Lower bounds for testing function isomorphism

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Abstract—We prove new lower bounds in the area of property testing of boolean functions. Specifically, we study the problem of testing whether a boolean function f is isomorphic to a fixed function g (i.e., is equal to g up to permutation of the input variables). The analogous problem for testing graphs was solved by Fischer in 2005. The setting of boolean functions, however, appears to be more difficult, and no progress has been made since the initial study of the problem by Fischer et al. in 2004.

Our first result shows that any non-adaptive algorithm for testing isomorphism to a function that "strongly" depends on k variables requires $\log k - O(1)$ queries (assuming k/n is bounded away from 1). This lower bound affirms and strengthens a conjecture appearing in the 2004 work of Fischer et al. Its proof relies on total variation bounds between hypergeometric distributions which may be of independent interest.

Our second result concerns the simplest interesting case not covered by our first result: non-adaptively testing isomorphism to the Majority function on k variables. Here we show that $\Omega(k^{1/12})$ queries are necessary (again assuming k/n is bounded away from 1). The proof of this result relies on recently developed multidimensional invariance principle tools.

Keywords-Boolean functions, property testing, lower bounds

I. INTRODUCTION

This paper is concerned with the field of property testing for boolean functions. Let us recall the standard framework, as originally introduced by Rubinfeld and Sudan [17].

Definition 1.1. Let \mathcal{P} be a class of functions $\{0,1\}^n \to \{0,1\}$. We say that a randomized query algorithm \mathcal{T} with black-box access to an unknown function $f:\{0,1\}^n \to \{0,1\}$ is an (ϵ,q) -tester for \mathcal{P} if it makes at most q queries to $f:\{0,1\}^n \to \{0,1\}$ and then:

- Accepts with probability at least 2/3 when f is in \mathcal{P} ; and,
- Rejects with probability at least 2/3 when f is ϵ -far from every function $f' \in \mathcal{P}$.

Here we say that f and f' are ϵ -far if they differ on at least an ϵ fraction of the inputs in $\{0,1\}^n$, and are ϵ -close otherwise. When the algorithm chooses all of its queries in advance it is *non-adaptive*; otherwise we say it is *adaptive*.

Definition 1.2. For a fixed $\epsilon > 0$ and choice of adaptivity, the *query complexity* of \mathcal{P} is the minimum value of q for which there is an (ϵ, q) -tester. Following standard conventions, when the query complexity q is independent of n for every $\epsilon > 0$, we say that \mathcal{P} is *easy to test*; otherwise we say that it is *hard to test*. This notion is independent of the choice of adaptivity, since non-adaptive query complexity can be (exponentially) bounded in terms of adaptive query complexity.

The last five years have seen great strides in understanding the testability of graph properties. This is the special case in which $f:\{0,1\}^{\binom{V}{2}} \to \{0,1\}$ encodes the adjacency matrix of a graph on vertex set V, and $\mathcal P$ is a property that is closed under graph symmetries; i.e., permutations of V. Indeed, the works [2], [3], [1], [4] have to a large extent characterized the testability of graph properties.

However the characterization problem for general boolean functions is very far from understood and remains a long-standing open problem. In this paper we revisit a major subproblem, introduced in the early work of Fischer, Kindler, Ron, Safra, and Samorodnitsky [7]: the difficulty of testing isomorphism to a function given in advance.

A. Testing g-isomorphism

Two boolean functions $f:\{0,1\}^n \to \{0,1\}$ and $g:\{0,1\}^n \to \{0,1\}$ are said to be *isomorphic* to each other if they are identical up to reordering input variables. More precisely, we say that f and g are isomorphic to each other if there is a permutation σ on [n] such that for every $x=(x_1,x_2,\ldots,x_n)\in\{0,1\}^n$, $f(x_1,x_2,\ldots,x_n)=g(x_{\sigma(1)},x_{\sigma(2)},\ldots,x_{\sigma(n)})$.

For each function $g:\{0,1\}^n \to \{0,1\}$, we let \mathcal{P}_g denote the class of all functions isomorphic to g. This gives a natural

¹Formally speaking, this makes sense only for a family of properties $(\mathcal{P}_n)_n$, one for each input length.

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testing problem: testing whether an unknown function f is isomorphic to the known function g. This is called the problem of *testing g-isomorphism*. Fischer et al. [7] proposed the following question, a significant component of the research program to characterize testability of boolean functions:

Research goal: Classify all boolean functions g according to whether testing g-isomorphism is easy or hard.

We remark that Fischer [6] solved the analogous problem for *graph properties* soon after; see also [8]. But he called the general case of boolean functions "rather hard", and indeed the authors are not aware of any additional progress specifically on this problem.

In this paper we make progress towards a characterization by showing hardness of testing g-isomorphism for a large class of functions g.

B. Prior work

When the problem of testing function isomorphism was first raised by Fischer et al. [7], a few simple cases were already well understood. First, it is easy to see that for any totally symmetric function $g:\{0,1\}^n \to \{0,1\}$, testing g-isomorphism is easy – no other functions are isomorphic to g, and testing function *identity* requires only $O(1/\epsilon)$ queries.

Another instance of the g-isomorphism testing problem that is well understood is one where $g(x) = x_i$ for some $i \in \{1, \ldots, n\}$. Then the g-isomorphism problem is equivalent to the well-studied dictatorship testing problem at the heart of PCP constructions (first studied in [5]). The query complexity of the dictatorship testing problem is $O(1/\epsilon)$, so this special case of the function isomorphism problem is also easy. Parnas, Ron, and Samorodnitsky [15] also showed that testing g-isomorphism is easy when g is an AND function on any number of variables.

The paper of Fischer et al. [7] introduced a strong new upper bound: they showed that for any k, if g is a k-junta – meaning that g depends on at most k variables – then it is possible to ϵ -test g-isomorphism (non-adaptively, even) with $\operatorname{poly}(k/\epsilon)$ queries. Therefore, testing isomorphism to any O(1)-junta is easy.

Regarding hardness results, prior to this work the only known lower bound for testing g-isomorphism was for the case of g being the Parity_k function, where $\operatorname{Parity}_k: \{0,1\}^n \to \{0,1\}$ is defined by $\operatorname{Parity}_k(x) = x_1 \oplus x_2 \oplus \cdots \oplus x_k$. Fischer et al. [7] showed that when $k \leq o(\sqrt{n})$, ϵ -testing Parity_k -isomorphism non-adaptively requires $\tilde{\Omega}(\sqrt{k}/\epsilon)$ queries. This result implies that testing Parity_k -isomorphism is hard for $\omega(1) \leq k \leq o(\sqrt{n})$.

C. Our results

It seems clear that more work needs to be done on the hardness side of testing g-isomorphism. A first direction

would be to investigate the following conjecture, stated in [7]: "If n is sufficiently large compared to k, and g is a k-junta which is ϵ -far from all (k-1)-juntas, then ϵ -testing q-isomorphism requires $\omega_k(1)$ queries."

Our first main result affirms and significantly strengthens this conjecture:

Theorem 1.3. Let $g: \{0,1\}^n \to \{0,1\}$ be a k-junta which is ϵ -far from being a (k-e)-junta for some $e \ge 1$. Then any non-adaptive ϵ -tester for g-isomorphism must make at least $\log_2(k'/e^2) - O(1)$ queries, where $k' = \min(k, n - k)$.

Qualitatively, Theorem 1.3 implies that testing g-isomorphism for k-juntas is hard whenever $\omega(1) \leq k \leq n - \omega(1)$, provided the k-junta g is far from all juntas on $k - o(\sqrt{k})$ variables. We discuss the possibility of improving this theorem in Section V.

As one very special case, Theorem 1.3 extends the Fischer et al. hardness result for testing Parity_k -isomorphism to all $\omega(1) \leq k \leq n - \omega(1)$ (albeit with a worse query complexity lower bound). It is important to note that the restriction $k \leq n - \omega(1)$ is not an artifact of our proof, but rather is inherent. This is because, e.g., testing Parity_k -isomorphism when k = n - O(1) is easy: if the tester XORs each query response with the parity of all bits in the query, it reduces testing Parity_k -isomorphism to testing $\operatorname{Parity}_{n-k}$ -isomorphism. Thus testing $\operatorname{Parity}_{n-O(1)}$ -isomorphism is easy, by the junta isomorphism-testing result of Fischer et al.

Our Theorem 1.3 is only useful for k-juntas g that are far from being $(k-o(\sqrt{k}))$ -juntas. Perhaps the most natural case not covered by this theorem is that of the majority function on k variables, Maj_k . A straightforward calculation shows that for all $\delta>0$, the function Maj_k is $o_\delta(1)$ -close to being a junta on $k-\delta k$ variables: namely $\mathrm{Maj}_{k-\delta k}$. Nevertheless, the second main result in our paper proves a strong hardness result for testing Maj_k -isomorphism:

Theorem 1.4. For every constant $\delta > 0$, there exists a constant $\epsilon > 0$ such that the following holds: Assuming $1/\epsilon \le k \le (1-\delta)n$, any non-adaptive algorithm for ϵ -testing Maj_k -isomorphism must make at least $\Omega((\delta k)^{1/12})$ queries.

Qualitatively, Theorem 1.4 implies that testing Maj_k -isomorphism is hard for every $\omega(1) \leq k \leq n - \Omega(n)$. Again, this range is optimal: the upper bound cannot be improved because as mentioned, $\mathrm{Maj}_{n-o(n)}$ is o(1)-close to Maj_n ; thus we can test $\mathrm{Maj}_{n-o(n)}$ -isomorphism using the easy Maj_n -isomorphism tester that follows from Maj_n being totally symmetric.

In this extended abstract, we prove Theorem 1.4 only in the case where $\delta=1/4$ (and assuming k is divisible by 3). The few tedious technical modifications needed to handle smaller values of δ are deferred to the full version of the

article. I.e., we prove:

Theorem 1.5. There is a universal $\epsilon_0 > 0$ such that whenever $k \leq (3/4)n$ (and is divisible by 3), any non-adaptive algorithm for ϵ_0 -testing Maj_k -isomorphism must make at least $\Omega(k^{1/12})$ queries.

We remark that the bound in Theorem 1.5 is exponentially stronger than the one in Theorem 1.3: it shows that any non-adaptive algorithm for testing Maj_k -isomorphism must make a number of queries *polynomial* in k. This bound is optimal (up to the right value of the exponent), since Fischer et al.'s [7] upper bound on the query complexity of testing isomorphism to k-juntas implies that testing isomorphism to the Maj_k function can be done with $\operatorname{poly}(k/\epsilon)$ queries.

The result of Theorem 1.4 is also interesting in light of the recent results of Matulef et al. on testing halfspaces: they showed that testing the class of halfspaces is easy [11], but testing a natural subclass of halfspaces – the class of ± 1 -weight halfspaces – is hard [12]. More precisely, they showed a non-adaptive query lower bound of $\Omega(\log n)$ for this class. Our Theorem 1.4 gives an exponentially improved lower bound for a similar subclass of halfspaces: $\Omega(n^{1/12})$ queries for the class of majority functions on, say, n/2 variables.

In light of Fischer [6]'s solution to the isomorphism testing problem for graph properties – i.e., boolean functions with a certain high degree of symmetry – we believe that characterizing testability of g-isomorphism for symmetric k-juntas is an approachable first step. Theorem 1.5 represents progress in this direction. We remark that our method of proving Theorem 1.5 can be extended to handle certain other symmetric k-juntas. However the general case of symmetric k-juntas g has some unexpected tricky aspects to it, which we discuss in Section V.

D. Our techniques

The proofs of Theorems 1.3 and 1.5 both use the standard approach for proving lower bounds in property testing: Yao's Minimax Principle [18]. That is, we prove both theorems by introducing distributions $\mathcal{F}_{\rm yes}$ and $\mathcal{F}_{\rm no}$ on functions that should be accepted and rejected, respectively, by algorithms testing g-isomorphism, and show a lower bound on the number of queries required by any *deterministic* testing algorithm. The main technical contribution of this research is in the design and the analysis of the distributions $\mathcal{F}_{\rm yes}$ and $\mathcal{F}_{\rm no}$.

The main challenge in proving Theorem 1.3 is that the lower bound applies to a very general class of functions g. To prove the theorem we need to design distributions that work without using any structural properties of the function g being tested. The key to doing this involves analyzing the statistical (total variation) distance between two multivariate

hypergeometric distributions. What follows is the main lemma we need; it may be of independent interest:

Lemma 1.6. Suppose $X \sim \operatorname{Hyp}(n,r,k)$ and $Y \sim \operatorname{Hyp}(n,r,\ell+e)$, where $\operatorname{Hyp}(n,r,\ell)$ denotes the (univariate) hypergeometric distribution: i.e., the number of red balls drawn when selecting ℓ balls randomly without replacement from an urn containing n balls, r of which are red. Then

$$d_{\text{TV}}(X, Y) \le .01$$

provided $(1-\frac{r}{n})\min(\ell, n-\ell) \ge Ce^2$, where C is a universal constant. Here $d_{\text{TV}}(\cdot, \cdot)$ denotes total variation distance.

In Section III-E we comment on why the somewhat complicated hypothesis on r, n, ℓ , and e is necessary. The proof of Theorem 1.3, as well as a more complete discussion of the techniques it requires, is presented in Section III.

The challenge in proving Theorem 1.5 is fundamentally different. For Theorem 1.3, we know that the k-junta g is far from all (k-1)-juntas, say, which means it is okay for our \mathcal{F}_{no} functions to be "small tweaks" to g. However when $g=\mathrm{Maj}_k$, small tweaks result in functions that are still close to Maj_k . Thus our \mathcal{F}_{no} functions must be somewhat drastically changed from Maj_k , yet still "look like" Maj_k . We arrange for this by making the \mathcal{F}_{no} functions very carefully constructed weighted majority functions on $\frac{4}{3}k$ coordinates. To show that such functions still "look like" Maj_k , we use recently developed multidimensional invariance principles [13], [9]. However these need to be adapted to the case of sums of random vectors which are not independent, but rather are drawn without replacement from a fixed pool of random vectors.

II. PRELIMINARIES AND DEFINITIONS

A. Probability theory

Given two random variables X,Y defined on a common discrete sample space Ω , the *total variation* distance between X and Y is

$$d_{\mathrm{TV}}(X,Y) = \frac{1}{2} \sum_{\omega \in \Omega} |\mathbf{Pr}[X = \omega] - \mathbf{Pr}[Y = \omega]|.$$

When $X=(X_1,\ldots,X_n)\in\mathbb{R}^n$ is a random vector and we let $\mu_i=\mathbf{E}[X_i]$ for $i=1,\ldots,n$, then the *mean* of X is the vector $\mathbf{E}[X]=(\mu_1,\ldots,\mu_n)$ and the *covariance matrix* of X is the $n\times n$ matrix $\mathbf{Cov}[X]$ whose (i,j)th entry is defined by

$$Cov[X]_{i,j} = E[(X_i - \mu_i)(X_j - \mu_j)] = E[X_i X_j] - \mu_i \mu_j.$$

The function $\Phi: \mathbb{R} \to [0,1]$ is used throughout this paper to denote the cumulative density function of the standard normal distribution $\mathcal{N}(0,1)$.

We write $\operatorname{Hyp}(n,m,k)$ to represent the hypergeometric distribution – the distribution on the number t of red balls

drawn when k balls are drawn without replacement from a set of n balls, m of which are red.

The following theorem of Höglund [10] provides useful bounds on the normal approximation of hypergeometric distributions.

Höglund Theorem. Let $S = X_1 + \cdots + X_n$, where X_1, \ldots, X_n are chosen uniformly at random without replacement from $A = \{x_1, \ldots, x_N\}$. Then for all $t \in \mathbb{R}$,

$$\left| \mathbf{Pr}[S \le t] - \Phi\left(\frac{t - n\mu}{\sigma\sqrt{n(1 - n/N)}}\right) \right| \le C \cdot \frac{\sum_{i=1}^{N} |x_i - \mu|^3 / N}{\sigma^3 \sqrt{n(1 - n/N)}},$$

where $\mu = \sum_{i=1}^{N} \frac{x_i}{N}$, $\sigma^2 = \sum_{i=1}^{N} \frac{(x_i - \mu)^2}{N}$, and C is an absolute constant.

Höglund's Theorem yields the following anticoncentration property of hypergeometric distributions.

Corollary 2.1. Let $S \sim \operatorname{Hyp}(n,m,k)$ and let $\sigma^2 = \frac{km}{n} \left(1 - \frac{m}{n}\right) \left(1 - \frac{k}{n}\right)$. Then for every $t \geq 0$,

$$\Pr[S = t] \le \frac{C}{\sigma}$$

where C is an absolute constant.

Proof: Let A be a set containing m ones and n-m zeros. The random variable $S=X_1+\cdots+X_k$ obtained by choosing k variables uniformly at random without replacement from A follows the $\mathrm{Hyp}(n,m,k)$ distributions. Fixing $\mu=km/n$, Höglund's Theorem implies that for any $t\in\mathbb{R}$,

$$\begin{split} \left|\mathbf{Pr}[S \leq t] - \Phi\left(\frac{t-\mu}{\sigma}\right)\right| &\leq c_0 \cdot \frac{\sum |x_i - m/n|^3}{\sum (x_i - m/n)^2} \cdot \frac{1}{\sigma} \leq \frac{c_0}{\sigma}. \\ \text{Since } \mathbf{Pr}[S = t] &= \mathbf{Pr}[S \leq t] - \mathbf{Pr}[S \leq t-1], \text{ we have } \\ \mathbf{Pr}[S = t] &\leq \Phi(\frac{t-\mu}{\sigma}) - \Phi(\frac{(t-1)\mu}{\sigma}) + 2\frac{c_0}{\sigma} \leq \frac{C}{\sigma}, \\ \text{where } C = 2c_0 + \sqrt{2/\pi}. \end{split}$$

B. Influence

The *influence* of the *i*th variable in the function $f:\{0,1\}^n \to \{0,1\}$ is $\mathrm{Inf}_f(i) = \mathbf{Pr}_x[f(x) \neq f(x^{(i)})]$, where the probability is taken over the uniform distribution of $x \in \{0,1\}^n$ and $x^{(i)}$ is the input formed by flipping the value of the *i*th bit in x.

When $\text{Inf}_f(i) > 0$, we say that the *i*th variable is *relevant* in f. A function that contains at most k relevant variables is called a k-junta.

III. GENERAL LOWER BOUND

In this section, we prove Theorem 1.3.

Theorem 1.3 (Restated). Let $g: \{0,1\}^n \to \{0,1\}$ be a k-junta which is ϵ -far from being a (k-e)-junta for some $e \geq 1$. Then any non-adaptive ϵ -tester for g-isomorphism must make at least $\log_2(k'/e^2) - O(1)$ queries, where $k' = \min(k, n-k)$.

On first reading, the reader is encouraged to focus on the simplest case, where e=1. In this case, Theorem 1.3 affirms a conjecture stated in [7], as it shows that for any $\omega(1) \leq k \leq n - \omega(1)$, when g is a k-junta and is ϵ -far from being a (k-1)-junta for some $\epsilon>0$, then testing g-isomorphism is hard.

We now begin the proof of Theorem 1.3. Let g, n, k, k', e, and ϵ be as in the statement of the theorem. Without loss of generality, we may assume that the k relevant coordinates for g are $[k] = \{1, 2, \ldots, k\}$. We write $g_{\text{core}} : \{0, 1\}^k \to \{0, 1\}$ for the restriction of g to these coordinates.

As is standard in property testing lower bounds, the proof of Theorem 1.3 uses Yao's Minimax Principle [18]. Specifically, we construct two probability distributions $\mathcal{F}_{\rm yes}$ and $\mathcal{F}_{\rm no}$ over functions isomorphic to g and functions ϵ -far from being isomorphic to g, respectively. We then show that any deterministic non-adaptive algorithm making $\ll \log_2(k'/e^2)$ queries cannot distinguish with probability at least 1/3 between functions drawn from $\mathcal{F}_{\rm ves}$ or from $\mathcal{F}_{\rm no}$.

A. The distributions \mathfrak{F}_{ves} and \mathfrak{F}_{no}

We define $\mathcal{F}_{\mathrm{yes}}$ in the most natural way, by randomly embedding g_{core} into [n]. More precisely, to obtain a draw $f_{\mathrm{yes}} \sim \mathcal{F}_{\mathrm{yes}}$, we first choose a uniformly random subset $J \subseteq [n]$ of cardinality k. Next, we choose a uniformly random bijection $\sigma: [k] \to J$. Finally, we define $f_{\mathrm{yes}}(x) = g_{\mathrm{core}}(x_{\sigma(1)}, \ldots, x_{\sigma(k)})$. It is clear that every such f_{yes} is isomorphic to g.

As for $\mathcal{F}_{\mathrm{no}}$, we define a draw $f_{\mathrm{no}} \sim \mathcal{F}_{\mathrm{no}}$ as follows: First, we choose a uniformly random subset $J \subseteq [n]$ of cardinality k-e. Next, we choose a uniformly random map $\sigma:[k] \to J$ from among those satisfying the following property: there is one $j_1 \in J$ with e+1 preimages under σ , and the remaining $j \in J \setminus \{j_1\}$ have a unique preimage. Finally, we define $f_{\mathrm{no}}(x) = g_{\mathrm{core}}(x_{\sigma(1)}, \ldots, x_{\sigma(k)})$. Each such f_{no} only depends on the coordinates J and hence is a (k-e)-junta. Thus by the assumption in Theorem 1.3, each such f_{no} is indeed ϵ -far from being isomorphic to g.

To prove Theorem 1.3, it suffices to prove the following:

Theorem 3.1. Let \mathcal{T} be any deterministic non-adaptive q-query testing algorithm for functions $\{0,1\}^n \to \{0,1\}$. Then

$$\left| \Pr_{f_{\text{ves}} \sim \mathcal{F}_{\text{ves}}} [\Upsilon \ acc. \ f_{\text{yes}}] - \Pr_{f_{\text{no}} \sim \mathcal{F}_{\text{no}}} [\Upsilon \ acc. \ f_{\text{no}}] \right| \le O(\frac{2^q e^2}{k'}) + .01.$$

Note that if $q < \log_2(k'/e^2) - c_0$ for a sufficiently large constant c_0 , then the upper bound in this theorem is at most 1/3. From this we deduce Theorem 1.3 immediately using Yao's Minimax Principle.

B. Distance between multivariate hypergeometrics

The typical way to prove a property testing bound such as Theorem 3.1 is as follows. First, we write the q queries

of tester \mathcal{T} as $x^1,\ldots,x^q\in\{0,1\}^n$. We then introduce the *response vector* random variables R_{yes} and R_{no} . Here $R_{\mathrm{yes}}\in\{0,1\}^q$ is defined by drawing $f_{\mathrm{yes}}\sim\mathcal{F}_{\mathrm{yes}}$ and letting $R_{\mathrm{yes}}=\langle f_{\mathrm{yes}}(x^1),\ldots,f_{\mathrm{yes}}(x^q)\rangle$, and R_{no} is defined analogously. Finally, we show that

$$d_{\text{TV}}(R_{\text{yes}}, R_{\text{no}}) \le 2^q \cdot \frac{O(e^2)}{k'} + .01.$$
 (1)

We will in fact prove a stronger statement. To understand it, let's reconsider the complete random processes \mathcal{P}_{yes} and \mathcal{P}_{no} by which the response vectors R_{yes} and R_{no} are generated. We begin by focusing on the "yes" process, \mathcal{P}_{ves} .

Given the tester \mathcal{T} 's queries $x^1, \ldots, x^q \in \{0, 1\}^n$, we think of them as row vectors and arrange them into a $q \times n$ query matrix Q. We will be especially interested in the columns of this matrix Q, the jth column consisting of the jth bits of all the query strings. Abstractly, we define the set of all possible column (types)

$$\mathfrak{C} = \{0, 1\}^q$$
.

Since $|\mathfrak{C}| = 2^q \ll n$, some columns will occur many times in the matrix Q. In fact, we will think of the query matrix Q as being an ordered *multiset* of columns from \mathfrak{C} .

Recalling the definition of $\mathcal{F}_{\mathrm{yes}}$, we think of the first step of $\mathcal{P}_{\mathrm{yes}}$ as choosing k column indices j_1,\ldots,j_k randomly and without replacement from [n]. We next extract columns j_1,\ldots,j_k from Q. We view this as a multiset of columns, and call it the argument multiset S_{yes} . Next, we randomly order the columns in S_{yes} , forming a $q \times k$ argument matrix A_{yes} . Finally, we produce the response vector R_{yes} by applying g_{core} to the argument matrix, row-wise.

The reader can easily verify this process $\mathcal{P}_{\mathrm{yes}}$ generates the correct distribution on the response vector random variable R_{yes} .

The "no" process $\mathcal{P}_{\mathrm{no}}$ is very similar, differing only in the way it generates the argument multiset from the query matrix. Recalling the definition of $\mathcal{F}_{\mathrm{no}}$, we think of $\mathcal{P}_{\mathrm{no}}$ as forming the argument multiset S_{no} by choosing $\ell=k-e$ random columns from Q without replacement, and including an additional e copies of the first-chosen column. The process $\mathcal{P}_{\mathrm{no}}$ then forms the argument matrix A_{no} by again randomly ordering the columns in the argument multiset, and finally produces the response rector R_{no} again by applying g_{core} to A_{no} , row-wise. The reader can again easily verify that $\mathcal{P}_{\mathrm{no}}$ generates the correct distribution on R_{no} .

Because the processes are identical after the argument multiset is formed, a coupling argument immediately implies that

$$d_{\text{TV}}(R_{\text{ves}}, R_{\text{no}}) \le d_{\text{TV}}(S_{\text{ves}}, S_{\text{no}}). \tag{2}$$

This inequality can be extremely lossy, depending on the function $g_{\rm core}$. However, since Theorem 1.3 applies for an extremely broad range of functions, we are almost forced to

design a proof of Theorem 3.1 that uses no properties of the function $g_{\rm core}$. That is, in the absence of additional restrictions on the class of functions considered, there is no obvious way to bound $d_{\rm TV}(R_{\rm yes},R_{\rm no})$ except by $d_{\rm TV}(S_{\rm yes},S_{\rm no})$.

Letting $S_{\rm yes}$ denote the subprocess of $\mathcal{P}_{\rm yes}$ generating $S_{\rm yes}$, and similarly for $S_{\rm no}$, we have reduced proving (1), and hence Theorem 1.3, to the following:

Theorem 3.2. For $S_{\rm yes} \sim S_{\rm yes}, S_{\rm no} \sim S_{\rm no}$, we have

$$d_{\text{TV}}(S_{\text{yes}}, S_{\text{no}}) \le |\mathfrak{C}| \cdot \frac{O(e^2)}{\min(k, n - k)} + .01.$$

The reader can see now why our query complexity lower bound in Theorem 1.3 is only logarithmic; we have $|\mathfrak{C}|=2^q$ competing against $\frac{1}{k}$ in the above bound. Indeed, we can never prove a better-than-logarithmic lower bound if our proof only involves showing statistical closeness of the argument multisets $S_{\rm yes}$ and $S_{\rm no}$. To see this, suppose k=n/2, so n-k=n/2 as well. Then if $2^q\gg n/2$, it is possible that every column in the query matrix is unique. In this case, the total variation distance between argument multisets $S_{\rm yes}$ and $S_{\rm no}$ will be 1 even in the case e=1, because $S_{\rm yes}$ will always consist of unique columns, whereas $S_{\rm no}$ will always have one column duplicated.

Notice that the ordering of the columns in the query matrix Q has proven to be unimportant; we can think of Q simply as an unordered multiset of columns from $\mathfrak C$. Thus Theorem 3.2 is really a statement about the total variation distance between certain multivariate hypergeometric random variables. Specifically, for each column $\mathfrak c \in \mathfrak C$, let $m(\mathfrak c)$ denote the number of copies of $\mathfrak c$ in Q. In process S_{yes} , we choose k random columns from Q without replacement and count the number of copies of each column (type) in the draw. Process S_{no} is similar, except we choose ℓ random columns from Q without replacement, and count an extra e copies of the first-drawn column.

C. Reduction of Theorem 3.2 to two lemmas

This preceding discussion motivates the following notation:

Definition 3.3. Given integers $N, e \ge 1$, $M, L \ge 0$, with $M, L + e \le N$, we define $\lambda_{N,M,L}(e) = d_{\mathrm{TV}}(X,Y)$, where $X \sim \mathrm{Hyp}(N,M,L+e)$ and $Y \sim \mathrm{Hyp}(N,M,L) + e$.

The proof of Theorem 3.2 relies on the following two lemmas. The first lemma is relatively straightforward, and relates the distance between $\$_{\rm yes}$ and $\$_{\rm no}$ to the total variation distance between hypergeometric distributions.

Lemma 3.4.

$$d_{\mathrm{TV}}(S_{\mathrm{yes}}, S_{\mathrm{no}}) \le \sum_{\mathfrak{c} \in \mathfrak{C}: m(\mathfrak{c}) \ne 0} \frac{m(\mathfrak{c})}{n} \cdot \lambda_{n-1, m(\mathfrak{c})-1, \ell-1}(e).$$

The second lemma is a total variation distance bound between (univariate) hypergeometric random variables which may be of independent interest.

Lemma 3.5. There is a universal constant $2 \le \kappa < \infty$ such that for any N, M, L, if $L' = \min(L, N - L)$ satisfies $\frac{ML'}{N} \ge \kappa e^2$, then $\lambda_{N,M,L}(e) \le .01$.

This lemma is in fact identical to the key Lemma 1.6: to see this, one only needs to replace M with r=N-M and use the obvious fact that $\operatorname{Hyp}(N,N-M,L)$ is the same distribution as $L-\operatorname{Hyp}(N,M,L)$.

We briefly comment on why the hypothesis $\frac{ML'}{N}\gg e^2$ is necessary to show that $\operatorname{Hyp}(N,M,L+e)$ and $\operatorname{Hyp}(N,M,L)+e$ are close in total variation distance. For simplicity, first suppose that e=1. It is necessary that that $\frac{ML}{N}\gg 1$; this quantity is the mean of $\operatorname{Hyp}(N,M,L)$, and if it is $\ll 1$ then $X\sim \operatorname{Hyp}(N,M,L+1)$ is likely to be 0 whereas $Y\sim \operatorname{Hyp}(N,M,L)+1$ is at least 1. Second, it is also necessary that $\frac{M(N-L)}{N}=M(1-\frac{L}{N})\gg 1$. To see this, note that if by way of contrast $1-\frac{L}{N}\ll \frac{1}{M}$, then X is concentrated at M and Y is concentrated at M+1. Finally, to understand the hypothesis's dependence on e, suppose M=N/2 and L is quite small. Then $\operatorname{Hyp}(N,M,L)$ is distributed very much like $\operatorname{Binomial}(L,e)$; hence we require $L\gg e^2$ or else the extra +e in Y will dominate the standard deviation of $\operatorname{Binomial}(L,e)$.

We prove Lemmas 3.4 and 3.5 in the next sections, but first we show how Theorem 3.2 follows from the lemmas.

Proof of Theorem 3.2: Note that we may freely assume $k \geq 2e+2$, as otherwise the bound we are trying to prove exceeds 1 (assuming the constant in the $O(\cdot)$ is large enough). Let us introduce the notation N=n-1, $M(\mathfrak{c})=m(\mathfrak{c})-1$, $L=\ell-1$, $L'=\min(L,N-L)$. Then by Lemma 3.4,

$$d_{\text{TV}}(S_{\text{yes}}, S_{\text{no}}) \leq \sum_{\mathbf{c} \in \mathfrak{C}: m(\mathbf{c}) \neq 0} \frac{m(\mathbf{c})}{n} \cdot \lambda_{n-1, m(\mathbf{c})-1, \ell-1}(e)$$

$$= \sum_{0 \leq \frac{M(\mathbf{c})}{N} < \frac{\kappa e^2}{L'}} \frac{m(\mathbf{c})}{n} \cdot \lambda_{N, M(\mathbf{c}), L}(e)$$

$$+ \sum_{\frac{M(\mathbf{c})}{N} \geq \frac{\kappa e^2}{L'}} \frac{m(\mathbf{c})}{n} \cdot \lambda_{N, M(\mathbf{c}), L}(e)$$

$$\leq \sum_{0 \leq \frac{M(\mathbf{c})}{N} < \frac{\kappa e^2}{L'}} \frac{m(\mathbf{c})}{n} + \sum_{\frac{M(\mathbf{c})}{N} \geq \frac{\kappa e^2}{L'}} \frac{m(\mathbf{c})}{n} \cdot .01,$$

where the last inequality uses Lemma 3.5. Since $\sum_{\mathfrak{c} \in \mathfrak{C}} m(\mathfrak{c}) = n$, the second sum above is at most .01. Thus it remains to bound the first sum by $|\mathfrak{C}| \frac{O(e^2)}{\min(k, n-k)}$. There are at most $|\mathfrak{C}|$ summands in this first sum, and for each we have

$$\frac{m(\mathfrak{c})}{n} \le \frac{M(\mathfrak{c}) + 1}{N} \le \frac{\kappa e^2}{L'} + \frac{1}{N} \le \frac{2\kappa e^2}{L'}$$

by the condition of the sum.

To complete the proof, it remains to show that $L'=\min(\ell-1,n-\ell)\geq\Omega(\min(k,n-k)).$ When $\ell-1\leq n-\ell$, then $L'=\ell-1=k-e-1\geq k/2$ by the fact that $k\geq 2e+2.$ And when $n-\ell<\ell-1$, then $L'=n-\ell=n-k+e\geq n-k.$ so $L'\geq \frac{1}{2}\min(k,n-k)$, as we wanted to show.

D. Proof of Lemma 3.4

Let us think of the experiment S_{yes} in an alternate way. We begin by choosing a first column from Q for S_{yes} — call it C_1 . We next decide how many additional copies of C_1 to include into S_{yes} . Call this quantity T. We have

$$T \mid (C_1 = \mathfrak{c}) \sim \text{Hyp}(n-1, m(\mathfrak{c}) - 1, k-1).$$

(Note that $m(\mathfrak{c})-1\geq 0$ always, because \mathfrak{c} won't be chosen if $m(\mathfrak{c})=0$.) So far, S_{yes} consists of T+1 copies of C_1 . Finally, we complete the draw of S_{yes} by choosing k-(T+1) columns without replacement from " $Q\setminus C_1$ ", meaning the multiset of columns formed from Q by removing all copies of C_1 .

We think of the experiment $S_{\rm no}$ in a similar way. Again, we begin by choosing a first column C_1 from Q for $S_{\rm no}$. We next determine how many additional copies of C_1 there will be from among the remaining $\ell-1$ choices. Calling this quantity U, we have

$$U \mid (C_1 = \mathfrak{c}) \sim \text{Hyp}(n-1, m(\mathfrak{c}) - 1, \ell - 1).$$

Recall, however, that in $\mathcal{S}_{\mathrm{no}}$, we include an additional e copies of C_1 into S_{no} . Hence S_{no} ends up with U+e+1 copies of C_1 . Finally, we complete S_{no} by adding $\ell-(U+1)$ columns drawn without replacement from $Q\setminus C_1$.

Let V=U+e. We claim that by coupling the random variables $T\mid (C_1=\mathfrak{c})$ and $V\mid (C_1=\mathfrak{c}),$ we couple S_{yes} and $S_{\mathrm{no}}.$ This follows immediately from the two descriptions, as then T+1=V+1=U+e+1, and $k-(T+1)=\ell+e-(V+1)=\ell-(U+1).$ Hence

$$\begin{split} d_{\mathrm{TV}}(S_{\mathrm{yes}}, S_{\mathrm{no}}) & \leq \sum_{\mathfrak{c} \in \mathfrak{C}} \mathbf{Pr}[C_1 = \mathfrak{c}] \cdot \\ d_{\mathrm{TV}}(T \mid (C_1 = \mathfrak{c}), V \mid (C_1 = \mathfrak{c})). \end{split}$$

On one hand, $\Pr[C_1 = \mathfrak{c}]$ is simply $\frac{m(\mathfrak{c})}{n}$. On the other hand, we have

$$T \mid (C_1 = \mathfrak{c}) \sim \text{Hyp}(n-1, m(\mathfrak{c}) - 1, \ell + e - 1),$$

 $V \mid (C_1 = \mathfrak{c}) \sim \text{Hyp}(n-1, m(\mathfrak{c}) - 1, \ell - 1) + e.$

So by definition, $d_{\text{TV}}(T \mid (C_1 = \mathfrak{c}), V \mid (C_1 = \mathfrak{c})) = \lambda_{n-1, m(\mathfrak{c})-1, \ell-1}(e)$, and hence

$$d_{\mathrm{TV}}(S_{\mathrm{yes}}, S_{\mathrm{no}}) \leq \sum_{\mathfrak{c} \in \mathfrak{C}: m(\mathfrak{c}) \neq 0} \frac{m(\mathfrak{c})}{n} \cdot \lambda_{n-1, m(\mathfrak{c})-1, \ell-1}(e),$$

as claimed.

E. Proof of Lemma 3.5

Recall that $L' = \min(L, N - L)$,

$$\frac{ML'}{N} \ge \kappa e^2,\tag{3}$$

and our goal is to bound $\lambda_{N,M,L}(e) = d_{\mathrm{TV}}(X,Y) \leq .01$, where $X \sim \mathrm{Hyp}(N,M,L+e)$ and $Y \sim \mathrm{Hyp}(N,M,L)+e$.

We begin by coupling X and Y, as follows. Imagine drawing balls randomly and without replacement from an urn containing N balls, M of which are white. We draw L+e balls from the urn. We let X be the number of white balls among all balls drawn; we let Y be the number of white balls among the first L balls drawn, plus e. Note that $X \leq Y$ always under this coupling.

Let us now compare the probability mass functions of X and Y. The integers u < e can be in X's range but not Y's; the integers $u > \min(M, L + e)$ can be in Y's range but not X's. The remaining integers are in the range of both X and Y, and we have

$$\frac{\mathbf{Pr}[X=u]}{\mathbf{Pr}[Y=u]} = \frac{\binom{M}{u}\binom{N-M}{L+e-u}}{\binom{N}{L+e}} / \frac{\binom{M}{u-e}\binom{N-M}{L+e-u}}{\binom{N}{L}}$$

$$= \frac{\binom{M}{u}}{\binom{M}{u-e}} \cdot \frac{\binom{N}{L}}{\binom{N}{L+e}}$$

$$= \frac{(M-u+e)(M-u+e-1)\cdots(M-u+1)}{u(u-1)\cdots(u-e+1)} \cdot \frac{\binom{N}{L}}{\binom{N}{L}}.$$

Evidently (and unsurprisingly), this ratio is a decreasing function of u. Letting t be the largest integer for which the ratio is at least 1, we conclude that

$$\mathbf{Pr}[X = u] \ge \mathbf{Pr}[Y = u] \text{ iff } u \le t.$$

It follows immediately that

$$d_{\text{TV}}(X, Y) = \mathbf{Pr}[X \le t] - \mathbf{Pr}[Y \le t].$$

But by our coupling,

$$\begin{aligned} \mathbf{Pr}[X \leq t] - \mathbf{Pr}[Y \leq t] &= \mathbf{Pr}[X \leq t \cap Y > t] \\ &- \mathbf{Pr}[X > t \cap Y \leq t] \\ &= \mathbf{Pr}[X \leq t \cap Y > t], \end{aligned}$$

since $X \leq Y$ always. Our goal, then, is to bound

$$d_{\text{TV}}(X,Y) = \mathbf{Pr}[X \le t \cap Y > t]. \tag{4}$$

We will in fact prove something slightly stronger: we will show that for any value of t, the right-hand side of (4) is small.

To analyze (4) we recall the ball and urn process defining X and Y. Having drawn L+e balls, let W be the number of white balls among the *last* e balls drawn, and let Z be the number of white balls among the first L. Thus X=W+Z and Y=e+Z. As a first observation, we may note that

if W = e then X = Y and hence the event in (4) does not occur. I.e.,

$$d_{\mathrm{TV}}(X,Y) \le \mathbf{Pr}[W \ne e] \le e(1 - \frac{M}{N}),\tag{5}$$

where we used a union bound over each of the last e balls being non-white. Now by (3),

$$e \le \sqrt{\frac{1}{\kappa} \frac{ML'}{N}} \le \sqrt{\frac{L'}{\kappa}} \le .001\sqrt{N},$$
 (6)

if we assume κ large enough. It follows that we may additionally assume

$$M \le N - .01\sqrt{N}$$
 \Leftrightarrow $1 - \frac{M}{N} \ge \frac{.01}{\sqrt{N}}$ (7)

because otherwise the bound in (5) is at most $.001\sqrt{N}$. $\frac{.01}{\sqrt{N}} = .00001$, which establishes the theorem with room to spare. We also use this opportunity to mention that

$$M \ge 2e$$
, $L' \ge 2e$, (and hence certainly $N \ge 2e$) (8)

follow easily from (3).

We next give a more refined upper bound on (4). By conditioning on ${\cal W}$ we have

$$\begin{split} d_{\text{TV}}(X,Y) &= \mathbf{Pr}[X \leq t \cap Y > t] \\ &= \sum_{i=0}^{e-1} \mathbf{Pr}[W = i] \, \mathbf{Pr}[t - e < Z \leq t - i \mid W = i]. \end{split}$$

Now $Z\mid (W=i)$ has distribution $\operatorname{Hyp}(N-e,M-i,L)$ (and note that $M-i\geq M-e\geq 0$ by (8)). Let us write $\sigma^2=L(1-\frac{L}{N-e})\frac{M-i}{N-e}(1-\frac{M-i}{N-e}).$ Applying Corollary 2.1 and a union bound we get

$$\begin{split} d_{\mathrm{TV}}(X,Y) & \leq & \sum_{i=0}^{e-1} \mathbf{Pr}[W=i] \cdot (e-i) \frac{C}{\sigma} \\ & \leq & \max_{0 \leq i < e} \left\{ \frac{C}{\sigma} \right\} \cdot \sum_{i=0}^{e-1} \mathbf{Pr}[W=i](e-i) \\ & = & \max_{0 \leq i < e} \left\{ \frac{C}{\sigma} \right\} \cdot \mathbf{E}[e-W]. \end{split}$$

We have $W \sim \mathrm{Hyp}(N,M,e),$ and thus $\mathbf{E}[e-W] = e(1-\frac{M}{N}).$ And by definition,

$$\max_{0 \le i < e} \left\{ \frac{C}{\sigma} \right\} = \max_{0 \le i < e} \left\{ \frac{C}{\sqrt{L(1 - \frac{L}{N-e})(\frac{M-i}{N-e})(1 - \frac{M-i}{N-e})}} \right\}$$

$$\le \frac{C}{\sqrt{L(1 - \frac{L}{N-e})(\frac{M-e}{N-e})(1 - \frac{M}{N-e})}}.$$

Thus we have established

$$d_{\text{TV}}(X,Y) \le \frac{Ce}{\sqrt{L(1-\frac{L}{N-e})}} \cdot \frac{1-\frac{M}{N}}{\sqrt{1-\frac{M}{N-e}}} \cdot \frac{1}{\sqrt{\frac{M-e}{N-e}}}.$$
 (9)

We will bound the three fractions in (9) one at a time. We begin with the middle one. Note first that

$$\frac{d}{dM} \left(\frac{1 - \frac{M}{N}}{\sqrt{1 - \frac{M}{N - e}}} \right) = -\frac{N - 2e - M}{2N\sqrt{1 - \frac{M}{N - e}}(N - e - M)}.$$

By combining (6) and (7) we get $M \leq N-10e < N-2e$. Hence the derivative above is always negative, implying that $(1-\frac{M}{N})/\sqrt{1-\frac{M}{N-e}}$ is a decreasing function of M on M's range. Hence we may upper-bound this fraction by taking M=0, giving an upper bound of 1. Substituting this into (9) gives

$$d_{\text{TV}}(X,Y) \le \frac{Ce}{\sqrt{L(1-\frac{L}{N-e})}} \cdot \frac{1}{\sqrt{\frac{M-e}{N-e}}}.$$
 (10)

We next examine the fraction on the right. It is at most

$$\frac{1}{\sqrt{\frac{M-e}{N}}} \leq \frac{1}{\sqrt{\frac{M/2}{N}}} = \sqrt{\frac{2N}{M}},$$

where we used (8). By virtue of (3), we can upper-bound this by $\sqrt{\frac{2}{\kappa}} \cdot \frac{\sqrt{L'}}{e}$. Substituting this upper bound into (10) yields

$$d_{\text{TV}}(X,Y) \le C\sqrt{\frac{2}{\kappa}} \cdot \sqrt{\frac{L'}{L(1 - \frac{L}{N - e})}}$$

$$\le .001\sqrt{\frac{L'}{L(1 - \frac{L}{N - e})}}, \tag{11}$$

assuming κ is sufficiently large compared with C.

Finally, we split into two cases, depending on whether $L \leq N/2$. If indeed $L \leq N/2$, then L' = L and we have

$$.001\sqrt{\frac{L'}{L(1-\frac{L}{N-e})}} = \frac{.001}{\sqrt{1-\frac{L}{N-e}}} \le \frac{.001}{\sqrt{1-\frac{N/2}{N-e}}}.$$

But $N-e \geq N-.001\sqrt{N} \geq (2/3)N$ (using (6) and $N \geq 2$ from (8)), so we upper-bound

$$d_{\text{TV}}(X, Y) \le \frac{.001}{\sqrt{1 - \frac{N/2}{(2/3)N}}} = .002 \le .01,$$

as needed. The second case is that $L \ge N/2$, in which case L = N - L' and the bound in (11) is

$$.001\sqrt{\frac{L'}{(N-L')(1-\frac{N-L'}{N-e})}} = .001\sqrt{\frac{L'}{(N-L')\frac{L'-e}{N-e}}}$$
$$= .001\sqrt{\frac{L'}{L'-e}}\sqrt{\frac{N-e}{N-L'}}.$$
(12)

But using (8),

$$\sqrt{\frac{L'}{L'-e}} \leq \sqrt{\frac{L'}{L'/2}} = \sqrt{2},$$

and using L' < N/2,

$$\sqrt{\frac{N-e}{N-L'}} \leq \sqrt{\frac{N}{N-L'}} \leq \sqrt{\frac{N}{N/2}} = \sqrt{2}.$$

Hence the upper bound (12) on $d_{\rm TV}(X,Y)$ is at most $.001\sqrt{2}\sqrt{2}<.01$, as needed.

This completes the proof of Lemma 3.5.

IV. MAJORITY FUNCTIONS

Recall that $\mathrm{Maj}_k: \{-1,1\}^n \to \{-1,1\}$ is defined by $\mathrm{Maj}_k(x) = \mathrm{sgn}(\sum_{i=1}^k x_i)$, where we define $\mathrm{sgn}(0)$ (arbitrarily) to be 1. We sometimes abuse notation by thinking of Maj_k also as a function on $\{-1,1\}^k$. In this section we prove our lower bound for testing Maj_k -isomorphism, Theorem 1.5, which we restate here for convenience:

Theorem 1.5 (Restated). There is a universal $\epsilon_0 > 0$ such that whenever $k \leq (3/4)n$ (and is divisible by 3), any non-adaptive algorithm for ϵ_0 -testing Maj_k -isomorphism must make at least $\Omega(k^{1/12})$ queries.

Note that the result of Theorem 1.5 cannot be handled by Theorem 1.3 since the Maj_k function is o(1)-close to being a (k-e)-junta whenever e=o(k). Specifically, the full version of this paper establishes the following proposition, which is very similar to the problem of computing the noise sensitivity of majority [16].

Proposition 4.1. For 0 < e < k/2, the Maj_k function is ϵ -close to the Maj_{k-e} function, where $\epsilon = 6(e/k)^{1/3}$.

Our proof of Theorem 1.5 reuses much of the framework introduced in the previous section in our testing lower bound for general g. As before, our goal is to construct probability distributions $\mathcal{F}_{\mathrm{yes}}$ and $\mathcal{F}_{\mathrm{no}}$ over functions isomorphic to Maj_k and functions ϵ_0 -far from being isomorphic to Maj_k (respectively) such that any deterministic non-adaptive testing algorithm making $o(k^{1/12})$ queries cannot distinguish with probability at least 1/3 between functions drawn from $\mathcal{F}_{\mathrm{yes}}$ or from $\mathcal{F}_{\mathrm{no}}$.

We define $\mathcal{F}_{\mathrm{yes}}$ just as we did in Section III: a function $f_{\mathrm{yes}} \sim \mathcal{F}_{\mathrm{yes}}$ is obtained by choosing j_1, \ldots, j_k randomly and without replacement from [n] and defining $f_{\mathrm{yes}}(x) = \mathrm{Maj}_k(x_{j_1}, \ldots, x_{j_k})$. The definition $\mathcal{F}_{\mathrm{no}}$, however, is very different from the definition we used in that section to ensure that it is supported on functions far from Maj_k .

To define \mathcal{F}_{no} , we first introduce a certain weighted majority function WgtMaj_k on $\frac{4}{3}k$ bits (note that $\frac{4}{3}k \leq n$):

$$WgtMaj_{k}(x_{1},...,x_{\frac{4}{3}k}) = sgn\left(\sum_{i=1}^{k/3} \left(\frac{1}{2}x_{4i-3} + \frac{1}{2}x_{4i-2} + \frac{1}{2}x_{4i-1} + \frac{3}{2}x_{4i}\right)\right).$$

$$(13)$$

I.e., WgtMaj_k gives k variables weight $\frac{1}{2}$ and k/3 variables weight $\frac{3}{2}$. This weight pattern is chosen very carefully; see the proof of Lemma 4.7 below. We then define $f_{\text{no}} \sim \mathcal{F}_{\text{no}}$ is obtained by choosing $j_1, \ldots, j_{\frac{4}{3}k}$ randomly and without replacement from [n] and taking $f_{\text{no}}(x) = \operatorname{WgtMaj}_k(x_{j_1}, \ldots, x_{j_{\frac{4}{3}k}})$.

The following proposition, whose proof is included in the full version of this paper, implies that functions in \mathcal{F}_{no} are ϵ -far from Maj_k functions for large enough values of ϵ :

Proposition 4.2. There exist universal constants $\epsilon_0 > 0$ and $k_0 \in \mathbb{N}$ such that when $k \geq k_0$, every function f_{no} in the support of \mathfrak{F}_{no} is ϵ_0 -far from being a k-junta.

Note that we may always assume $k \ge k_0$ as otherwise Theorem 1.5 is trivial. To complete the proof of Theorem 1.5, it suffices to prove the following:

Theorem 4.3. Let \mathcal{T} be any deterministic non-adaptive q-query algorithm for testing isomorphism to Maj_k . Then

$$\left| \Pr_{f_{\text{yes}} \sim \mathcal{F}_{\text{yes}}} [\mathfrak{T} \ \textit{acc.} \ f_{\text{yes}}] - \Pr_{f_{\text{no}} \sim \mathcal{F}_{\text{no}}} [\mathfrak{T} \ \textit{acc.} \ f_{\text{no}}] \right| \leq O\left(\frac{q^{3/2}}{k^{1/8}}\right).$$

To prove Theorem 4.3, we continue to recall the framework developed in Section III. Given a deterministic q-query tester \mathcal{T} , we arrange its q queries $x^1, \ldots, x^q \in \{-1, 1\}^n$ into a $q \times n$ query matrix Q. We again think of two processes $S_{\rm yes}$ and $S_{\rm no}$ for generating argument multisets $S_{\rm yes}$ and $S_{\rm no}$. However in the present case we simply have that S_{ves} chooses k columns at random from Q without replacement, and S_{no} chooses $\frac{4}{3}k$ columns at random from Q without replacement. Again, we imagine that the argument multisets are randomly ordered to form argument matrices: $A_{\rm yes}$ which is $q \times k$, and $A_{\rm no}$ which is $q \times \frac{4}{3}k$. Finally, we obtain the response vector random variable $R_{\text{ves}} \in \{-1, 1\}^q$ by applying Maj_k to A_{yes} row-wise, and the response vector $R_{\text{no}} \in \{-1, 1\}^q$ by applying WgtMaj_k to A_{no} rowwise. It is clear that this distribution on $R_{\rm yes}$ is equivalent to the one given by drawing $f_{
m yes} \sim {\it F}_{
m yes}$ and letting $R_{\text{yes}} = \langle f_{\text{yes}}(x^1), \dots, f_{\text{yes}}(x^q) \rangle$. The analogous statement is true for $R_{\rm no}$. Hence we can prove Theorem 4.3 by showing

$$d_{\text{TV}}(R_{\text{yes}}, R_{\text{no}}) \le O\left(\frac{q^{3/2}}{k^{1/8}}\right).$$
 (14)

We now come to the main difference between our Maj_k lower bound and the general lower bound from Section III. Obviously, we cannot proceed as in Section III by bounding the total variation distance between S_{yes} and S_{no} : this total variation distance is 1, since S_{yes} and S_{no} have disjoint support! (Specifically, the multiset S_{yes} has cardinality k whereas k_{no} has cardinality k_{no} involves adding up the columns in the argument matrix (in k_{no} involves adding up the certain weights), and this addition "loses a lot of information".

More precisely, suppose that we write X_1, \ldots, X_k for the (randomly chosen) columns in argument matrix A_{yes} and let

$$S = X_1 + \cdots + X_k$$
.

Then the response vector R_{yes} is given by taking the sgn of each entry of S; i.e., it is determined by the orthant of \mathbb{R}^q in which S lies. Similarly, if we write $Y_1,\ldots,Y_{\frac{4}{3}k}$ for the columns in argument matrix A_{no} , and let

$$T = \frac{1}{2}Y_1 + \frac{1}{2}Y_2 + \frac{1}{2}Y_3 + \frac{3}{2}Y_4 + \cdots + \frac{1}{2}Y_{\frac{4}{2}k-3} + \frac{1}{2}Y_{\frac{4}{2}k-2} + \frac{1}{2}Y_{\frac{4}{2}k-1} + \frac{3}{2}Y_{\frac{4}{2}k},$$

then $R_{\rm no}$ is determined by the orthant in which T lies. Hence we can establish (14) and thus Theorem 1.5 by proving the following:

Theorem 4.4. Let S and T be defined as above. Then for any union O of orthants in \mathbb{R}^d ,

$$|\mathbf{Pr}[S \in \mathcal{O}] - \mathbf{Pr}[T \in \mathcal{O}]| \le O(q^{3/2}/k^{1/8}).$$

A. Multidimensional invariance

To prove Theorem 4.4 we will use recently developed invariance principle tools [14], [13], [9]. In particular, we quote the following multidimensional results which essentially appear in [13], [9].

Lemma 4.5. (Essentially Theorem 4.1 in [13]; cf. [9].) Let $\psi: \mathbb{R}^q \to \mathbb{R}$ be a thrice continuously differentiable function with uniformly bounded third partial derivatives: $|\psi^{(J)}| \leq \beta$ for all multi-indices $J = (j_1, \ldots, j_q)$ with $|J| = j_1 + \cdots + j_q = 3$. Let $S = S_1 + \cdots + S_m$, where the S_i 's are independent \mathbb{R}^q -valued random variables, and let $T = T_1 + \cdots + T_m$ similarly. Assume that for each $i \in [m]$, S_i and T_i have matching means and covariance matrices: $\mathbf{E}[S_i] = \mathbf{E}[T_i]$ and $\mathbf{Cov}[S_i] = \mathbf{Cov}[T_i]$. Then

$$|\operatorname{\mathbf{E}}[\psi(S)] - \operatorname{\mathbf{E}}[\psi(T)]| \leq O(\beta) \cdot \sum_{i=1}^m \sum_{|J|=3} \left(\operatorname{\mathbf{E}}[|S_i^J|] + \operatorname{\mathbf{E}}[|T_i^J|]\right),$$

where U^J denotes $U_1^{j_1} \cdots U_q^{j_q}$ when $U \in \mathbb{R}^q$.

Lemma 4.6. (Essentially appears in [9].) Let 0 be any union of orthants in \mathbb{R}^q and let S, T be any \mathbb{R}^q -valued random variables. Let r>0. Then there is a certain smooth function ψ satisfying $|\psi^{(J)}| \leq O(1/r^3)$ for all multi-indices J with |J|=3 and such that

$$\begin{aligned} |\operatorname{\mathbf{Pr}}[S \in \mathfrak{O}] - \operatorname{\mathbf{Pr}}[T \in \mathfrak{O}]| &\leq \operatorname{\mathbf{Pr}}[S \in W_r] + |\operatorname{\mathbf{E}}[\psi(S)] - \operatorname{\mathbf{E}}[\psi(T)]|, \\ where \ W_r &= \{x \in \mathbb{R}^q : |x_i| \leq r/2 \ \text{for some} \ i \in [q]\}. \end{aligned}$$

At first, it does not appear as though these tools are of any help to us, because Lemma 4.5 very crucially uses the fact that the random vectors being summed are independent. Whereas, in our Theorem 4.4 the random vectors X_1, \ldots, X_k are certainly not independent, being drawn randomly without replacement from the fixed population Q. The same goes for $Y_1, \ldots, Y_{\frac{4}{3}k}$. Nevertheless, we can still reduce to Lemma 4.5 using a trick: finding random vectors which are conditionally independent.

B. How to handle drawing without replacement

Let us recap the scenario in Theorem 4.4. We have a fixed multiset Q of n columns (vectors) from $\{-1,1\}^q$. We draw k columns randomly from Q without replacement, yielding the vector-valued random variables X_1, \ldots, X_k ; we also define

$$S = X_1 + \cdots + X_k$$
.

Similarly, the vector-valued random variables $Y_1, \ldots, Y_{\frac{4}{3}k}$ are drawn randomly from Q without replacement, and

$$T = \frac{1}{2}Y_1 + \frac{1}{2}Y_2 + \frac{1}{2}Y_3 + \frac{3}{2}Y_4 + \cdots + \frac{1}{2}Y_{\frac{4}{3}k-3} + \frac{1}{2}Y_{\frac{4}{3}k-2} + \frac{1}{2}Y_{\frac{4}{3}k-1} + \frac{3}{2}Y_{\frac{4}{3}k}.$$

To introduce conditional independence, we reimagine how the X_i 's and Y_i 's are drawn. Specifically, we can couple the drawing of these random vectors as follows:

- 1) Define m = k/3 (an integer).
- 2) Randomly partition the columns of Q into m parts Q_1, \ldots, Q_m , each of cardinality 4, along with a leftover set of cardinality $n 4m \ge 0$.
- 3) Independently for each $i \in [m]$, choose $X_{3i-2}, X_{3i-1}, X_{3i}$ randomly without replacement from Q_i . Define also $S_i = X_{3i-2} + X_{3i-1} + X_{3i}$.
- 4) Independently for each $i \in [m]$, choose Y_{4i-3} , Y_{4i-2} , Y_{4i-1} , Y_{4i} , randomly without replacement from Q_i (i.e., choose them by randomly ordering the vectors in Q_i). Define also $T_i = \frac{1}{2}Y_{4i-3} + \frac{1}{2}Y_{4i-1} + \frac{1}{2}Y_{4i-2} + \frac{3}{2}Y_{4i}$.

It is easy to see that this coupling gives the correct marginal distributions on X_1,\ldots,X_k and $Y_1,\ldots,Y_{\frac{4}{3}k}$. We also have $S=S_1+\cdots+S_m$ and $T=T_1+\cdots+T_m$. And crucially, S_1,\cdots,S_m are independent *conditioned on any choice of the partition* (Q_1,\ldots,Q_m) , and similarly for T_1,\ldots,T_m . The following lemma will allow us to apply Lemma 4.5; it also explains the choice of the weight pattern $(\frac{1}{2},\frac{1}{2},\frac{1}{2},\frac{3}{2})$:

Lemma 4.7. Conditioned on any choice of the partition (Q_1, \ldots, Q_m) , we have $\mathbf{E}[S_i] = \mathbf{E}[T_i]$ and $\mathbf{Cov}[S_i] = \mathbf{Cov}[T_i]$ for each $i \in [m]$.

Let $W_r = \{x \in \mathbb{R}^d : \exists j \in [d] \ s.t. \ |x_j| \le r \}$ represent the region around the orthant boundaries. The following Lemma gives an upper bound on the probability that our random vector S lands near any orthant boundary:

Lemma 4.8. For $r \ge 1$ it holds that

$$\mathbf{Pr}[S \in W_r] \le O\left(\frac{qr}{\sqrt{m}}\right).$$

We present the proofs of Lemmas 4.7 and 4.8 below. First, let us now combine all these results to complete the proof of Theorem 4.4 and hence Theorem 1.5:

Proof of Theorem 4.4: Let us first condition on a particular partition (Q_1,\ldots,Q_m) . Having done so, S_1,\ldots,S_m become independent, as do T_1,\ldots,T_m . By Lemma 4.7, we

may apply Lemma 4.5. Doing so with the function ψ from Lemma 4.6 (with $r \ge 1$ to be chosen later) yields

$$|\mathbf{E}[\psi(S)] - \mathbf{E}[\psi(T)]| \le O\left(\frac{1}{r^3}\right) \cdot \sum_{i=1}^m \sum_{|J|=3} \left(\mathbf{E}[|S_i^J|] + \mathbf{E}[|T_i^J|]\right). \tag{15}$$

We emphasize that (15) is conditional on a particular (Q_1,\ldots,Q_m) . However, note that we can bound the quantities $\mathbf{E}[|S_i^J|]$ and $\mathbf{E}[|T_i^J|]$ uniformly in (Q_1,\ldots,Q_m) : Each coordinate of S_i is at most 1+1+1=3 in absolute value, and similarly each coordinate of T_i is at most $\frac{1}{2}+\frac{1}{2}+\frac{1}{2}+\frac{3}{2}=3$ in absolute value. Hence each expectation is at most 27, and we can therefore upper-bound the right-hand side of (15) by $O(mq^3/r^3)$, since there are at most q^3 many J's.

Now averaging over the choice of partition (Q_1, \ldots, Q_m) , the triangle inequality implies

$$|\mathbf{E}[\psi(S)] - \mathbf{E}[\psi(T)]| \le O(mq^3/r^3).$$

Here the expectation is over the whole definition of S and T. Substituting this into Lemma 4.6 gives

$$|\mathbf{Pr}[S \in \mathcal{O}] - \mathbf{Pr}[T \in \mathcal{O}]| \le \mathbf{Pr}[S \in W_r] + O(mq^3/r^3).$$

Applying Lemma 4.8 we can bound this by $O(qr/\sqrt{m})+O(mq^3/r^3)$. We optimize by taking $r=m^{3/8}d^{1/2}$, yielding a final upper bound of $O(q^{3/2}/m^{1/8})=O(q^{3/2}/k^{1/8})$ and completing the proof.

C. Proof of Lemma 4.7

It suffices to check the claim for i = 1. Let $\mu \in \mathbb{R}^q$ denote the average of the four vectors in Q_1 . Then

$$\mathbf{E}[S_1] = \mathbf{E}[X_1 + X_2 + X_3] = 3\mathbf{E}[X_1] = 3\mu,$$

using the fact that X_1, X_2, X_3 are identically distributed, and similarly

$$\mathbf{E}[T_1] = \mathbf{E}\left[\frac{1}{2}Y_1 + \frac{1}{2}Y_2 + \frac{1}{2}Y_3 + \frac{3}{2}Y_4\right]$$

= $\left(\frac{1}{2} + \frac{1}{2} + \frac{1}{2} + \frac{3}{2}\right)\mathbf{E}[Y_1] = 3\mu,$

verifying the first claim.

We now verify that $\mathbf{Cov}[S_i] = \mathbf{Cov}[T_i]$. Fix $j, j' \in [q]$ and let us write x_1, x_2, x_3 for the j-coordinate of X_1, X_2, X_3 respectively, and write x'_1, x'_2, x'_3 for the j'-coordinate of X_1, X_2, X_3 . Then

$$\mathbf{E}[(S_1)_j(S_1)_{j'}] = \mathbf{E}[(x_1 + x_2 + x_3)(x_1' + x_2' + x_3')]$$

= $3 \mathbf{E}[x_1 x_1'] + 6 \mathbf{E}[x_1 x_2'],$

where we use the facts that (x_1,x_1) has the same distribution as (x_2,x_2) and (x_3,x_3') , and that (x_1,x_2') has the same distribution as $(x_\ell,x_{\ell'}')$ for any $\ell\neq\ell'\in\{1,2,3\}$. So

$$\mathbf{Cov}[S_1]_{i,i'} = 3 \mathbf{E}[x_1 x_1'] + 6 \mathbf{E}[x_1 x_2'] - \mu_i \mu_{i'}.$$

Similarly, we define y_1, \ldots, y_4 and y'_1, \ldots, y'_4 as the jth and j'th coordinates of Y_1, \ldots, Y_4 , respectively. Then

$$\mathbf{E}[(T_1)_j(T_1)_{j'}] = \mathbf{E}[(\frac{1}{2}y_1 + \frac{1}{2}y_2 + \frac{1}{2}y_3 + \frac{3}{2}y_4) \cdot (\frac{1}{2}y'_1 + \frac{1}{2}y'_2 + \frac{1}{2}y'_3 + \frac{3}{2}y'_4)]$$

$$= (3 \cdot (\frac{1}{2})^2 + (\frac{3}{2})^2) \mathbf{E}[y_1y'_1] + (6 \cdot (\frac{1}{2})^2 + 6 \cdot \frac{1}{2} \cdot \frac{3}{2}) \mathbf{E}[y_1y'_2]$$

$$= 3 \mathbf{E}[y_1y'_1] + 6 \mathbf{E}[y_1y'_2],$$

where we use the facts that (y_1, y_1') has the same distribution as (y_ℓ, y_ℓ') for any $\ell = 2, 3, 4$ and (y_1, y_2') has the same distribution as $(y_\ell, y_{\ell'}')$ for any $\ell \neq \ell' \in \{1, 2, 3, 4\}$. So

$$\mathbf{Cov}[T_1]_{j,j'} = 3 \mathbf{E}[y_1 y_1'] + 6 \mathbf{E}[y_1 y_2'] - \mu_j \mu_{j'}.$$

Noting that (X_1, X_2) and (Y_1, Y_2) have the same distribution, we get that $\mathbf{Cov}[S_1] = \mathbf{Cov}[T_1]$.

D. Proof of Lemma 4.8

By union-bounding over the q coordinates, Lemma 4.8 reduces to proving the following statement:

Lemma 4.9. Let $r \geq 1$. Suppose we fix any query row $(x_1, \ldots, x_n) \in \{-1, 1\}^n$ from Q and form the random variable

$$s = x_{i_1} + \dots + x_{i_{3m}},$$

where the sequence i_1, \ldots, i_{3m} is drawn randomly without replacement from [n]. Then

$$\mathbf{Pr}[|s| \le r/2] \le O(r/\sqrt{m}).$$

Proof: Let us recall that $k = 3m \le (3/4)n$. Let u denote the number of 1's among x_1, \ldots, x_n . The statement to be proved is precisely equivalent to the following: Let $Z \sim \text{Hyp}(n, u, k)$. Then

$$\Pr[Z \in [k/2 - r/4, k/2 + r/4]] \le O(r/\sqrt{k}). \tag{16}$$

We divide into two cases.

Case 1: $1/4 \le u/n \le 3/4$.: In this case we use Corollary 2.1 and a union bound over the at most r/2 + 1 integers in the range $\lceil k/2 - r/4, k/2 + r/4 \rceil$ to deduce

$$\Pr[Z \in [k/2 - r/4, k/2 + r/4]] \le O(r)/\sigma_{n,u,k},$$

where $\sigma_{n,u,k} = \sqrt{k(1-k/n)(u/n)(1-u/n)}$. We have $1-k/n \ge 1/4$ and also $u/n, 1-u/n \ge 1/4$. Thus $\sigma_{n,u,k} = \Omega(\sqrt{k})$, establishing (16).

Case 2: $u/n \notin [1/4, 3/4]$.: By symmetry, it suffices to treat just one of the cases u/n < 1/4 or u/n > 3/4; say, the former. In this case we have $\mathbf{E}[Z] = k(u/n) \le k/4$, and we have $\mathbf{Var}[Z] = ku/n(1-u/n)(1-k/n) \le k$. Finally, we may assume that $r \le k/2$, as otherwise (16) is trivial. Thus by Chebyshev's Inequality,

$$\Pr[Z \ge k/2 - r/4] \le \Pr[Z \ge (3/8)k] \le \frac{k}{(k/8)^2} = O(\frac{1}{k}),$$

establishing (16) with room to spare.

V. DISCUSSION

We conclude this work by discussing what we feel are promising directions towards closing the basic research problem of characterizing the functions g for which g-isomorphism is testable.

It is possible that Fischer et al. [7]'s positive result, that testing g-isomorphism is easy when g is an O(1)-junta is mostly best possible. We pose the following question:

Question: Suppose $g: \{0,1\}^n \to \{0,1\}$ is an $(n-\ell)$ -junta which is ϵ -far from being an ℓ -junta. Is it true that ϵ -testing g-isomorphism requires $\omega_{\ell}(1)$ queries?

We are not quite bold enough to conjecture that this is true, but we do not know any g which rules it out. Proving the result seems like it might be difficult, but we believe the problem is approachable for the special case when g is a *symmetric* k-junta. Our Theorem 1.4 establishes the result for the simplest symmetric function, Maj_k . It is not too hard to extend our methods to deal with similar symmetric functions; for example, we are able to show (proof omitted) that testing g-isomorphism is hard for a function $g:\{0,1\}^n \to \{0,1\}$ such as

$$g(x) = \begin{cases} 1 & \text{if } k/2 - \sqrt{k} \le \sum_{i=1}^k x_i \le k/2 + \sqrt{k}, \\ 0 & \text{else.} \end{cases}$$

Roughly speaking, we can handle this case because it has only a constant number of "jumps" (just 2, in fact) between 0 and 1 on the main range of $\sum_{i=1}^k x_i$, namely $k/2\pm O(\sqrt{k})$. If, on the other hand, g is a symmetric k-junta with "many" jumps between 0 and 1 on the main range of $\sum_{i=1}^k x_i$ (e.g., if g is Parity_k), then the techniques we used for our general lower bound Theorem 1.3 may begin to apply.

However, we wish to close by drawing attention to a peculiar intermediate case. Suppose g is the following symmetric k-junta:

$$g(x) = \begin{cases} 0 & \text{if } \sum_{i=1}^{k} x_i = 1 \text{ or } 3 \le \sum_{i=1}^{k} x_i < k/2, \\ 1 & \text{if } \sum_{i=1}^{k} x_i \in \{0, 2\} \text{ or } k/2 \le \sum_{i=1}^{k} x_i \le k, \end{cases}$$

In this case g is $o_k(1)$ -close to Maj_k . Hence it would seem at first blush that the few jumps between 0 and 1 when $\sum_{i=1}^k \leq 3$ are irrelevant, and we should have an $\omega_k(1)$ lower bound for testing isomorphism to this g.

But oddly, this is not clear. Because g is so close to Maj_k , it seems we would need to use " $\mathcal{F}_{\operatorname{no}}$ functions" which are fairly different from Maj_k , like the weighted majority function WgtMaj_k introduced in Section IV. However there is a *one-query* test that distinguishes between a function isomorphic to the above g and a function isomorphic to WgtMaj_k : simply query the string $(0,0,\ldots,0)$, which has

²I.e., is the query complexity of the problem bounded below by $\Omega(\log^* \ell)$? Or by $\Omega(\log \ell)$? Or even by $\Omega(\operatorname{poly}(\ell))$?

value 1 under g and value 0 under $\operatorname{WgtMaj}_k!$ We could fix this by changing $\operatorname{WgtMaj}_k(0,\ldots,0)$ to 0, but there are still problems. For example, the tester could query random strings having a 1/k fraction of 1's. For such strings x, $\Pr[g(x)=1]$ will be noticeably higher than $\Pr[\operatorname{Maj}_k(x)=1]$, because there is a good chance that the string x will contain exactly two 1's among the k coordinates on which g depends.

So strangely, even though strings in $\{0,1\}^k$ with zero, one, or two 1's constitute only an $o_k(1) \ll \epsilon$ probability mass, a clever tester can exploit them for its advantage. This makes proving untestability for functions isomorphic to the above g somewhat tricky, and we leave it as a problem for future research.

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