

Imbalanced Data From U.S. Cotton Variety Test

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Harvested acres of cotton in the U.S. in 2017. Image by the USDA Economic Research Service.

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Since the 1960s, the USDA National Cotton Variety Test (NCVT) has tested more than 1,300 varieties and breeding lines over six growing regions in the U.S. Cotton Belt. Data from this rich multilenvironment trial (MET) are used to compare entries and evaluate the genetic and agronomic development, referred to as the longsterm trend, in cotton production over the decades.

In the literature, linear mixed models (LMM) have been widely used to analyze the short the literature, linear mixed models (LMM) have been widely used to analyze the short the literature, linear mixed models (LMM) have been widely used to analyze the short the literature, linear mixed models (LMM) have been widely used to analyze the short the literature, linear mixed models (LMM) have been widely used to analyze the short the mostly below and literature, linear mixed models (LMM) have been widely used to analyze the short the short data, mostly balanced (i.e., all varieties tested at all environments). However, for a longiterm project such as NCVT, many old variety entries drop out each year and are replaced by new entries, leading to highly missing variety by the literature combinations. The resulting extreme imbalance challenges the validity of the results from fitting an

LMM.

A simulation study investigated how the imbalance caused by various dropouts affects the estimation of model parameters and prediction of the variety's overall and local performances. Estimation of longsterm trends was found to be most affected by the imbalance among all parameters. Including the longsterm trend in the LMM is crucial for cotton performance prediction and should be implemented in other longsterm crop variety test trials.

Adapted from

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