





BACKGROUND

Strategies for predicting ustekinumab (UST) trough levels with machine learning techniques can improve personalized care and aid in decision making for UST initiation or scheduling.

AIMS

 The aim of this study was to identify variables capable of predicting an adequate UST response through a gradient boosted decision trees (GBDT) model.

METHODS

- A retrospective cohort of Crohn's disease (CD) patients from our quaternary referral center being treated with UST were reviewed for variables including age, gender, ethnicity, BMI, dosing schedule, time passed since starting UST, previously used biologics, disease duration, age of diagnosis, disease location, disease behavior, and measurements of inflammation.
- As part of feature selection, a univariate analysis was conducted to determine which features significantly correlated with UST trough levels.
- These features were then used to train a multivariate GBDT model, which was then evaluated using a nested crossvalidation framework.
- The gini importance of the features included in each model was then ranked and then averaged across the different models.

Gradient boosted decision tree to model ustekinumab trough levels in **Crohn's Disease**

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RESULTS



TABLE 1. Gini importance ranking of variables from different models

Variable	All	CV + previous biologic exposure + CD characteristics	CV + CD Characteristi cs	CV + previous biologic exposure	CV + Inflammato ry Markers	CV	Aver age
Core Variables (CV)							
Gender	9	4	4	6	5	3	4
Dosing Schedule	5	2	3	3	4	2	2
Time on UST	1	1	1	1	1	1	1
Previous Biologic Exposures							
Failed Adalimumab	6	7	-	4	-	-	6
Failed Infliximab	10	6	-	2	-	-	5
Failed Certolizumab	7	8	-	5	-	-	7
CD Characteristics							
Non-stricturing, non-penetrating	12	10	6	-	-	-	11
Penetrating	11	5	5	-	-	-	10
Ileal Disease	8	9	7	-	-	-	12
Ileocolonic Disease	4	3	2	-	-	-	3
Inflammatory Markers							
CRP	2	-	-	-	2	-	8
ESR	3	-	-	-	3	-	9

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RESULTS

155 CD patients were identified in our hort with UST trough levels obtained. nivariate analysis determined the llowing variables to be significant sitive predictors of adequate UST sponse: female gender, higher equency of dosing, time on UST, and e Montreal classifications B1, B3, L1,

ne following were negative predictors adequate UST response: ESR, CRP, iled adalimumab, failed infliximab, iled certolizumab

esults of various input variable mbinations are outlined in figure 1. ne gini importance of features in each odel is included in table 1.

the generated GBDT models, core ariables only, and core variables with evious biologic exposure and CD naracteristics were the best performing odels with mean AUC of 0.72 ± 0.10 nd 0.71 ± 0.09 respectively.

CONCLUSIONS

nis proof-of-concept study

emonstrates how predictive models in be used to understand variables portant for UST response, and hen additional doses of UST might necessary to achieve therapeutic vels.

ur proof-of-concept models seem to ustrate that the most predictive riables for UST trough levels were ne passed since starting UST, osing schedule, ileocolonic disease, nd previously failed anti-tumor ecrosis factor agents.