

Combinatorial Hypothesis Testing

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1 Introduction



Suppose we observe an n -dimensional vector $\mathbf{X} = (X_1, \dots, X_n)$. The null hypothesis H_0 is that the components of \mathbf{X} are independent and identically distributed (i.i.d.) standard normal random variables. We denote the probability measure and expectation under H_0 by \mathbb{P}_0 and \mathbb{E}_0 , respectively.

Combinatorics kicks in as we consider the alternative hypotheses, by introducing a class \mathcal{C} with some combinatorial structure: consider a class $\mathcal{C} = \{S_1, \dots, S_N\}$ of N sets of indices such that $S_k \subset \{1, \dots, n\}$ for all $k = 1, \dots, N$. Under H_1 , there exists an $S \in \mathcal{C}$ such that X_i has a distribution determined by whether i is in S :

1. In its simplest form, as discussed in [1, 3, 4], we consider

$$X_i \text{ has distribution } \begin{cases} \mathcal{N}(0, 1), & \text{if } i \notin S \\ \mathcal{N}(\mu, 1), & \text{if } i \in S \end{cases}$$

where $\mu > 0$ is a positive parameter and components of \mathbf{X} are independent.

2. In testing correlations [2], we consider

$$\text{Cov}(X_i, X_j) = \begin{cases} 1, & \text{if } i = j \\ \rho, & \text{if } i \neq j \text{ with } i, j \in S \\ 0, & \text{otherwise} \end{cases}$$

For each $S \in \mathcal{C}$, we denote the probability measure and expectation by \mathbb{P}_S and \mathbb{E}_S , respectively. Many interesting examples of \mathcal{C} arises for this scenario: subsets of size K , cliques, perfect matchings, spanning trees, and clusters.

A *test* is a binary-valued function $f : \mathbb{R}^n \rightarrow \{0, 1\}$. If $f(X) = 0$, then the test accepts the null hypothesis H_0 ; otherwise H_0 is rejected by f . We measure the performance of a test based on the *minimax risk*:

$$R_*^{\max} := \inf_f R^{\max}(f).$$

where $R^{\max}(f)$ is the worst-case risk over the class of interest \mathcal{C} , formally defined by

$$R^{\max}(f) = \mathbb{P}_0\{f(X) = 1\} + \max_{S \in \mathcal{C}} \mathbb{P}_S\{f(X) = 0\}.$$

In this report, we discuss the techniques introduced in [1–3] to derive the asymptotic upper and lower bounds of R_*^{\max} , as well as more recent extensions.

2 Lower Bounds



2.1 Moment Methods



A standard way of obtaining lower bounds for the minimax risk is by putting a prior on the class \mathcal{C} and obtaining a lower bound on the corresponding *Bayesian risk*, which never exceeds the worst-case risk. Because this is true for any prior, the idea is to find one that is hardest (often called *least favorable*). Consider the uniform prior on \mathcal{C} , giving rise to the following *average risk*:

$$R(f) = \mathbb{P}_0\{f(X) = 1\} + \mathbb{P}_1\{f(X) = 0\},$$

where

$$\mathbb{P}_1\{f(X) = 0\} := \frac{1}{N} \sum_{S \in \mathcal{C}} \mathbb{P}_S\{f(X) = 0\},$$

and $N := |\mathcal{C}|$ is the cardinality of \mathcal{C} . The advantage of considering the average risk over the worst-case risk is that we know an optimal test for the former, which, by the Neyman–Pearson fundamental lemma, is the likelihood ratio test, denoted f^* . Introducing $L(X)$, the likelihood ratio between H_0 and H_1 , the optimal test becomes

$$f^*(x) = 0 \quad \text{if and only if} \quad L(x) \leq 1.$$

The (average) risk $R^* = R(f^*)$ of the optimal test is called the *Bayes risk* and it satisfies

$$R^* = 1 - \frac{1}{2} \mathbb{E}_0 |L(X) - 1|$$

3 Clusters



4 Extension



4 References



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