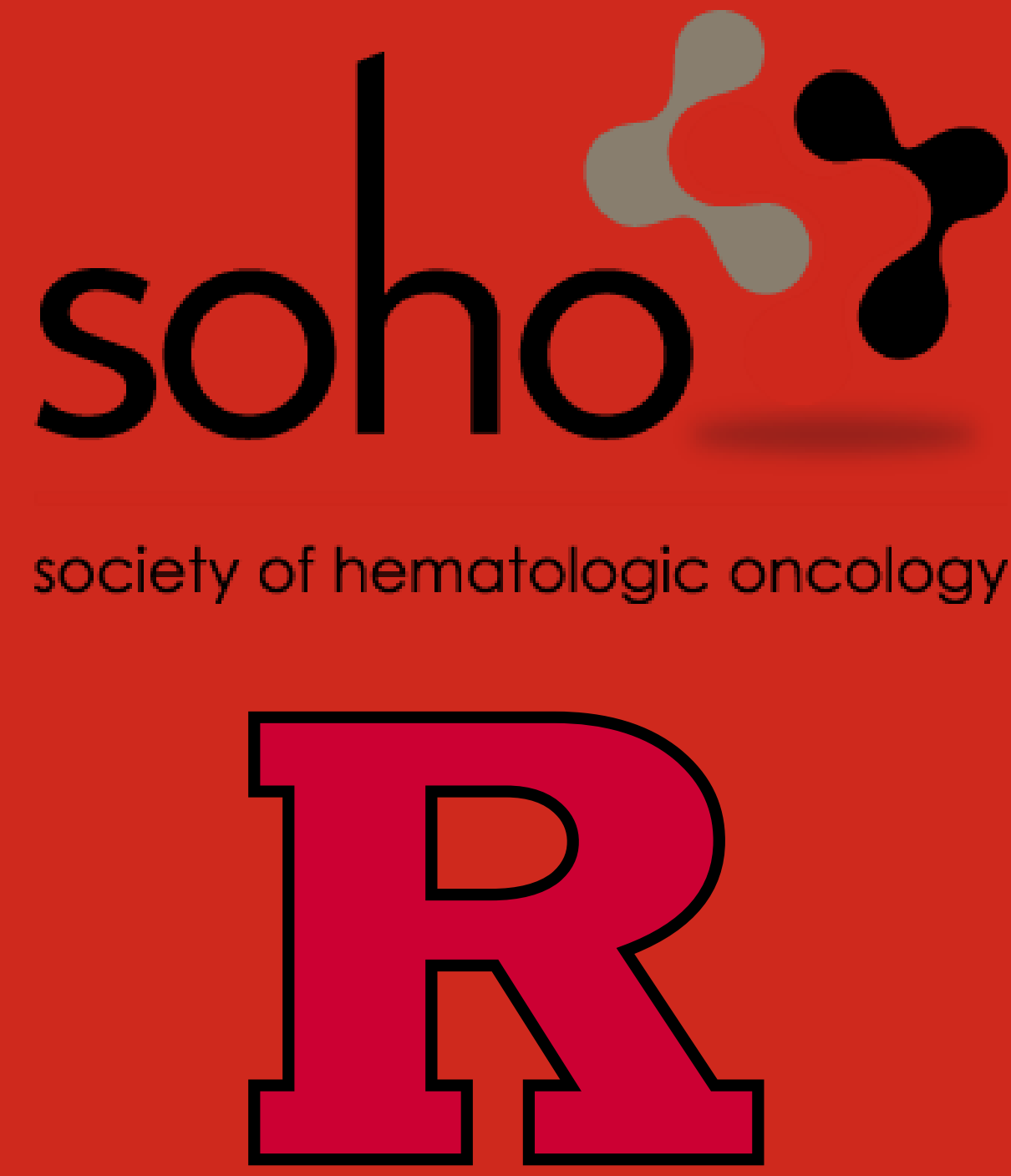


Asciminib Versus Dasatinib/Nilotinib as First-Line Tyrosine Kinase Inhibitor Therapy in Chronic Myeloid Leukemia: A Propensity Score-Matched Analysis of Disease Progression and Safety Outcomes



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SOHO 2026 · Abstract #913 · Chronic Myeloid Leukemia (CML)

INTRODUCTION

- Asciminib targets the ABL myristoyl pocket, bypassing the ATP-binding site used by dasatinib and nilotinib.¹
- The phase 3 ASC4FIRST trial demonstrated superior 48-week major molecular response (MMR) rates for asciminib versus investigator-selected TKIs (**67.7% vs. 49.0%**).^{2,3}
- Real-world comparative data against second-generation TKIs remain limited.

AIM

To compare disease progression and safety outcomes between asciminib and dasatinib/nilotinib as first-line TKI therapy in chronic myeloid leukemia (CML) using real-world data.

METHODS

- Data source.** TriNetX Research Network — 112 healthcare organizations, approximately 147.5 million patients.⁴
- Population.** Adults with BCR-ABL–positive CML initiating asciminib (n=327) or dasatinib/nilotinib (n=5,789) as their only TKI within 90 days of diagnosis.
- Matching.** Propensity score matching (1:1) on age, sex, race, ethnicity, diabetes, chronic kidney disease, heart failure and hepatic fibrosis/cirrhosis yielded 315 patients per cohort.
- Primary endpoint.** Disease progression between days 90 and 270, defined as escalation to ponatinib or omacetaxine, hematopoietic stem cell transplantation, or transformation to acute leukemia.
- Secondary safety outcomes.** Cytopenia, pleural effusion, cardiovascular events and pancreatitis.
- Outcomes were extracted from structured EHR data without manual chart review.

CONCLUSIONS

- The cytopenia reduction with asciminib was substantial and statistically significant.**
- The progression difference did not reach statistical significance, partly reflecting insufficient power to detect a moderate effect.
- Retrospective claims-level ascertainment of progression is imperfect, as some therapy switches may not represent true disease advancement.
- The safety profile represents the most defensible argument for asciminib in the frontline setting** — a consideration of particular relevance when managing a largely chronic disease population.

RESULTS

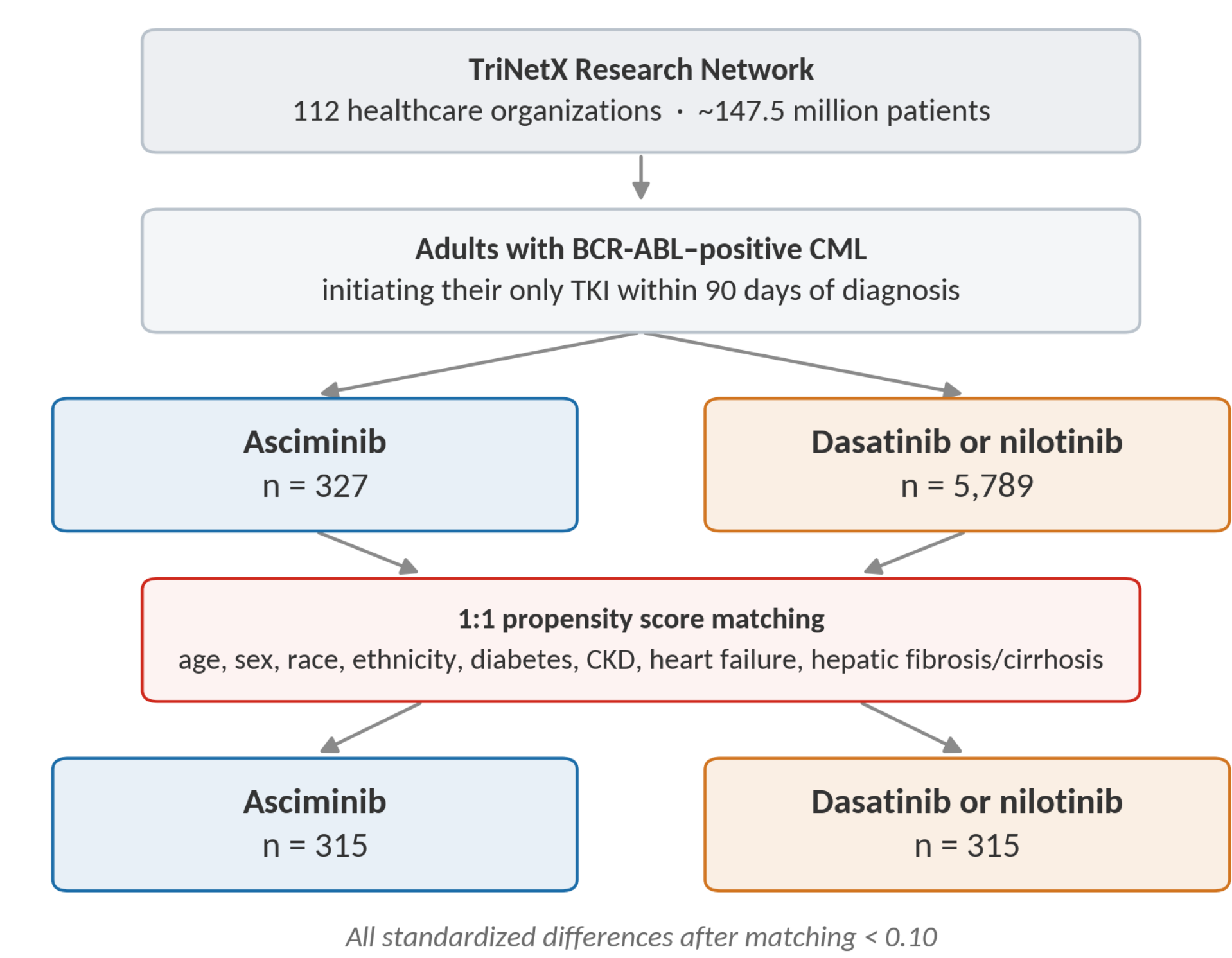


Figure 1. Cohort selection and propensity score matching.

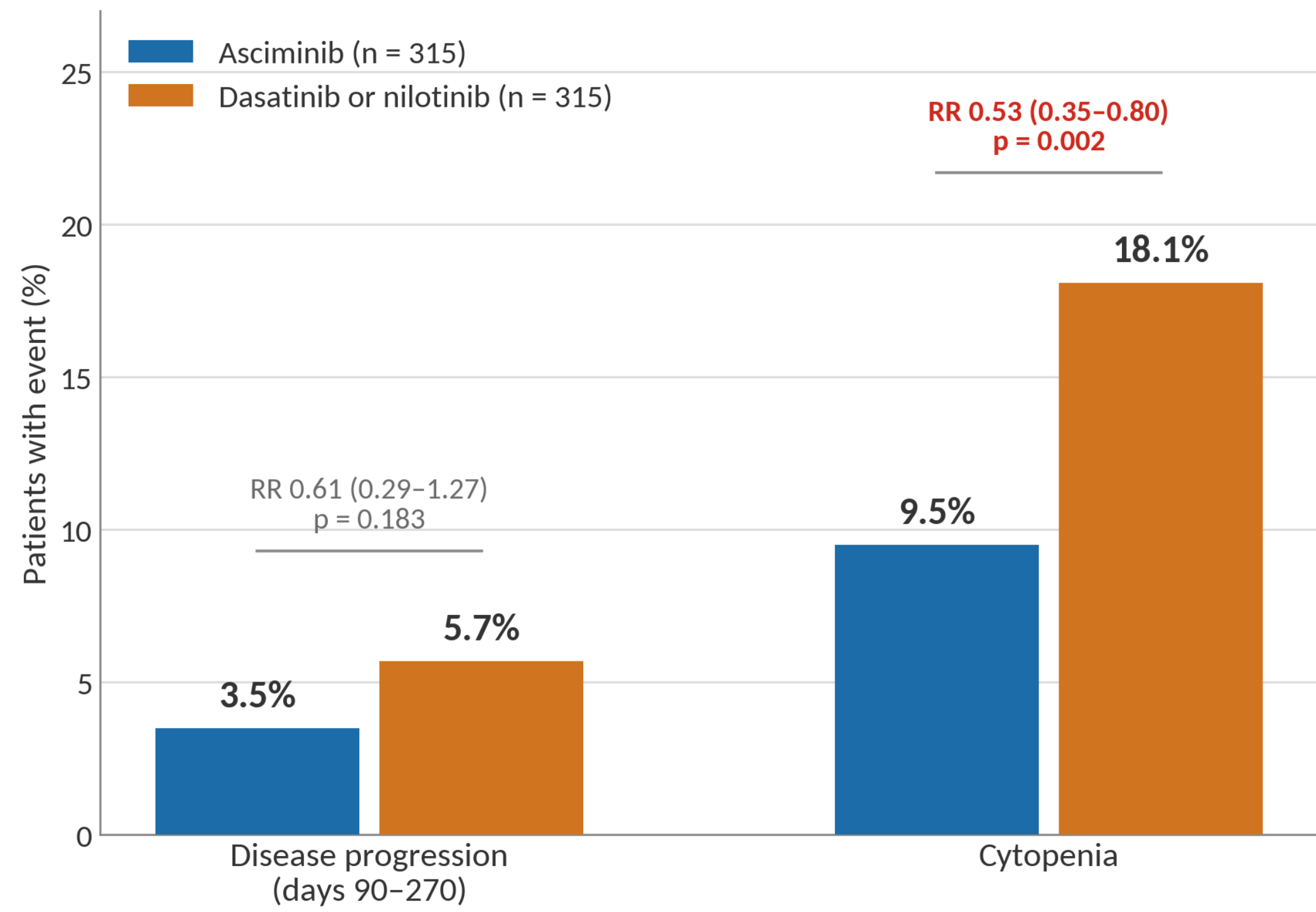
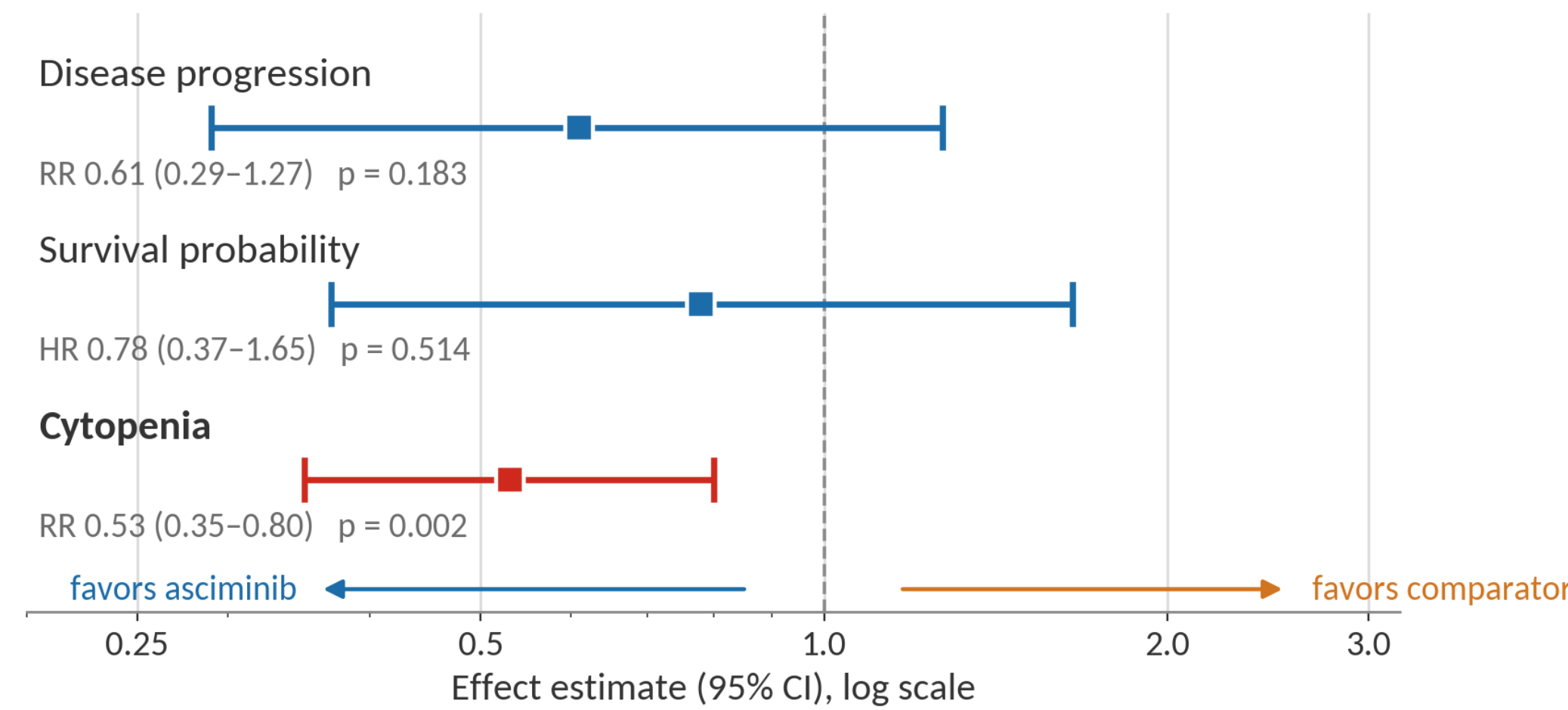


Figure 2. Event rates by cohort. Cytopenia differed significantly; progression did not.

KEY RESULTS

- Post-matching balance was acceptable across all covariates (standardized differences < 0.10).
- Eleven asciminib patients (3.5%) progressed versus 18 (5.7%) in the comparator cohort — RR 0.61 (95% CI 0.29–1.27), p = 0.183.
- Survival probabilities were 94.5% versus 93.4% — HR 0.78 (95% CI 0.37–1.65), log-rank p = 0.514.
- Cytopenia occurred in 9.5% versus 18.1% — RR 0.53 (95% CI 0.35–0.80), p = 0.002.**
- Pleural effusion and cardiovascular events were rare in both groups, with cell counts too low for meaningful subgroup analysis. No pancreatitis events were recorded in either cohort.

Figure 3. Effect estimates with 95% confidence intervals.



LIMITATIONS

- Retrospective claims-level ascertainment of progression is imperfect, as some therapy switches may not represent true disease advancement.
- The comparison had insufficient power to detect a moderate effect on disease progression.
- Pleural effusion and cardiovascular events were rare in both groups, with cell counts too low for meaningful subgroup analysis.
- Outcomes were extracted from structured EHR data without manual chart review.

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Table 1. Progression and safety outcomes in the matched cohorts.

Outcome	Asciminib (n = 315)	Dasatinib or nilotinib (n = 315)	Effect estimate (95% CI)	p
Disease progression, days 90–270	11 (3.5%)	18 (5.7%)	RR 0.61 (0.29–1.27)	0.183
Survival probability (Kaplan-Meier)	94.5%	93.4%	HR 0.78 (0.37–1.65)	0.514
Cytopenia	9.5%	18.1%	RR 0.53 (0.35–0.80)	0.002
Pleural effusion	rare*	rare*	—	—
Cardiovascular events	rare*	rare*	—	—
Pancreatitis	0	0	—	—

* Rare in both groups; cell counts too low for meaningful subgroup analysis.