The relationship between infliximab concentrations, antibodies to infliximab and disease activity in Crohn’s disease

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ABSTRACT
Objective Although low infliximab trough concentrations and antibodies to infliximab (ATI) are associated with poor outcomes in patients with Crohn’s disease (CD), the clinical relevance of ATI in patients with adequate infliximab concentrations is uncertain. We evaluated this question using an assay sensitive for identification of ATI in the presence of infliximab.

Design In an observational study, 1487 trough serum samples from 483 patients with CD who participated in four clinical studies of maintenance infliximab therapy were analysed using a fluid phase mobility shift assay. Infliximab and ATI concentrations most discriminating for remission, defined as a C-reactive protein concentration of <5 mg/L, were determined by receiver operating characteristic curves. A multivariable regression model evaluated these factors as independent predictors of remission.

Results Based upon analysis of 1487 samples, 77.1% of patients had detectable and 22.9% had undetectable infliximab concentrations, of which 9.5% and 71.8%, respectively, were positive for ATI. An infliximab concentration of >2.79 μg/mL (area under the curve (AUC)=0.681; 95% CI 0.632 to 0.731) and ATI concentration of >3.15 U/mL (AUC=0.632; 95% CI 0.589 to 0.676) were associated with remission. Multivariable analysis showed that concentrations of both infliximab trough (OR 1.8; 95% CI 1.3 to 2.5; p<0.001) and ATI (OR 0.57; 95% CI 0.39 to 0.81; p=0.002) were independent predictors of remission.

Conclusions The development of ATI increases the probability of active disease even at low concentrations and in the presence of a therapeutic concentration of drug during infliximab maintenance therapy. Evaluation of strategies to prevent ATI formation, including therapeutic drug monitoring with selective infliximab dose intensification, is needed.

INTRODUCTION
Over the past decade, considerable evidence has accumulated that sensitisation to biological drugs is an important clinical problem. In 2003, Baert and colleagues performed a prospective cohort study of 125 patients with Crohn’s disease (CD), which identified that the presence of an antibody to infliximab (ATI) concentration ≥8 μg/mL was associated

Significance of this study
What is already known on this subject?
▸ C-reactive protein is a marker of disease activity in patients with Crohn’s disease.
▸ Low or undetectable serum trough concentrations of infliximab (IFX) are associated with worse clinical outcomes.
▸ Antibodies to infliximab (ATI) increase the clearance of drug and are an important cause of low-serum IFX trough concentrations.

What are the new findings?
▸ Based on a combined analysis of patient-level data from four studies that evaluated 483 patients with Crohn’s disease using a fluid phase mobility shift assay, IFX trough concentrations >2.79 μg/mL during maintenance therapy were associated with remission as measured by C-reactive protein concentration.
▸ Detectable ATI were associated with greater disease activity as measured by C-reactive protein concentration even in the presence of an adequate IFX trough concentration.
▸ ATI impair the drug’s activity through an alternative mechanism than only by affecting its clearance.

How might it impact on clinical practice in the foreseeable future?
▸ These results support the role of therapeutic drug monitoring in patients with Crohn’s disease receiving IFX. We speculate that if the presence of ATI has a negative effect on the pharmacodynamics of IFX independent of the trough concentration, other pharmacokinetics determinants such as the area under the curve and maximum serum concentration (Cmax) of the drug might also be important predictors of clinical efficacy. Therapeutic drug monitoring with dose intensification in patients with undetectable or low IFX trough concentrations should be evaluated as a strategy to prevent development of ATI.
with shorter time to relapse compared with patients with low-titre ATL. Furthermore, patients with AT1 had a greater risk of hypersensitivity reactions. Although subsequent studies have attempted to define an AT1 concentration that correlates with clinical outcomes, differences in assay design, the sometimes transient nature of AT1 expression and the inability of conventional assays to measure AT1 in the presence of drug have made identification of such a relationship problematic. Nevertheless, a recent meta-analysis that evaluated 1378 patients with IBD who received treatment with infliximab (IFX) showed that the pooled risk ratio for loss of clinical response in patients with AT1 was 3.2 (95% CI 2.0 to 4.9; p<0.0001) in comparison with control patients without AT1. Considerable evidence exists that higher IFX concentrations are associated with worse clinical outcomes. As a consequence, the use of therapeutic drug monitoring (TDM) has been advocated to improve clinical decision making in patients with a secondary loss of efficacy in patients with IBD. For example, in ACCENT I, a multicentre trial that evaluated IFX induction therapy in patients with active CD, patients who failed to respond to therapy had lower serum IFX concentrations than those with a sustained response (1.9 and 4.0 μg/mL, respectively; p=0.03). Collectively, these data indicate that both presence of AT1 and low IFX concentrations are associated with worse clinical outcomes. As a consequence, the use of therapeutic drug monitoring (TDM) has been advocated to improve clinical decision making in patients with a secondary loss of response to IFX. These observations also hold out the possibility that dose optimisation based on TDM during induction, prior to recurrence of symptoms, and possibly prior to AT1 formation might result in better outcomes. However, analysis of the impact of AT1 on IFX pharmacokinetics has been hindered by the inability of most assays to detect AT1 in the presence of IFX. Thus, it has not been possible to determine whether the negative effects of AT1 on clinical efficacy are solely the consequence of low-titre drug concentrations (i.e. whether patients with AT1 and low drug concentrations are entirely a subset of patients with low-drug concentrations) or whether AT1 might also impair other measures of drug exposure, such as the maximum serum concentration (Cmax) or the area under the concentration time curve (AUC), which are critical to the pharmacodynamics of IFX (even when drug is present). Therefore, an enhanced understanding of the relevance of AT1 in patients with a detectable serum concentration of IFX is potentially important.

Based on these considerations, we used a homogenous mobility shift drug-tolerant assay able to measure AT1 in the presence of IFX to evaluate the relationship between serum IFX trough concentration, AT1 concentration and disease activity in a large population of patients with CD.

MATERIALS AND METHODS

Study objectives

Our study had two objectives. First, we wished to determine the IFX trough concentration during maintenance therapy that best correlated with C-reactive protein (CRP)-defined remission as a surrogate for clinical remission. A CRP concentration of ≤5 mg/L was defined as remission. CRP was chosen to define disease activity, in distinction to clinical symptoms, because numerous studies have demonstrated that a poor correlation exists between symptoms and inflammation defined by colonoscopy, serum acute-phase reactants such as CRP or faecal biomarkers such as calprotectin. A second goal was to evaluate the relationship between CRP concentration as a measure of disease activity, the IFX trough concentration and the presence of AT1 in four distinct groups of patients: (1) IFX trough concentration ≥3 μg/mL and AT1 negative; (2) IFX trough concentration <3 μg/mL and AT1 negative; (3) IFX trough concentration ≥3 μg/mL and AT1 positive; and (4) IFX trough concentration <3 μg/mL and AT1 positive. The rationale for these sub-group analyses is based on the hypothesis that the indirect correlation between IFX and disease activity differs by AT1 status. Specifically, we wished to determine whether AT1 was a predictor of disease activity independent of IFX trough concentration.

Study design and conduct

Assays were performed on 1487 trough serum samples from 483 patients with CD treated with maintenance IFX therapy who had previously participated in one of four prospective randomised controlled trials or cohort studies: the COMMIT trial, the Leuven transient AT1 study, the Canadian Multicentre study and the Leuven antimetabolite withdrawal study. The COMMIT study examined the effect of transient AT1 formation on IFX pharmacokinetics and clinical response. The Canadian Multicentre study was a prospective analysis of the relationship between IFX concentrations, AT1 concentrations and clinical disease activity in patients receiving maintenance therapy. The fourth study evaluated the efficacy of continued antimalbolite therapy in patients, after receiving combination therapy with IFX during the first six months of therapy. These parent studies enrolled adult patients with moderate-to-severe CD. Demographic data collected included the age at diagnosis, the age at first IFX infusion and the use of concomitant treatment with azathioprine or methotrexate, as well as the serum sample collection date. Each of these studies collected trough serum samples for analysis of serum IFX, AT1 and CRP concentration. Serum CRP was measured with the Vascular Injury Panel 2 (human) V-PLEX Kit (Meso Scale Discovery, Rockville, Maryland, USA) with a lower limit of quantitation (LLOQ) of 0.7 μg/L. Serum IFX and AT1 were measured using a previously described commercially available homogeneous mobility shift assay (Prometheus Laboratories, San Diego, California, USA). The LLOQ for the presence of IFX and AT1 in this assay was 0.98 μg/mL and 3.13 U/mL, respectively. By using a preanalytical acid dissociation step, this drug-tolerant assay can detect AT1 in the presence of up to 60 μg/mL IFX. To reduce variability, all of the samples were reanalysed in batch.

Statistical analyses

Receiver operating characteristic (ROC) curves were used to determine the IFX and AT1 concentration thresholds that best discriminated remission, defined as a CRP concentration of ≤5 mg/L. The AUC of the ROC curve along with the 95% CI were adjusted for repeated observations per subject. A multivariable logistic model assessed multiple clinical and laboratory factors as independent predictors of a CRP concentration of >5 mg/L. Factors included in the modelling process were study, gender, age at diagnosis, age at collection, days since first IFX infusion, age at first IFX, methotrexate use, azathioprine use, IFX concentration <3 μg/mL, AT1 concentration ≥3.13 U/mL and an IFX–AT1 interaction term. Factors not significant at the 0.05 level were removed by backwards elimination. A statistical model was validated in a separate training set.
multivariable logistic model evaluated IFX concentration <3 µg/mL, ATI concentration ≥3.13 U/mL and interval (days in-between samples) in a ‘current’ sample as independent predictors of a CRP concentration of >5 mg/L in a ‘future’ sample. Only samples with an interval of 56±14 days in-between ‘current’ and ‘future’ sample were taken into account for each patient. Two-way and three-way interactions between all the variables IFX, ATI and interval between samples were included in the model to investigate whether the effects of IFX and ATI were dependent on each other and dependent on time. The model was adjusted for study, days between first IFX and first sample and baseline CRP in the ‘current’ sample. All models were adjusted for multiple observations per subject, as a random effect.

Ethical considerations
All serum samples analysed were obtained during the execution of four separate parent clinical trials. The informed consent process complied with the International Conference on Harmonisation—Good clinical practice and all applicable regulatory requirement(s). The consent of subjects included the use of the collected data and serum for other medical purposes. Therefore, additional consent for the current study was not obtained. Each study was reviewed by a research ethics board at the respective site where the initial study was conducted (Western University for the COMMIT Trial and the Canadian Multicentre Trial, and the University of Leuven for the transient ATI study and the antimetabolite withdrawal study). All subject information used in this study was de-identified with respect to the subject identification number and investigational site. The serum samples were assayed following conclusion of the four original parent studies and did not affect patients’ treatment and/or well-being.

RESULTS
Study population
Samples from 483 patients were analysed, of whom 236 (48.9%) were men. The mean age of the patients at diagnosis was 28.5±12.9 years. Eighty-five patients (17.6%) received concomitant treatment with methotrexate while 140 patients (29%) received concomitant azathioprine. A total of 1487 serum samples were analysed with a median of 2 (range 1–26) samples per patient. Demographic information and sample characteristics are summarised in Table 1.

Overall the median (IQR) CRP concentration of the sample cohort was 3.6 (IQR 1.0–12.7) mg/L. Of the 1487 serum samples, 6.5% (n=96) were IFX and ATI negative, 69.8% (n=1038) were IFX positive and ATI negative, 16.4% (n=244) were IFX negative and ATI positive and 7.3% (n=109) were both IFX and ATI positive (Figure 1). Samples that were positive for ATI had a significantly lower likelihood of having detectable IFX compared with ATI-negative samples (OR 0.041; 95% CI 0.030 to 0.056; p<0.001).

Outcomes
Based on ROC curve analysis, an IFX concentration cut-off of >2.79 µg/mL (AUC=0.681; 95% CI 0.632 to 0.731) was associated with remission (77.6% specificity, 52.5% sensitivity), as seen in Figure 2. When the IFX trough concentration was divided into quartiles, a concentration-dependent effect was observed on CRP concentration (p<0.001) (Figure 3A) and a linear trend was observed (p<0.001).

Based on ROC curve analysis, an ATI concentration cut-off of <3.15 U/mL (AUC=0.632; 95% CI 0.589 to 0.676) was associated with remission (87.4% specificity, 38.0% sensitivity). Only taking ATI-positive samples into account (n=342), after stratification per quartile, a concentration-dependent trend was observed on CRP concentration (p=0.07) (Figure 3B). Moreover, a significant linear relationship was found, explaining that at least a part of the association with CRP can be explained by a trend across the four ATI quartiles (p=0.025). Combining both factors IFX and ATI as predictors of disease activity increased the AUC of the ROC curve to 0.701.

Overall, 117 out of 483 patients (24.2%) had detectable ATI on at least one time point during follow-up. The median CRP concentration was lower in ATI-negative samples compared with ATI-positive samples (respectively 2.4 vs. 11.9 mg/L; p<0.001). In ATI-negative samples stratified according to IFX concentration, the median CRP concentration was significantly lower when the serum IFX concentration was ≥3 µg/mL compared with <3 µg/mL (respectively 2.0 vs. 6.0 mg/L; p<0.001). However, in ATI-positive samples, the median CRP concentration was similar in these two groups (respectively 11.9 vs. 11.6 mg/L), although significantly different after correcting for multiple observations per patient (p=0.019) (Figure 4). In an exploratory analysis, omitting the samples from the Leuven transient ATI study, we found that CRP concentrations were numerically lower but the same effect was observed, in that CRP concentrations were elevated in ATI-positive samples despite an IFX concentration of ≥3 µg/mL (7.9 mg/L) or <3 µg/mL (9.2 mg/L) (p=0.004).

Univariate analyses demonstrated significant associations between originating study, presence of ATI and IFX concentration and the presence of active disease (CRP concentration >5 mg/L). However, multivariable analysis retained IFX <3 µg/mL (OR 1.8; 95% CI 1.3 to 2.5; p<0.001) and detectable ATI (≥3.15 U/mL) (OR 1.8; 95% CI 1.2 to 2.6; p=0.002) as the only independent predictors of this outcome. The interaction between IFX and ATI concentrations was also evaluated and was not significant (p=0.116). For the multivariable logistic model evaluating IFX, ATI and interval (between samples) in a ‘current’ sample as independent predictors of a CRP concentration of >5 mg/L in a ‘future’ sample, there were 454 patients with a complete set of IFX, ATI and CRP on at least two consecutive time points with no more than 56±14 days in-between samples. After correcting for study, days between first IFX and first sample and baseline CRP we observed that both IFX <3 µg/mL (p=0.035) and ATI ≥3.13 U/mL (p=0.006) in a ‘current’ sample were able to predict a CRP concentration of >5 mg/L in a ‘future’ sample. The interval between ‘current’ and ‘future’ sample was not found to be a predictor of CRP (p=0.466). IFX and ATI were found to independently predict CRP and independent of the time in-between samples as none of the two-way or three-way interactions between the variables IFX, ATI and interval between samples were significant.

DISCUSSION
This study, which evaluated 1487 trough serum samples from 483 patients with CD who received maintenance IFX therapy in four parent studies, to our knowledge is the largest evaluation of the relationship between IFX trough concentrations, ATI and clinical efficacy that has been reported. Importantly, the use of a drug-tolerant assay allowed us to assess the relevance of detection of ATI in the presence of an adequate IFX trough concentration.

These results confirm the existence of a strong inverse correlation between IFX trough concentration and CRP-defined remission. Notably, a linear test for trend was highly significant
for these factors supporting the presence of an ‘exposure-effect relationship’. In ROC curve analysis, an IFX trough concentration of \( \leq 2.79 \, \mu\text{g/mL} \) best discriminated disease activity (OR 2.13; 95% CI 1.54 to 2.96), suggesting that this value can be used as a benchmark in clinical practice; however, confirmatory studies are needed. Prospectively acquired data re-confirming this threshold would provide additional validity.

This study also provides new insights into the relationship between ATI and disease activity. Use of the drug-tolerant fluid phase mobility shift assay allowed us to assess the role of ATI in the presence of IFX. A strong correlation was observed between the presence of ATI and the likelihood of active disease defined by a CRP concentration of >5 mg/L. Interestingly, a linear test for trend for CRP concentration across different ATI quartiles was significant, suggesting a ‘causal relationship’ between both these factors. ROC curve analysis revealed an ATI concentration of \( \geq 3.15 \, \text{U/mL} \) best discriminated active disease (OR 2.22; 95% CI 1.56 to 3.15). However, it was notable that in the subset of samples that were positive for ATI and also had an IFX trough concentration \( \geq 3 \, \mu\text{g/mL} \), a similar proportion of samples had CRP >5 mg/L compared with the subset of samples that were ATI positive and had an IFX trough concentration <3 \( \mu\text{g/mL} \).

Table 1  Patient demographics within each study and the combined data set*

<table>
<thead>
<tr>
<th>Patients</th>
<th>COMMIT</th>
<th>Canadian multicentre</th>
<th>Transient ATI</th>
<th>IMM withdrawal</th>
<th>Combined</th>
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</thead>
<tbody>
<tr>
<td>Patients—no.</td>
<td>49</td>
<td>327</td>
<td>25</td>
<td>82</td>
<td>483</td>
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<tr>
<td>Men—no. (%)</td>
<td>28 (57)</td>
<td>157 (48)</td>
<td>11 (44)</td>
<td>40 (49)</td>
<td>236 (49)</td>
</tr>
<tr>
<td>Age at diagnosis</td>
<td>29.5±13.3</td>
<td>29.3±13.2</td>
<td>26.8±11.6</td>
<td>25.5±11.3</td>
<td>28.5±12.9</td>
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<tr>
<td>Age at first sample collection</td>
<td>39.0±14.0</td>
<td>39.9±12.9</td>
<td>40.3±12.7</td>
<td>35.7±13.8</td>
<td>39.1±13.2</td>
</tr>
<tr>
<td>Age at first infliximab</td>
<td>38.6±14.1</td>
<td>38.5±13.0</td>
<td>39.9±12.5</td>
<td>33.7±13.8</td>
<td>37.7±13.3</td>
</tr>
<tr>
<td>Patients ever IFX ( \geq 3 , \mu\text{g/mL} )—no. (%)</td>
<td>44 (90)</td>
<td>237 (73)</td>
<td>15 (60)</td>
<td>75 (92)</td>
<td>371 (77)</td>
</tr>
<tr>
<td>Patients ever ATI positive—no. (%)</td>
<td>9 (18)</td>
<td>73 (22)</td>
<td>23 (92)</td>
<td>12 (15)</td>
<td>117 (24)</td>
</tr>
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<td>Patients ever receiving MTX—no. (%)</td>
<td>23 (47)</td>
<td>30 (9)</td>
<td>9 (36)</td>
<td>23 (28)</td>
<td>85 (18)</td>
</tr>
<tr>
<td>Patients ever receiving AZA—no. (%)</td>
<td>0 (0)</td>
<td>81 (25)</td>
<td>0 (0)</td>
<td>59 (72)</td>
<td>140 (29)</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Serum samples</th>
<th>Serum samples—no.</th>
<th>Samples per subject†</th>
<th>Time between first infliximab and sample—months†</th>
<th>Serum samples IFX ( \geq 3 , \mu\text{g/mL} )—no. (%)</th>
<th>Serum samples ATI positive—no. (%)</th>
<th>CRP—mg/L</th>
<th>CRP†—mg/L</th>
</tr>
</thead>
<tbody>
<tr>
<td>Serum samples—no.</td>
<td>169</td>
<td>4 (1–4)</td>
<td>7 (6.5–9)</td>
<td>135 (80)</td>
<td>19 (11)</td>
<td>5.9±11.1</td>
<td>2.1 (0.8–6.2)</td>
</tr>
<tr>
<td>Serum samples</td>
<td>637</td>
<td>2 (1–2)</td>
<td>17 (11–26)</td>
<td>417 (66)</td>
<td>115 (18)</td>
<td>11.2±22.9</td>
<td>3.0 (0.8–10.0)</td>
</tr>
<tr>
<td>Serum samples IFX ( \geq 3 , \mu\text{g/mL} )—no. (%)</td>
<td>314</td>
<td>11 (2–26)</td>
<td>17 (9–40)</td>
<td>102 (33)</td>
<td>183 (58)</td>
<td>24.4±32.5</td>
<td>11.7 (3.1–29.7)</td>
</tr>
<tr>
<td>Serum samples ATI positive—no. (%)</td>
<td>367</td>
<td>4 (1–13)</td>
<td>24 (17–50)</td>
<td>295 (80)</td>
<td>36 (10)</td>
<td>8.5±13.8</td>
<td>2.4 (0.7–9.7)</td>
</tr>
<tr>
<td>CRP—mg/L</td>
<td>2.1 (0.8–6.2)</td>
<td>1.8 (0.8–6.2)</td>
<td>3.0 (0.8–10.0)</td>
<td>11.7 (3.1–29.7)</td>
<td>2.4 (0.7–9.7)</td>
<td>3.6 (1.0–12.7)</td>
<td></td>
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*Mean±SD.
†Values are median and IQR.
ATI, antibodies to infliximab; AZA, azathioprine; CRP, C-reactive protein; IFX, infliximab; IMM, immunosuppressive; MTX, methotrexate.

Figure 1  XY plot of infliximab (IFX) and antibody to infliximab (ATI) concentration of all 1487 samples. Lower limit of quantitation (LLOQ) for IFX and ATI using the homogenous mobility shift assay was respectively 0.98 \( \mu\text{g/mL} \) and 3.13 U/mL. Seventeen data points were outside the axis limits: 4/17 samples had IFX >35 \( \mu\text{g/mL} \) (none were positive for ATI) and 13/17 samples had ATI >90 U/mL (none were positive for IFX). Positive (+) and negative (−) signs represent respectively samples with detectable and undetectable IFX or ATI.

Figure 2  Receiver operating characteristic curve analysis showing the infliximab threshold that best discriminated disease activity, as measured by C-reactive protein (concentration \( \leq 5 \, \text{mg/mL} \) was defined as inactive disease).
active disease in the multivariable model. This observation suggests that ATI negatively influence the efficacy of IFX even in low concentrations and in the presence of an adequate trough drug concentration. These results point towards an alternative mechanism of action of how ATI impair drug activity as it was previously thought that this was mainly driven by an effect on the pharmacokinetics of the drug causing a faster clearance. Our results support the hypothesis that at least in some patients the effect of ATI is mediated through binding of the ATI to the idio-
type of IFX, thereby blocking the binding to tumour necrosis factor (TNF) and directly neutralising the drug’s activity. Most likely, in the majority of patients, it is a combination of both mechanisms that will ultimately lead to impaired drug activity. In fact, it is believed that when only one or two endogenous antidrug antibodies bind to an exogenous protein, this has no effect on its half-life and may actually increase and approach that of endogenous IgG. On the other hand, when three or more endogenous antidrug antibodies bind to the exogenous protein simultaneously, the resulting immune complex will be eliminated rapidly through phagocytosis, increasing the clearance and decreasing the half-life of the drug.29 Aside from characterising the neutralising capacity of endogenous antidrug antibodies and complex formation in vivo, future research should evaluate the role of other pharmacokinetic parameters such as the AUC10 and Cmax31 as these might prove to be important determinants of the pharmacokinetic/pharmacodynamic relationship that are negatively affected by ATI.

Our study has several clinical implications. Identification of optimum IFX trough and ATI threshold concentrations should facilitate TDM using a drug-tolerant assay. Currently, the most common application of TDM is for the evaluation of patients with a secondary loss of response. In this paradigm, patients with subtherapeutic IFX concentrations who lack ATI may benefit from dose intensification. A second group consists of patients who are ATI negative and have an adequate IFX trough concentration. Although scientific evidence is lacking, inflammation in these individuals may be driven by processes that are not TNF dependent. Accordingly, a switch to an ‘out of class’ agent such as vedolizumab might be optimal.32 In a third group of patients, those with absent drug and a high concentration of ATI, a switch to another TNF antagonist is likely optimal since the patient originally responded to this class of drugs and, for the most part, ATI does not neutralise other agents. Finally, little is known regarding the optimal management of patients with detectable concentrations of both IFX and ATI. Dose escalation may be considered in those patients as well as a switch to another TNF antagonist as from our data it was observed that even low ATI in the presence of IFX are an independent predictor for elevated CRP (>5 mg/L).

Apart from the emerging role in the management of patients with secondary loss of response, other important applications of TDM are likely to evolve including prospective dose optimisation with selective dose intensification or reduction. This strategy could improve therapeutic efficacy, reduce the risk of ATI formation and decrease the cost of care.18 31 34 Studies such as the TAXIT and TAILORIX trials, in which patients are randomly assigned to standard care or to prospective dose optimisation based on TDM, have been initiated. The TAXIT trial showed that, during maintenance therapy with IFX, better clinical outcomes were achieved when a threshold IFX trough concentration of ≥3 μg/mL was targeted and that patients with an IFX trough concentration of >7 μg/mL could be safely dose de-escalated, resulting in significantly lower drug costs without affecting the proportion of patients in remission.35 After this
initial dose optimisation, no additional benefit was observed to continue concentration-based dosing over clinically based dosing throughout the first year. The prospectively acquired data from TAXIT confirm the IFX trough concentration threshold that was found here, and its validity across different assay formats as in the TAXIT trial a clinically validated ELISA was used to measure IFX concentrations.

Our study had several limitations. First, it was retrospectively designed. However, data were derived from four parent cohorts that evaluated patients with CD receiving IFX maintenance therapy and where trough serum samples were prospectively collected. Second, disease activity was measured by CRP concentration, a biomarker for inflammation, rather than by the gold standard of colonoscopy. During the time the parent studies were designed, endoscopic evaluation was not routinely used as an outcome measure in clinical research. Nevertheless, CRP has previously been shown to be an accurate marker to predict reactivation of disease activity in patients with CD treated with IFX therapy. Finally, the IFX and ATI thresholds were not independently confirmed in a separate population of patient samples. This is an important limitation that will be redressed in a follow-up study.

In conclusion, we have defined trough IFX and ATI concentration thresholds that are independently associated with remission in CD. Further investigation into the mechanisms by which ATI negatively affect the efficacy of IFX and strategies to prevent the development of sensitisation are a research priority.

**Contributors** NVC: acquisition of data, analysis and interpretation of data, drafting of the manuscript; RK: drafting of the manuscript; BGL and GRG: critical revision of the manuscript for important intellectual content; LS and GY2: statistical analysis; SS and LO: technical or material support; SL: statistical analysis and interpretation of data; PIR: study concept and design, critical revision of the manuscript for important intellectual content; AG and WJS: interpretation of the data and critical revision of the manuscript for important intellectual content; SV: study concept and design, interpretation of the data and critical revision of the manuscript for important intellectual content and study supervision.

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