higher baseline anti-CCP titers compared with lower titers [13]. Furthermore, some studies suggest that treatment with abatacept in combination with MTX may lead to a reduction in autoantibody levels over time, and that conversion to ACPA seronegative status may be

associated with better clinical outcomes [20, 21]. Consistent data from real-world studies have also been observed. In the ACTION (AbataCepT In rOutiNe clinical practice) study, RF/anti-CCP+ status was associated with greater clinical effectiveness and retention on treatment in patients receiving abatacept [22]. Analyses from the Corrona RA Registry [14] and a pan-European registry [23] have shown greater effectiveness with abatacept in patients who were ACPA+ or ACPA+/RF+, respectively. Additionally, a meta-analysis of 18 studies showed ACPA positivity to be associated with better EULAR responses in patients receiving abatacept but not in those receiving a TNFi [24].

A major genetic risk factor for RA is the shared epitope (SE), which is located on the β 1 chain of the human leukocyte antigen-DR major histocompatibility complex Class II cell-surface receptor [25, 26]. SE is associated with the production of ACPAs in patients with RA (the majority of patients with anti-CCP+RA also exhibit SE) [27], and studies have shown an association between SE and increased disease activity [26, 28, 29]. Additionally, abatacept has significantly higher clinical efficacy among patients with SE+RA compared with SE–RA [30, 31]. Ongoing studies are exploring the impact of SE on treatment outcomes.

This study has several strengths. Corrona is a US-based registry, which includes a large number of patients with RA with validated outcome measures and is representative of patients found in clinical practice in the US [32]. Previous analyses have compared Medicare patients with RA enrolled in Corrona to those not part of the registry and have found similar demographic and comorbidity characteristics, supporting the generalizability of the Corrona registry [32]. Although the Corrona RA Registry patient population is managed by rheumatologists across the US, with no centralized laboratory testing for serological status, all patients derived from the CERTAIN study had centralized measurement of CCP status. Data were collected prospectively at patient visits; therefore, differential bias in responses by treatment assignment were not expected. The results from the current study complement those of previous studies that have provided data regarding the

clinical effectiveness of abatacept in patients with RA who were anti-CCP+ at baseline [13, 14, 18, 19, 21, 22, 30, 31]. Our study is methodologically strong: In contrast to previous studies [14, 18, 19, 21, 22, 30, 31], patients were grouped into anti-CCP2 quartiles (quartile 1, 11–94 U/ml; quartile 2, 95–296 U/ml; quartile 3, 297–876 U/ml; and quartile 4, > 876 U/ml) and we sought to correlate outcomes by quartile. We also performed a sensitivity analysis using the typical laboratory cut-off points for baseline anti-CCP2 available to physicians (> 10–250 U/ml and > 250 U/ml). Additionally, we compared the utility of anti-CCP status with TNFi treatment. As such, our study provides real-world, clinically relevant evidence that may assist clinicians in making more informed individualized treatment decisions taking into account the effect of ACPA status on treatment response [4].

As with all real-world studies, there are limitations to our study. Patients included in the registry were diagnosed by different rheumatologists across the US. Treatment selection in any real-world setting is not random, as physicians are influenced by a number of factors when prescribing treatments, including individual patient profiles. However, to help offset such potential selection bias, propensity score matching was used in the present study to identify patients with similar disease activity within each line of therapy prior to anti-CCP2 testing. The study sample size was relatively small, and the duration of follow-up (6 months) was relatively short. Treatments were not blinded; as such, patients may have been aware if they were treated with a new class of biologic, which may have influenced PROs and CDAI scores.

CONCLUSIONS

In conclusion, among bDMARD-experienced patients with RA initiating treatment with abatacept, but not a TNFi, higher anti-CCP2 concentrations at baseline were associated with numerically, but not statistically, greater improvements in CDAI score and significantly greater improvement in PROs in the adjusted

model after 6 months of treatment. This association was not observed in a similar sample size of patients initiating treatment with TNFi. The association between genetic factors and treatment outcome in patients with RA is under further investigation.

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statisticians completed all analyses. All authors drafted the article or critically revised the article for important intellectual content. All authors were involved in the decision to submit the article for publication and approved the final version of the article for submission.

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Compliance with Ethics Guidelines. The study was conducted in accordance with the Declaration of Helsinki and International Conference on Harmonization Good Clinical Practice. The protocol and patient's informed consent received institutional review board/independent ethics committee approval prior to initiation of the study. All participating investigators were required to obtain full board approval for conducting research involving human subjects. Sponsor approval and continuing review was obtained through a central IRB (New England Independent Review Board, NEIRB No. 120160610). For academic

investigative sites that did not receive a waiver to use the central IRB, approval was obtained from the respective governing IRBs and documentation of approval was submitted to the Sponsor prior to initiating any study procedures. All registry subjects were required to provide written informed consent prior to participating.

Data Availability. The Corrona RA Registry dataset is based on a large US multicenter study adhering to a number of institutional review boards, with complex logistics. Patients did not provide consent to raw data sharing during the data collection for this purpose, and the CorEvitas data sharing policies do not permit raw data sharing for this purpose. An aggregated limited dataset from the current analyses is available to qualified investigators with an approved protocol. Data requests may be sent to CorEvitas, represented by Dr. Jeffrey D. Greenberg, MD, MPH, NYU School of Medicine, New York, NY, e-mail jgreenberg@corevitas.com.

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