

# Real-World Cost-Effectiveness of Publicly Reimbursed Multi-Gene Panel Sequencing to Inform Therapeutic Decisions for Advanced Non-Small Cell Lung Cancer in British Columbia, Canada

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## Background

- Multi-gene panel sequencing streamlines treatment selection for advanced non-small cell lung cancer (NSCLC). Implementation continues to be uneven across jurisdictions, in part due to uncertain clinical and economic impacts.
- This study determined the population-level cost-effectiveness of publicly reimbursed multi-gene panel sequencing compared to single-gene EGFR testing for advanced NSCLC.

## Methods

- In British Columbia (BC), Canada, the public healthcare system reimbursed a multi-gene panel in September 2016.
- Our population-based retrospective study design used comprehensive patient-level cancer control and linked administrative health databases. We considered adult BC residents with an advanced lung cancer diagnosis between September 2016 and December 2018.
- Using a machine learning approach, we conducted 1:1 genetic algorithm matching of recipients receiving multi-gene panel sequencing to controls receiving single-gene testing, maximizing balance on observed demographic and clinical characteristics.
- Following matching, we estimated mean three-year survival time and costs (public healthcare payer perspective; 2021 CAD) and calculated the incremental net monetary benefit (INMB) for life-years gained (LYG) at conventional willingness-to-pay thresholds using inverse probability of censoring weighted linear regression and nonparametric bootstrapping.

## Results

- We matched 858 panel-eligible advanced NSCLC patients to controls, achieving balance for the 16 included covariates.
- Average test turnaround times were 18.6 days for multi-gene panel sequencing and 7.0 days for single-gene testing.
- After matching, mean  $\Delta$  costs were \$3,529 (95%CI: -\$4,268, \$10,942) and mean  $\Delta$  LYG were 0.08 (95%CI: - 0.04, 0.18).
- The INMB was \$523 (95%CI: -\$6,256, \$7,023) at \$50,000/LYG, with a 57.5% probability of being cost-effective, and \$4,575 (95%CI: -\$5,468, \$14,064) at \$100,000/LYG, with an 84.0% probability of being cost-effective.



Using population-level real-world data, we found multi-gene panels in advanced NSCLC were cost-effective at higher thresholds, even with differences in survival and costs that were not statistically significant compared to single-gene testing

Figure 1. Balance of Characteristics between Patients Receiving Single-gene or Multi-gene Panel Testing, Before and After Matching.

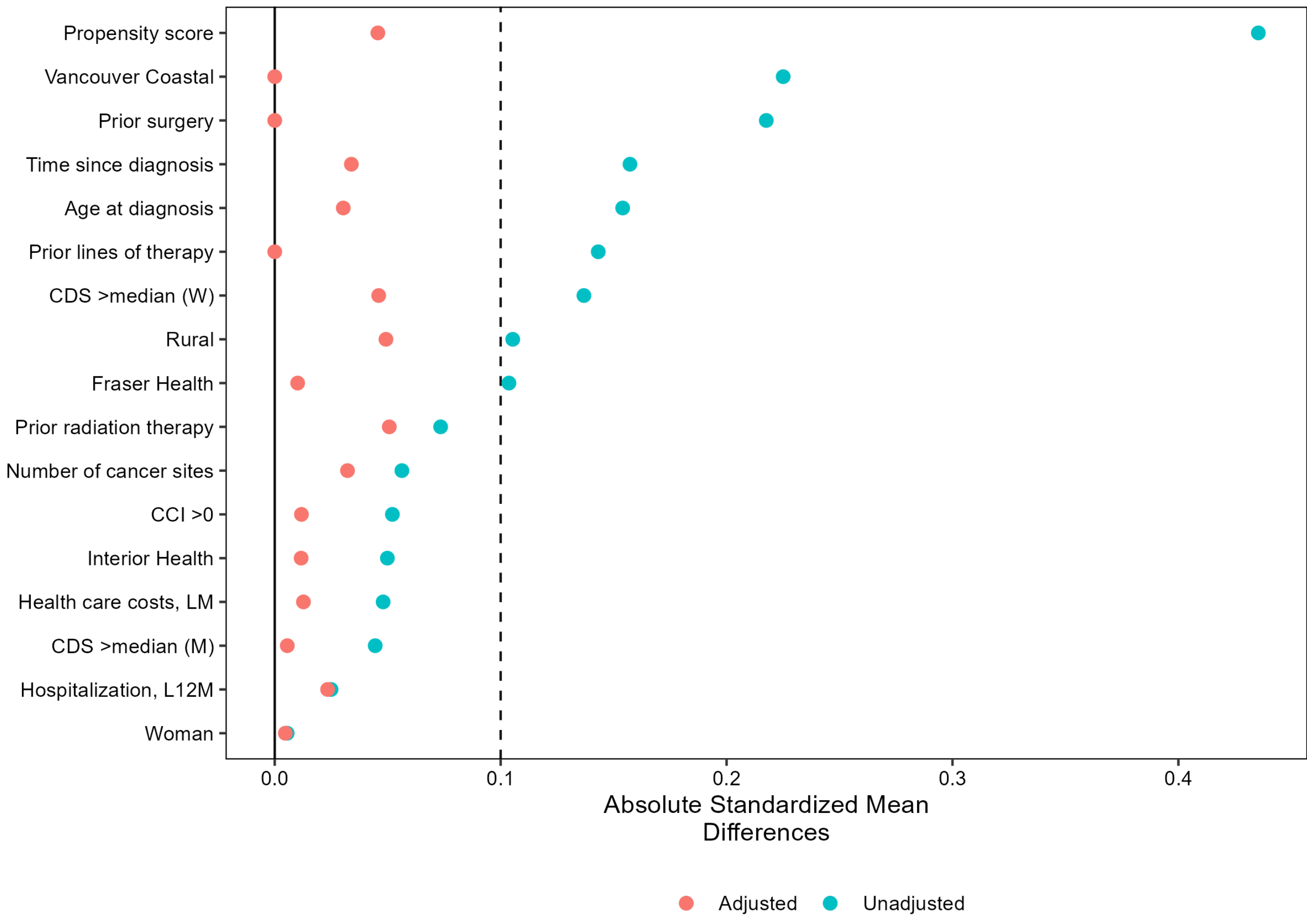
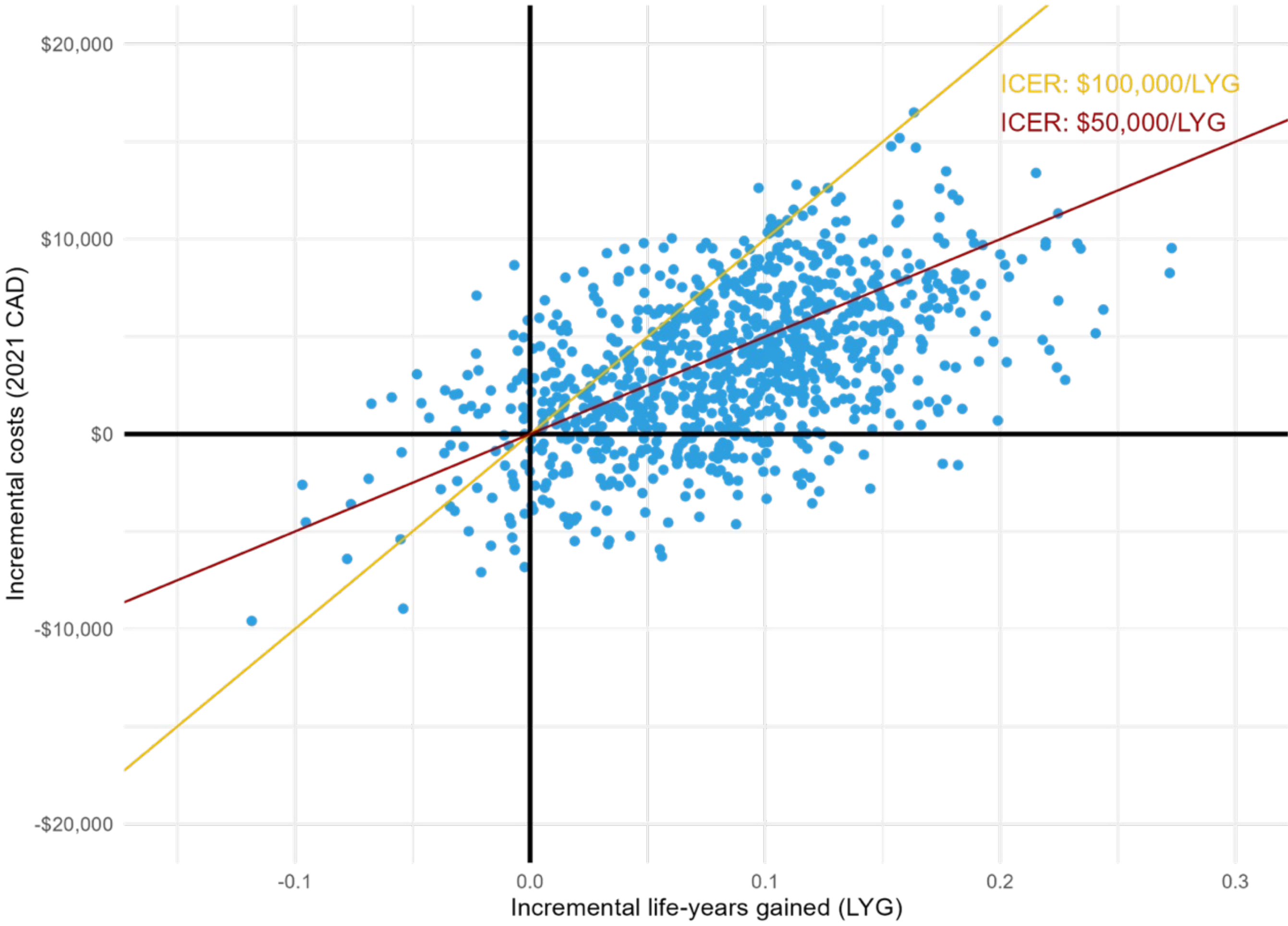


Figure 2. Cost-effectiveness plane for multi-gene panel testing compared to single-gene EGFR testing.



## Conclusion

- We used machine learning-based quasi-experimental methods to identify a well-balanced counterfactual.
- We found a high probability that panel-based testing in advanced NSCLC would be cost-effective at higher thresholds, even with differences in survival and costs that were not statistically significant compared to single-gene EGFR testing.

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Access to data provided by the Data Steward(s) is subject to approval, but can be requested for research projects through the Data Steward(s) or their designated service providers. All inferences, opinions, and conclusions drawn in this publication are those of the authors, and do not reflect the opinions or policies of the Data Steward(s).

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