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# Discussion of Sesia et al. 2018 and the Knockoff Framework

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#### 1. On the problem

The authors of Sesia et al. (2018) set out to design a procedure for variable selection with provable statistical guarantees. The *knockoff* algorithm proposed by Sesia et al. (2018), provably controls the FDR of conditionally independent variables. Denoting with x and y the predictor and outcome variables, respectively. The *false detection proportion*, a.k.a. the *false selection proportion*, is defined as V/R where R is the number of variables selected, and V is the number of such variables where  $y|x_{-j}$  is independent of  $x_j$ . The knockoff algorithm of Sesia et al. (2018) provably control the  $FDR := \mathbb{E}[FDP]$ , at some user selected magnitude.

Crucially for our comments: (1) The FDR is an expectation with respect to variability in x and y, i.e., a  $random\ design\ guarantee$ . (2) The procedure is  $model\ free$ , or non-parametric in that nothing is assumed on the parametric form of y|x. (3) The proofs assume full knowledge of  $F_x$ , i.e., the joint distribution of predictors, marginalized over y. (4) The method aims at good variable selection, not prediction.

## 2. On the method

The fundamental idea of the method is to generate variables that have all the properties of the original  $x_j$ , only that they are conditionally uncorrelated to y. These are termed knockoff variables. The method then proceeds to compute a test statistic that captures the difference in the strength of the dependence of the knockoff, and the original variable. This statistic is then compared to it resampling distribution: the distribution over resampled knockoffs.

- 2.1. Other Variable Selection Methods
  - 2.2. Other Knockoffs
- 2.3. Permutation Testing and Symmetries
  - 3. FUTURE RESEARCH
  - 4. On the hypotheses
  - 5. On the problem setup

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## REFERENCES

SESIA, M., SABATTI, C. & CANDES, E. J. (2018). Gene hunting with hidden markov model knockoffs. *Biometrika* 00000.

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