Designing Studies to Evaluate Biomarkers for Selecting Patient Treatment

Holly Janes^{*}, Fred Hutchinson Cancer Research Center Margaret Pepe, Fred Hutchinson Cancer Research Center Ying Huang, Fred Hutchinson Cancer Research Center

Abstract

Biomarkers associated with patient response to treatment have the potential to improve clinical outcomes by restricting treatments to the patients most likely to benefit. The ideal setting for evaluating a treatment selection biomarker is a randomized controlled trial. The biomarker may be measured at baseline on the entire trial population, or on a subset of participants potentially selected on the basis of treatment response. Existing study design methodology is limited, and focuses on evaluating biomarkers by testing for a statistical interaction between biomarker and treatment assignment. We propose an approach which powers the biomarker study to evaluate the impact of a marker-based treatment policy on the population response rate. We provide methods for determining the required number of patients and optimal treatment allocation and for sub-sampling from the trial population based on treatment response. We compare these designs with those powered to test for statistical interactions and illustrate the approach using a study to evaluate the Oncotype DX marker for selecting adjuvant chemotherapy to treat estrogen-receptor positive breast cancer.

Keywords: Biomarker; Treatment selection; Clinical trial; Interaction.

Presenting author