

Development of prediction tools for diarrhea and rash in breast cancer patients receiving lapatinib in combination with capecitabine: rule of thumb in prediction studies

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To the Editor,

I was interested to read the paper by Dranitsaris G and colleagues published in *Breast Cancer Res Treat* 2014 Oct. The purpose of the authors was to predict the risk of \geq grade 2 diarrhea and rash prior to each cycle of Lapatinib and Capecitabine (L-CAP) in HER-2-positive patients with metastatic breast cancer (MBC) [1].

They used generalized estimating equations to develop the risk models using a backward elimination process. A receiver operating characteristic curve (ROC) analysis was undertaken to measure the predictive accuracy of the scoring algorithms [1].

For prediction studies, we need two different cohort datasets or at least one cohort dataset splitting that to develop our prediction model and then to validate it. Without validation of prediction models, most of the times, misleading results may be the main outcome of such researches [2–5].

They concluded that their models provide patient-specific risk information that could be helpful in assessing the risks and benefits of L-CAP in the MBC patients. Such a

conclusion may have limited generalizability due to no validation of prediction model which is a rule of thumb in prediction studies.

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