

# Clinical Utility of Therapy Selection Informed by Predicted Nonresponse to Tumor Necrosis Factor- $\alpha$ Inhibitors: An Analysis from the Study to Accelerate Information of Molecular Signatures (AIMS) in Rheumatoid Arthritis

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## INTRODUCTION

### Background

- Rheumatoid arthritis (RA) is a systemic autoimmune disease characterized by chronic, symmetrical articular inflammation, pain, and disability
- If left untreated, RA results in progressive joint destruction with subsequent disability and increased mortality
- ACR50 response rates at 24 weeks with TNFi therapy in RA patients with inadequate response to methotrexate range from 27% to 38%
- In the absence of a precision medicine test to inform treatment selection, two-thirds of RA patients fail to achieve ACR50 responses and therefore are unlikely to reach the treat-to-target goal of remission

### PrismRA<sup>®</sup> Test Description

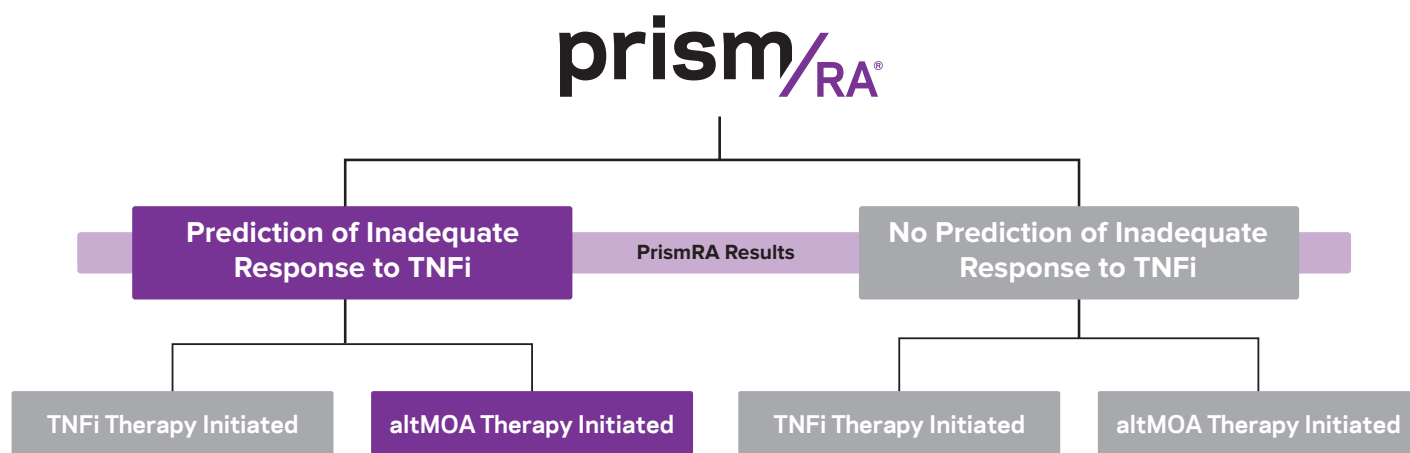
- A blood-based molecular signature response classifier that uses RNA sequencing data combined with clinical factors to predict the likelihood that an individual RA patient will inadequately respond to TNFi therapy

### METHODS

- This study reports an interim analysis from the AIMS study, which is a large scale, prospective study to collect longitudinal patient clinical and molecular data to evaluate clinical utility
- Multicenter study with >70 academic and private rheumatology sites in the US (35 sites were included for this interim analysis)
- Eligibility criteria included age  $\geq 18$  years and a clinical diagnosis of RA, with low to high disease activity
- Data was collected at baseline, treatment initiation, and 12- and 24-week follow-up visits
- The primary endpoint of the outcome analysis was therapeutic responsiveness defined by achievement of ACR50 at 24 weeks
- The secondary endpoint evaluated improvement in baseline CDAI scores exceeding MID\*

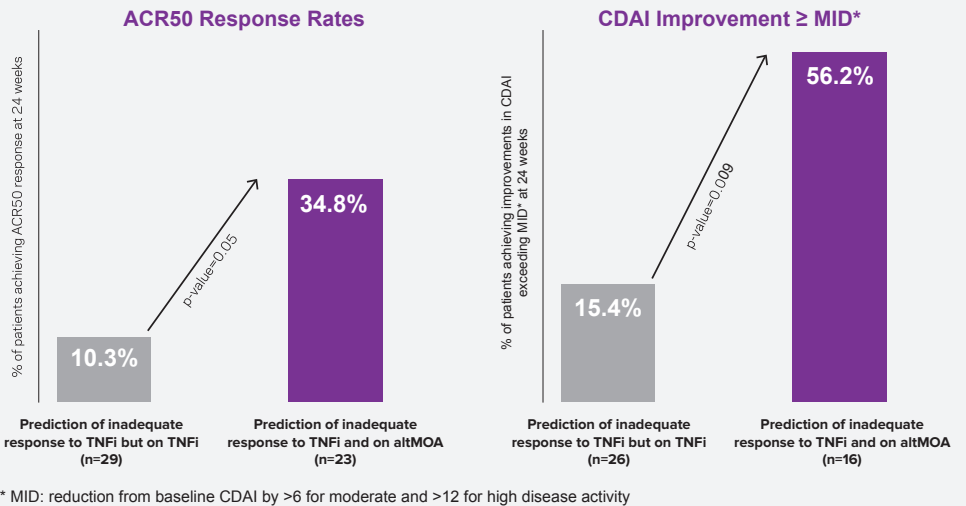
\* Minimally important difference (MID): reduction from baseline CDAI by >6 for moderate and >12 for high disease activity

## CLINICAL OUTCOMES COHORT



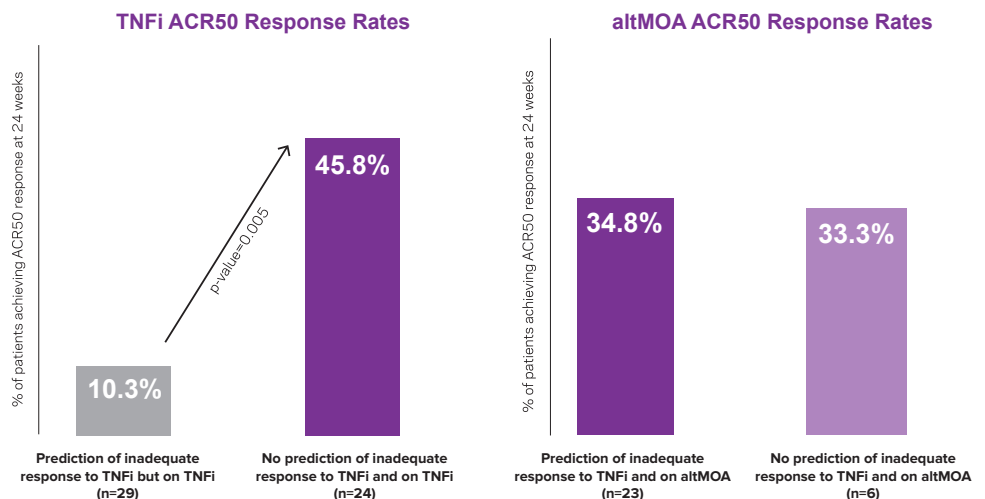
## Therapeutic Responses in Patients with Prediction of Inadequate Response

- Patients who were predicted inadequate responders (molecular signature detected) who were prescribed an altMOA instead of a TNFi therapy, experienced significantly improved therapeutic response



## Therapeutic Responses in Patients with Prediction vs. No Prediction of Inadequate Response

- Patients with a molecular signature of inadequate response (predicted inadequate responders to TNFi) had significantly lower therapeutic responses to TNFi therapies when compared with patients lacking the molecular signature
- No significant differences in therapeutic responses between patients with or without a molecular signature who received an altMOA.



## CONCLUSION

- This study demonstrated the clinical utility of adopting PrismRA into rheumatology practice; it may reduce inappropriate use of TNFi and increase responses across the RA population
- Patients with a molecular signature of inadequate response prescribed an altMOA, instead of a TNFi therapy, experienced significantly improved therapeutic response (34.8% achieved ACR50 on altMOA while 10.3% reached ACR50 on TNFi)
- 90% of patients with a molecular signature of inadequate response according to PrismRA are likely inadequate responders to TNFi therapies and according to a treat-to-target approach should be directed to altMOAs

**With PrismRA, providers can evaluate therapy ineffectiveness before a patient presents clinically, thus PrismRA-informed care shows an improvement in clinical outcomes over current care**