Derandomizing Algorithms on Product Distributions and Other Applications of Order-Based Extraction

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Abstract: Getting the deterministic complexity closer to the best known randomized complexity is an important goal in algorithms and communication protocols. In this work, we investigate the case where instead of one input, the algorithm/protocol is given multiple inputs sampled independently from an *arbitrary unknown* distribution. We show that in this case a strong and generic derandomization result can be obtained by a simple argument.

Our method relies on extracting randomness from "same-source" product distributions, which are distributions generated from multiple independent samples from the same source. The extraction process succeeds even for arbitrarily low min-entropy, and is based on the order of the values and not on the values themselves (this may be seen as a generalization of the classical method of Von-Neumann [26] extended by Elias [8] for extracting randomness from a biased coin.)

The tools developed in the paper are generic, and can be used in several other problems. We present applications to streaming algorithms, and to *implicit probe search* [9]. We also refine our method to handle product distributions, where the *i*'th sample comes from one of several arbitrary unknown distributions. This requires creating a new set of tools, which may also be of independent interest.

Keywords: extractors; derandomization

1 Introduction

A central goal in complexity theory is achieving *de*randomization in as many settings as possible. The object of derandomization is to take computational tasks that can be achieved with the aid of randomness, and find ways to perform them using less randomness, or ideally none at all. We want to achieve derandomization without increasing the use of other resources by much. For example, we would like the amount of time, space, communication, etc. used in the deterministic solution to be similar to the corresponding quantities in the original randomized solution.

In this paper we deal with both algorithms for decision problems and communication complexity protocols. In the first case, a long line of work initiated by [5, 14, 21, 24, 28] shows that, assuming certain circuit lower bounds, any randomized polynomial time algorithm can be converted into a deterministic polynomial time algorithm However, proving such lower bounds seems well beyond reach and in fact, Kabanets and Impagliazzo[17] building on Impagliazzo et. al [13] show that proving lower bounds is necessary for proving such results. For communication complexity, there are exponential separations between deterministic and randomized protocols (see [19]).

It thus seems well motivated to look for relaxed (but still interesting) models where derandomization can be achieved. Consider the case of time-bounded algorithms. A first (very naïve) attempt at such a relaxation might be to require that instead of succeeding on *every* input, we succeed with high probability on any distribution of inputs. Of course, this is no relaxation at all, as we can consider distributions concentrated on one hard input. A natural way to further relax this is to require high-probability of success only on distributions of inputs that can be efficiently sampled. Impagliazzo and Wigderson [15], followed by Trevisan and Vadhan[25], give conditional derandomizations (and unconditional Gap Theorems for BPP) in this model. Another type of relaxation, which we investigate here, is to allow arbitrary distributions on individual inputs, but to require multiple independent samples from the same distribution¹. In this setting, when receiving k inputs for large enough k, we would like our deterministic algorithm to solve all k inputs correctly, at a running time close to k-times the run-

¹We also consider the case of multiple samples that are *not* from the same distribution. Moreover, one might want to consider the case of multiple samples that are correlated in some way, and this might be a direction for further work.

ning time of the randomized algorithm. Note that in the case of a distribution concentrated on one hard input, the running time on this input will be amortized over k instances. Similarly, we would like a deterministic communication protocol that when receiving kinputs from an arbitrary distribution (over the inputs of *both* parties) solves all instances correctly with a number of communication bits that is close to k times the number of bits used by the randomized protocol. We show that such results can be achieved by simple argument. We show that our constructions are almost optimal, in some sense. Here is a concrete example, which gives the feel of the parameters.

1.1 A Motivating Example and Result

Consider the equality problem in communication complexity: Alice and Bob receive *n*-bit strings xand y, respectively. They want to decide whether x = y. The deterministic communication complexity is *n*, and shared randomness reduces this to O(1). Repeating the randomized protocol we get that for any k, $O(\log k)$ communication bits suffice such that Alice and Bob will have the incorrect answer with probability at most 1/100k.

Consider the setting discussed above: Alice and Bob are now given k-tuples of instances (x_1, \ldots, x_k) and (y_1, \ldots, y_k) respectively, such that each pair (x_i, y_i) is sampled independently from the same arbitrary unknown distribution D. Obviously, we have a deterministic protocol that uses $k \cdot n$ communication bits for solving the entire sequence correctly, and a public coin randomized protocol using $O(k \log k)$ communication bits solving the entire sequence correctly with high probability. We show that when $k > c \cdot n \log n$ for some universal constant c, there exists a deterministic protocol using $O(k \log k)$ communication bits, which solves all instances correctly with probability 2/3. This result is almost optimal if $k < n/\log n$ facing the same hard input k times any deterministic protocol must send more than $k \log k$ bits to succeed.

1.2 Main Results

The parameters presented above are derived from the following theorem.

Theorem 1.1. Let $f : \{0,1\}^n \times \{0,1\}^n \to \{0,1\}$ be any function. Let P_R be a public coin randomized protocol with error ϵ for f using c_r communication bits and r random bits. Let P_D be a deterministic protocol for f using c_d communication bits. For every integer $k \ge \min\{10 \cdot r \cdot n, 100 \cdot r^2 \cdot (c_d/c_r)\}$, there exists a deterministic protocol P using at most $k \cdot (c_r + \log r + 6)$ communication bits, such that for any distribution Don $\{0,1\}^n \times \{0,1\}^n$,

$$\Pr(P((x_1, y_1), \dots, (x_k, y_k))) = (f(x_1, y_1), \dots, f(x_k, y_k)))$$

 $\geq 1 - (\epsilon \cdot k + 2^{-r})$, where $(x_1, y_1), \ldots, (x_k, y_k)$ are drawn independently from D.

We get a similar theorem in the case of algorithms for decision problems.

Theorem 1.2. Let C be the class of product distributions on $(\{0,1\}^n)^k$. Let $f : \{0,1\}^n \to \{0,1\}$ be any function. Let A_R be a randomized, two-sided error, algorithm with error ϵ for f, running in time t_r and using r random bits, and let A_D be a deterministic algorithm for f running in time t_d . For every integer $k \ge 10 \cdot (t_d/t_r) \cdot r$, there exists a deterministic algorithm A that runs in time $k \cdot t_r + \tilde{O}(n \cdot k)$, such that for any distribution D on $\{0,1\}^n$,

 $\Pr(A(x_1, \dots, x_k) = (f(x_1), \dots, f(x_k)) \ge 1 - (\epsilon \cdot k + 2^{-r})$

where (x_1, \ldots, x_k) are drawn independently from D.

Remark 1.3. The reader may wonder whether Theorem 1.2 is interesting as in case the original deterministic algorithm A_D is exponential, we will require an exponential number of independent inputs to use the theorem, and thus still need exponential time. We note again that nothing better is possible in this model (unless a worst case derandomization is achieved). Also one gets more interesting instantiations in the case where A_D 's running time is a larger polynomial than A_R (this is the case in the currently known algorithms for primality testing), and in cases where A_D is polynomial or linear and A_R is sublinear - as is the case in many property testing algorithms. That said, we agree that the communication complexity setting of Theorem 1.1 is probably more convincing.

We also consider the case where the inputs or sampled *several* arbitrary distributions. To formally present our results, we need the following definition.

Definition 1.4. Let D_1, \ldots, D_d be any distributions on $(\{0,1\}^n \times \{0,1\}^n)$. A d-part product distribution (defined using D_1, \ldots, D_d) on $(\{0,1\}^n \times \{0,1\}^n)^k$, is a distribution $X = (X_1, \ldots, X_k)$ such that the X_i 's are all independent, and for each $1 \le i \le k$, X_i is distributed according to D_j , for some $1 \le j \le d$.

Our main theorem for d-part product distributions is as follows.

Theorem 1.5. Fix any positive integers d, n and kand let C be the class of d-part product distributions on $(\{0,1\}^n)^k$. Let $f: \{0,1\}^n \to \{0,1\}$ be any function. Let A_R and A_D be algorithms for f, similarly to Theorem 1.2. For any $0 < \gamma < 1$ and any $k \ge (t_d/t_r) \cdot ((r \cdot 8(d+1)) + \frac{16 \cdot (2d)^5}{\gamma})$, there exists a deterministic algorithm A that runs in time at most $k \cdot t_r + O(n \cdot k \cdot d^2)$ that solves f on C with error $\epsilon \cdot k + \gamma$.

The analogous theorem for communication protocols can be found in the full version.

1.3 Overview of Technique — using 'Content Independent' Extraction

We sketch the proof of Theorem 1.2. The proof of Theorem 1.1 is similar but requires additional tech-We are given a sequence of inputs nical details. (x_1,\ldots,x_k) and we want to deterministically compute $f(x_1), \ldots, f(x_k)$ very efficiently. We distinguish between two cases. In the first there are 'few' distinct inputs among x_1, \ldots, x_k . In this case we simply run the deterministic algorithm A_D on all these inputs and as there are few of them, it will not take too \log^2 In the second case, we have 'many'³ distinct inputs among x_1, \ldots, x_k . In this case, we extract a random string from the sequence, and use that random string to run the randomized algorithm A_R on each input. Let $\{z_1, \ldots, z_s\}$ be the distinct values among x_1, \ldots, x_k . A potential problem with this approach would be that the random string we are using depends on the values z_1, \ldots, z_s and thus might be a 'bad' string for some z_i with high probability. This does not occur as our extraction method is essentially independent of the actual values of the inputs. More specifically, the random string we extract is simply a function of the *order* in which (the potentially multiple instances of) z_1, \ldots, z_s appear in the sequence. This may be seen as a generalization of the classical method of Von-Neumann [26] extended by Elias [8] on extracting randomness from a biased coin. (see also the work of Peres[22])

Remark 1.6. As the inputs in the sequence are independent, a more straightforward approach might have been to apply a (deterministic) multi-source extractor on the inputs. However, multi-source extractors require that each input be sampled from a distribution having a certain min-entropy. Thus, to use a multisource extractor we would have needed to assume the individual inputs come from such a distribution, and would not get results for arbitrary distributions.

Generalizing to multiple distributions We now sketch the ideas used to prove Theorem 1.5. As in the above, the problem essentially reduces to extracting randomness from *d*-part product distributions conditioned on seeing 'many' distinct values. Moreover, the extraction procedure should be independent of the actual values and depend only on their order.

Consider the following simple example: We are given 3 independent samples, such that the first and third

are sampled from a distribution D_1 distributed on values $a, b \in \{0, 1\}^n$. The second sample comes from a distribution D_2 that gives probability one to a value $c \in \{0,1\}^n$ such that $c \neq a, b$. In our terminology, this is a 2-part product distribution D on $(\{0,1\}^n)^3$. Let us look at D conditioned on seeing 3 distinct values. In this case we have a uniform distribution on the sequences (a, c, b) and (b, c, a) (note that we indeed have a *uniform* distribution on these sequences no matter how D_1 is distributed on a and b). This suggests the following method for extracting one bit: Given $x \in (\{0, 1\}^n)^3$, for each pair of indices $i < j \in$ $\{1, \ldots, 3\}$, let $z_{i,j}$ be 1 if $x_i < x_j$ by lexicographical ordering, and 0 otherwise. Now output the sum mod 2 of the $z_{i,j}$'s. Let us call this function the 'all-pairs compare' (APC) function. The APC function has the property that if (x_1, x_2, x_3) are all distinct then any substitution of the order of a pair of elements changes the output value. Note that it is essential that *all* the x_i 's are distinct. For example, it is easy to check that for any a < b, APC(a, a, b) = APC(b, a, a). Thus to extract many random bits, we need many 'blocks' where all inputs are distinct. This suggests the following extraction scheme for *d*-part product distributions conditioned on seeing many distinct values: Given a sequence of inputs, delete the values that appear 'too many times' in the sequence. Now divide the (possibly trimmed) sequence into blocks of d + 1 inputs each. Count the number of blocks such that all inputs in the block are distinct. If there are at least msuch blocks - where m is the number of bits we want to extract - output the APC function on each block. It can be shown that if we start out with enough distinct values (where the exact number is a function of d and m) with high probability we will indeed have m blocks of distinct inputs.

1.4 Related Work

Goldreich and Wigderson [11], using an observation of Noam Nisan, attain results similar to ours for the case of the uniform distribution⁴. Their technique uses seeded extractors, and their correctness argument is different (and would not work for product distributions of arbitrary distributions). Barak, Braverman, Chen and Rao [4] show that randomized communication protocols require about k-times the communication bits to solve k instances with high probability over product distributions. Together with our result this shows that deterministic and randomized protocols have approximately⁵ the same compu-

²In the case of Theorem 1.1 there is an additional complication here of having Alice and Bob conclude what indeed are the distinct input pairs (x_i, y_i) with small communication.

³'many' in this sketch roughly corresponds to the number of random bits used by the randomized algorithm for f.

 $^{^{4}\}mathrm{In}$ fact, for this case their probability of error is smaller than ours.

⁵ with the exception that in our result deterministic algorithms only work for large enough k.

tational power in this setting.

1.5 Applications

Beside our main results, we present two applications of extracting randomness based on the order of elements in a sequence.

1.5.1 Implicit Probe Search

For domain size m and table size n, *implicit probe* search is the problem of searching for an element $x \in [m]$ in a table T containing n elements from [m]using as few queries as possible to T. Arranging T by the regular ordering of [m] and using binary search we can always use at most $\log n$ queries. Yao [27] showed that when m is allowed to be arbitrarily large as a function of n, $\log n$ queries are necessary. Fiat and Naor [9] showed that when m = poly(n), T can be efficiently arranged such that a constant number of queries suffice⁶. The results of [9] are obtained by reducing this problem to the one of explicitly constructing *rainbows*, which may be viewed as a kind of randomness extraction problem (details in the full version). Using this reduction we extend their results, showing that for any $m \leq 2^n$, $O(\frac{\log n}{\log \log n}) = o(\log n)$ queries suffice. Thus, we show that even when the domain is exponentially large there is a scheme significantly better than binary search.

1.5.2 Streaming Algorithms

The data stream model was introduced by Munro et al. [20] (see also the seminal work by Alon, Matias and Szegedy [1]). In this model, an algorithm is presented with a sequence of n elements, and its goal is to estimate a function of it, when it is allowed to pass over the data just once. The algorithm runs in bounded space, usually poly-logarithmic in n. We restrict our attention to algorithms which perform in poly-logarithmic space, and compute a frequency moment of the input. For this problem, it is known that even when the order of the appearance of elements in the stream is chosen in an adversarial manner, the algorithm can approximate the p'th moment for $0 \le p \le 2$ [1, 16], and that this is not possible for moments p > 2, see [3, 7].

A relaxation of the problem assumes that the adversary chooses the values of the elements, but they are presented to the algorithm in a random ordering [2, 6, 12]. For a random ordering of the elements, known bounds only imply that one cannot approximate moments larger than 2.5, although it is believed

that the right lower bound is 2, as in the adversarial ordering case. We show a strong derandomization result in this model, which enables concentrating on proving lower bounds for deterministic algorithms⁷.

We briefly sketch the proof, showing how any randomized algorithm can be simulated by a deterministic one. Let R be a randomized algorithm which approximates (to within any constant) a moment p > 2, with any constant success probability. We present a deterministic algorithm D with the same success probability and approximation ratio, up to an $(1 + n^{-\alpha})$ factor, for a constant $\alpha < 0.25$. To simulate R, D first extracts randomness from the beginning of the stream, using the extractors presented later. If the number of elements required to extract enough randomness is small, it uses this randomness to load a pseudo random generator against space bounded machines, and uses this to simulate the random algorithm on the rest of the input; we prove that with high probability this does not change the quality of the approximation by much. If the randomness requires many elements, the deterministic algorithm simply counts the number of appearances of the first polylog n different elements in the stream; we prove that with high probability this is sufficient to approximate the frequency moments over the entire stream ⁸. Details appear in the full version.

2 Preliminaries

For background on communication complexity we refer the reader to [19]. The following definitions will be useful for discussing high probability of success on a sequence on inputs.

Definition 2.1. Let C be a class of distributions on $(\{0,1\}^n)^k$. Let $f : \{0,1\}^n \to \{0,1\}$ be any function. We say that a deterministic algorithm A solves f on C with error ϵ , if for any distribution X in C, when sampling a sequence (x_1, \ldots, x_k) according to X, A answers correctly on all inputs in the sequence with probability at least $1 - \epsilon$. That is,

$$\Pr(A(x_1,\ldots,x_k) = (f(x_1),\ldots,f(x_k))) \ge 1 - \epsilon,$$

when (x_1, \ldots, x_k) is sampled from X.

⁶Gabizon and Shaltiel[10] showed that for $m = n^{\text{polylog }n}$ a constant number of queries also suffice, although with today's dispersers [9] could have gotten the same result.

⁷Approximating frequency moments is perhaps the most common studied problem in this model. Our results apply to other problems as well. For adversarial order there are separations between randomized and deterministic algorithms

⁸We note that it is impossible to use PRG's and exhaust over the seeds, as the stream appears just once (in the adversarial order model deterministic algorithm are provably weaker than random ones). Also, the coins used by the random algorithm should be uncorrelated with the ordering of the elements; this is the reason for running it only on part of the stream. Getting a strong result requires some fine tuning of the parameters.

We define extractors for families of distributions. Note that we are talking only about *deterministic* extractors.

Definition 2.2. Let C be a class of distributions on a set Ω . A function $E : \Omega \to \{0, 1\}^m$ is an extractor for C with error γ (also called a γ -extractor for C), if for every distribution X in C, E(X) is γ -close to uniform.

3 The Main Result

A *product distribution* consists of multiple independent samples from an *arbitrary* distribution.

Definition 3.1 (Product Distributions). A distribution $X = (X_1, \ldots, X_k)$ on $(\{0, 1\}^n)^k$ is a product distribution if it consists of k independent samples from the same distribution D, where D can be any distribution over $\{0, 1\}^n$.

Our method relies on the fact that product distributions can be written as convex combinations of distributions that just permute a fixed set of values. We now define these distributions.

Definition 3.2 (Multinomial distributions). The class of multinomial distributions on $(\{0,1\}^n)^k$ consists of all distributions of the following form:

Let $z_1, \ldots, z_s \in \{0, 1\}^n$ be distinct strings and let a_1, \ldots, a_s be non-zero positive integers such that $\sum_{i=1}^s a_i = k$. The multinomial distribution X on $(\{0, 1\}^n)^k$ defined by $z_1, \ldots, z_s, a_1, \ldots, a_s$, is the uniform distribution on sequences of n-bit strings of length k such that for $1 \leq i \leq k$, the string z_i appears a_i times in the sequence. Moreover, we call such a distribution X an s-valued multinomial distribution.

Lemma 3.3. Any product distribution is a convex combination of multinomial distributions.

Proof. Let $X = (X_1, \ldots, X_k)$ be a product distribution on $(\{0, 1\}^n)^k$. For any distinct $z_1, \ldots, z_s \in \{0, 1\}^n$ and positive integers a_1, \ldots, a_s such that $\sum_{i=1}^s a_i = k$. Condition X on the event that the distinct strings outputted were z_1, \ldots, z_s and z_i appears a_i times. Given this conditioning X, because of the independence of X_1, \ldots, X_k , any sequence where z_i appears a_i times has equal probability, and therefore we get a multinomial distribution. Writing X as a convex combination of such conditional distributions, the lemma follows.

For positive integers k, a_1, \ldots, a_s such that $\sum_{i=1}^{s} a_i = k$, the multinomial coefficient $\binom{k}{a_1,\ldots,a_s}$ is the number of different sequences of length k consisting of s distinct elements such that the *i*'th element

appears a_i times: $\binom{k}{a_1,\ldots,a_s} = \frac{k!}{a_1!\cdots a_s!}$. We use the following estimate:

Lemma 3.4. For any integers $s \leq k$ with $k \geq 32$ and $s \geq 4$ we have $\log \binom{k}{a_1,..,a_s} \geq \frac{s \cdot \log k}{4}$.

The following claim will enable us to convert uniform distributions over arbitrary sized sets into distributions over binary strings that are close to uniform. A proof can be found in [18].

Claim 3.5. Let N > M be any integers. Suppose that R is uniformly distributed over $\{1, \ldots, N\}$. Then $R \mod M$ is $\frac{1}{\lfloor N/M \rfloor}$ -close to uniform on $\{0, \ldots, M-1\}$.

Lemma 3.6. Fix integers $s \leq k$ with $k \geq 32$ and $s \geq 4$, and let $t = \lfloor \frac{s \cdot \log k}{8} \rfloor$. There exists an extractor $E : (\{0, 1\}^n)^k \to \{0, 1\}^t$ for the class of s-valued multinomial distributions with error $\gamma = 2^{-t}$. E is computable in time $\tilde{O}(k \cdot n)$.

Proof. Given a sequence (x_1, \ldots, x_k) , let $z_1 < z_2 \ldots < z_s$ be the distinct elements that appear in the sequence, where < denotes the lexicographical ordering. For $i = 1, \ldots, s$ denote by a_i the number of times z_i appears in the sequence. Let S be the set of sequences of length k over $\{1, \ldots, s\}$ such that i appears a_i times. Then $|S| = \binom{k}{a_1, \ldots, a_s}$. The work of Ryabko and Matchikina[23] gives a correspondence of S with $\{1, \ldots, \binom{k}{a_1, \ldots, a_s}\}$ computable in time $\tilde{O}(k \cdot n)$. ⁹ Let r be the image of the sequence (x_1, \ldots, x_k) in $\{1, \ldots, \binom{k}{a_1, \ldots, a_s}\}$ through this correspondence. Define $E(x_1, \ldots, x_k) \triangleq r \pmod{2^t}$. For any s-valued multinomial distribution X, r is uniformly distributed. Thus using Claim 3.5, E(X) will be γ -close to uniform for $\gamma \leq \frac{1}{\lfloor \binom{k}{a_1, \ldots, a_s} \rfloor / 2^t \rfloor} \leq \frac{1}{2^t}$, where we used the definition of t and Claim 3.4 in the second inequality.

A basic principle in this work, is that when we restrict our input distribution to a component that only 'reorders' a fixed set of values, we can use randomness extracted from the input to run our algorithm or protocol. The following definition and two lemmata formalize this.

Definition 3.7. We say a distribution X on $(\{0,1\}^n)^k$ is same-valued, if there is a fixed set of values $\{z_1,\ldots,z_s\} \subseteq \{0,1\}^n$, such that the support of X consists of sequences x_1,\ldots,x_k such that the set of distinct values in each sequence is exactly $\{z_1,\ldots,z_s\}$.

 $^{^9[23]}$ do this for the case s=2, but it is easy to reduce to this case.

Lemma 3.8. Let C be a class of same-valued distributions on $(\{0,1\}^n)^k$. Let $E: (\{0,1\}^n)^k \to \{0,1\}^m$ be a γ -extractor for C computable in time t_E . Fix any $f: \{0,1\}^n \to \{0,1\}$ and any $\epsilon > 0$, and let A_R be a randomized algorithm computing f with error ϵ running in time t_r . Then, there exists a deterministic algorithm A running in time $k \cdot t_r + t_E$ that solves f on C with error $\epsilon \cdot k + \gamma$.

Proof. Given $x_1, \ldots, x_k \in (\{0, 1\}^n)^k$, A computes $r = E(x_1, \ldots, x_k)$ and outputs $A_R(x_i, r)$ for every $i \in [k]$. The probability that a uniformly chosen $r \in \{0, 1\}^m$ is bad for some x_i is at most $\epsilon \cdot k$. Thus, the probability that r = E(X) is bad for some x_i is at most $\epsilon \cdot k + \gamma$.

We are now ready to prove the theorems stated in the introduction.

Proof of Theorem 1.2. Given k instances x_1, \ldots, x_k , let $z_1, z_2 \ldots, z_s$ be the distinct elements that appear in x_1, \ldots, x_k . The algorithm A works as follows. Denote $t = \lfloor \frac{s \cdot \log k}{8} \rfloor$. If $t \leq r$ or $s \leq 4$, we run A_D on z_i for every $i \in [s]$. This takes time $t_d \cdot s \leq t_d \cdot 10 \cdot r \leq k \cdot t_r$, which satisfies the theorem for the chosen value of k. Thus we can assume $t \ge r$ and $s \ge 4$. In this case, we compute $y = E(x_1, \ldots, x_k)$, where E is the extractor for multinomial distributions from Lemma $3.6.^{10}$ We now apply the randomized algorithm A_R on all inputs using y as randomness. Let X be a product distribution on $(\{0,1\}^n)^k$. By Lemma 3.3, X is a convex combination of multinomial distributions. Thus, it is enough to prove the theorem in the case that X itself is a multinomial distribution. But in this case, as a multinomial distribution is same-valued the claim follows immediately from Lemma 3.8. As the running time of E is $\tilde{O}(n \cdot k)$, the total running time of A is at most $k \cdot t_r + \tilde{O}(n \cdot k)$.

For communication protocols the proof requires additional details, as Alice and Bob need to communicate to find out what are the distinct input pairs (x, y). Details appear in the full version.

3.1 On the Optimality of our Scheme

Using the notation of Theorem 1.2, our method works given at least $k = O((t_d/t_r) \cdot r)$ samples. Can we get a similar result for smaller k? It is easy to see that to get running time $k \cdot t_r$ we need $k \ge t_d/t_r$. In the full version we prove a stronger lower bound for a restricted type of scheme. We also show that our extraction scheme is almost optimal. **Theorem 3.9.** Let C be the class of product distributions on $(\{0,1\}^n)^k$ conditioned on having at least s distinct values. Then C contains a distribution X with $H_{\infty}(X) \leq O(s \cdot \log k)$. Thus any extractor for C with error $\epsilon \leq 1/2$ can extract at most $O(s \cdot \log k)$ bits.

Proof. As a first step to construct X, we define a distribution D on $\{0,1\}^n$ as follows: Fix distinct elements $z_0, \ldots, z_{2s} \in \{0,1\}^n$. D will give z_0 probability $1 - \frac{2s}{k}$ and, for $1 \leq i \leq 2s$, D gives z_i probability 1/k. Let us denote by X the product distribution $D^{\oplus k}$ conditioned on seeing at least s distinct elements. Denote by X' the distribution X conditioned on having between s and 3s appearances of z_1, \ldots, z_{2s} . As the min-entropy of a distribution is at most the log of its support size, using the bound of Lemma 3.4, we have $H_{\infty}(X') \leq O(s \cdot \log k + \log s) = O(s \cdot \log k)$. Note that the log s term came from having 2s + 1 options as to how many appearances of z_1, \ldots, z_{2s} we have.

By Chebychev, with probability at least 1 - 2/swe have between s to 3s appearances of the elements z_1, \ldots, z_{2s} in a sequence of k independent samples from D. Thus, X' has mass at least 1 - 2/s in X and therefore $H_{\infty}(X) = H_{\infty}(X') + O(\log s) = O(s \cdot \log k)$.

4 Handling Multiple Distributions

In this section we show that a similar derandomization for can be achieved when the sequence of inputs is sampled independently from *several* distributions. It is convenient to view d-part product distributions (see Definition 1.4) as convex combinations of certain same-valued distributions.

Definition 4.1. A d-multinomial distribution on $(\{0,1\}^n)^k$ is a distribution $X = (X_1, \ldots, X_k)$ such that there is a partition $C_1 \cup \ldots \cup C_d = [k]$ into disjoint subsets such that for every $i \neq j \in [d]$, $X|_{C_i}$ and $X|_{C_j}$ are independent, and for every $i \in [d]$ X_{C_i} is a multinomial source. It will be convenient to allow some of the C_i 's to be empty. Thus, every d'-multinomial distribution for some $1 \leq d' \leq d$ is also a d-multinomial distribution.

For distinct strings $z_1, \ldots, z_s \subseteq \{0, 1\}^n$ and positive integers a_1, \ldots, a_s such that $\sum_{i=1}^s a_i = k$ denote by $D_{z_1,\ldots,z_s,a_1,\ldots,a_s}^d$ the set of *d*-multinomial distributions whose support consists of sequences where z_i appears a_i times.

Lemma 4.2. A *d*-part product distribution is a convex combination of *d*-multinomial sources.

¹⁰Note that using $t_d, r > 1$ (otherwise the claim is trivial), we get $k \ge 10 \cdot t_d \cdot r \ge 40$, and thus can use Lemma 3.6.

4.1 The All Pairs Extractor

In the following definition, for strings $x, y \in \{0, 1\}^n$ we denote by (x < y) the value 1 if x < y (by lexicographical ordering of strings) and 0 otherwise. For an integer l, define the *l*-string-all-pairs compare function $APC: (\{0,1\}^n)^l \to \{0,1\}$ by

$$APC(x_1, \ldots, x_l) \triangleq \bigoplus_{1 \le i < j \le l} (x_i < x_j),$$

where \oplus denotes addition modulo 2. That is, we take the parity of comparisons between all pairs.

Claim 4.3. Fix integers l and n. The l-string allpairs compare function APC : $(\{0,1\}^n)^l \rightarrow \{0,1\}$ is an extractor with error $\epsilon = 0$ for the subclass $D_{z_1,...,z_l,1,...,1}^d$ of d-multinomial distributions for any d < l.

Proof. We first prove the following claim. Let $x = (x_1, \ldots, x_l) \in (\{0, 1\}^n)^l$ be a sequence such that $x_i \neq x_j$ for all $i < j \in [l]$. Denote by $x_{i \leftrightarrow j}$ the sequence obtained from x by swapping x_i and x_j . We show that for all $i < j \in [l]$, $APC(x) \neq APC(x_{i \leftrightarrow j})$: To see this¹¹ notice that swapping adjacent values changes the value of E. That is, for every $1 \leq i \leq l-1$, $APC(x) \neq APC(x_{i \leftrightarrow i+1})$. Loosely speaking, this is because one comparison has changed and the rest have stayed the same. Formally,

$$\begin{aligned} APC(x_{i\leftrightarrow i+1}) &= APC(x) \oplus (x_i < x_{i+1}) \oplus (x_{i+1} < x_i) \\ &= APC(x) \oplus 1 \end{aligned}$$

Now note that $x_{i \leftrightarrow j}$ can obtained from x by an odd number of swap operations performed on adjacent places: j - (i + 1) swap operations to move x_i to the (j-1)'th position and another j-i operations to move x_i to the *i*'th position. Thus we have shown that $APC(x) \neq APC(x_{i \leftrightarrow j})$ for all $i < j \in [l]$. Returning to the original claim, let X be a distribution in $D^{d}_{z_1,\ldots,z_l,1,\ldots,1}$. Recall that this means there are disjoint subsets $C_1 \cup \ldots \cup C_d = [l]$ such that $X|_{C_i}$ is a multinomial distribution. As d < l there must be an i such that $|C_i| > 1$. Assume w.l.g. that $|C_1| > 1$, and fix two indices $i < j \in C_1$. Look at the distribution X conditioned on a fixing of values in all positions except i and j. Under such a conditioning, we are left we two distinct strings z and z' that are to be assigned in these positions, and as $X|_{C_1}$ is a multinomial distribution, each of the two possible assignments has probability half. From our previous argument it follows that the different assignments will lead to different values of E. Thus, under any such conditioning APC(X) is uniform. Viewing X as a convex combination of such conditional distributions finishes the proof.

4.2 Reducing to All Pairs

It will be useful to talk about *d*-multinomial distributions where 'no value appears too frequently'. The following definition formalizes such a notion.

Definition 4.4. Let X be a d-multinomial distribution on $(\{0,1\}^n)^k$. We say that X is δ -bounded if X belongs to a subclass $D^d_{z_1,\ldots,z_s,a_1,\ldots,a_s}$ of d-multinomial distributions such that for every $i \in [s]$ $a_i \leq \delta \cdot k$. That is, no value z_i appears in more than a δ -fraction of the indices.

The following lemma shows that a general dmultinomial distribution can be converted into a δ bounded one, provided it has enough distinct values.

Lemma 4.5. Fix any $0 < \delta < 1$ and integers n and k. There is a deterministic algorithm F such that for any s-valued d-multinomial distribution X on $(\{0,1\}^n)^k$ with $s \ge (1/\delta) \cdot \log k$, the distribution F(X) is a convex combination of s'-valued δ -bounded d-multinomial distributions on $(\{0,1\}^n)^{k'}$, for some $s' \ge s - (1/\delta) \cdot \log k$ and $k' \le k$.

Proof. Given $x = (x_1, \ldots, x_k)$, F operates as follows:

- 1. Check if there exists a value $z \in \{0, 1\}^n$ such that $x_i = z$ for more than a δ -fraction of the x_i 's. If so, let z be the most common value in the sequence and remove all x_i 's with $x_i = z$.
- 2. If sequence was changed and it is non-empty, repeat first step on the newly obtained sequence.

Each application of the first step on a *d*-multinomial distribution, results in a convex combination of *d*-multinomial distributions. After *m* repetitions of the first step we are left with at most $(1-\delta)^m \cdot k < e^{-\delta \cdot m} \cdot k$ strings. Thus, after $(1/\delta) \cdot \log k$ repetitions we are left with an empty sequence, and therefore the number of repetitions is bounded by $(1/\delta) \cdot \log k$. Since each repetition reduces the number of values by one, the final components are s'-valued for some $s' \geq s - (1/\delta) \cdot \log k$, as required.

Theorem 4.6 shows how to extract many random bits from δ -bounded *d*-multinomial distributions.

Theorem 4.6. Fix any integers n, m, d and k such that (d + 1)|k and $m \leq \frac{k}{d+1}$. The exactor E: $(\{0,1\}^n)^k \rightarrow \{0,1\}^m$ be defined as follows. Given input $x \in (\{0,1\}^n)^k$, first partition x into $\frac{k}{d+1}$ blocks $x^1, \ldots, x^{\frac{k}{d+1}}$, each containing (d+1) n-bit strings. We say a block is good if all (d + 1) n-bit strings in the

¹¹Another way to see this is that if x_1, \ldots, x_l are distinct, the APC function just corresponds to the sign of the permutation which sorts the values. Swapping two elements changes the sign.

block are distinct. If there are at least m good blocks x^{i_1}, \ldots, x^{i_m} , E outputs the all-pairs compare function on each one, $E(x) = APC(x^{i_1}), \ldots, APC(x^{i_m})$. Otherwise, E outputs the 0-string.

Fix any $0 < \gamma < 1$ and let $\delta = \frac{\gamma}{16d^5}$. E is a γ -extractor for the class of s-valued δ -bounded dmultinomial distributions on $(\{0,1\}^n)^k$, whenever $s \ge m \cdot 4(d+1)$.

The theorem will follow easily from the following lemma.

Lemma 4.7. Fix any integers n, s, d and k such that (d+1)|k. Fix any $0 < \gamma < 1$ and let $\delta = \frac{\gamma}{8 \cdot (d+1) \cdot d^4}$. Let X be an s-valued δ -bounded d-multinomial distribution on $(\{0,1\}^n)^k$. Partitioning a string $x \in (\{0,1\}^n)^k$ and defining a good block as in Theorem 4.6, we have

$$\Pr_{x \leftarrow X}\left(x \text{ has less than } \frac{s}{4 \cdot (d+1)} \text{ good blocks}\right) \leq \gamma.$$

Proof. Let $C_1 \cup \ldots \cup C_d = [k]$ be the subsets defining X. That is, $X|_{C_i}$ is some multinomial distribution. We show that after removing frequent values as in Lemma 4.5, each one of the underlying distributions $X|_{C_i}$ is either rare or bounded. Note that if for some $0 < \eta < 1$ and $i \in [d] X|_{C_i}$ is not η -bounded, then $|C_i| \leq \frac{\delta}{\eta} \cdot k$. Taking $\eta = 2\delta \cdot d \cdot (d+1)^2$ we get that at most $\left(\frac{\delta}{2\delta \cdot d \cdot (d+1)^2} \cdot d\right) \cdot k = \frac{1}{2(d+1)^2} \cdot k$ indices belong to sets C_i such that $X|_{C_i}$ is not η -bounded. Thus, at most $(d+1) \cdot \left(\frac{1}{2(d+1)^2} \cdot k\right) = \frac{k}{2(d+1)}$ blocks contain an index $j \in [k]$ belonging to a subset C_i where $X|_{C_i}$ is not η -bounded. Therefore, we have at least $\frac{k}{d+1} - \frac{k}{2(d+1)} = \frac{k}{2(d+1)}$ blocks such that all indices in the block belong to a set C_i such that $X|_{C_i}$ is η bounded. Assume without loss of generality that the first $\frac{k}{2(d+1)}$ blocks have this property. For each i = $1, \ldots, \frac{k}{2(d+1)}$ define a random variable Z_i by $Z_i = 1$ if X^i is bad, and 0 otherwise.

Note that $E(Z_i) = \Pr(Z_i = 1) \leq \frac{(d+1)\cdot d}{2} \cdot \eta =$ $\delta \cdot (d^2)(d+1)^3$. Define $Z = \sum_{i=1}^{\frac{k}{2} \cdot (d+1)} Z_i$. Then, $E(Z) \leq \delta \cdot (d^2)(d+1)^3 \cdot \frac{k}{2 \cdot (d+1)} =$ $\frac{\delta \cdot d^2 \cdot (d+1)^2}{2} \cdot k \leq \delta \cdot 2d^4 \cdot k.$

Therefore, using Markov's inequality, for any $0 < \gamma < 1$, $\Pr(Z > \frac{1}{\gamma} \cdot (\delta \cdot 2d^4 \cdot k)) \leq \gamma$. Conversely, with probability at least $1 - \gamma$, we have at least $\frac{k}{2 \cdot (d+1)} - \frac{\delta}{\gamma} \cdot 2d^4 \cdot k$ good blocks. Finally, noting that $k \geq s$ and using the value of δ we get

$$\frac{k}{2\cdot (d+1)} - \frac{\delta}{\gamma} \cdot 2d^4 \cdot k \geq \frac{s}{2\cdot (d+1)} - \frac{\delta}{\gamma} \cdot 2d^4 \cdot s \geq \frac{s}{4\cdot (d+1)}$$

and the lemma follows.

Proof of Theorem 4.6. Let X be a δ -bounded svalued d-multinomial distribution on $(\{0,1\}^n)^k$. Let $l = \frac{k}{d+1}$. For subsets $Z_1, \ldots, Z_l \subseteq \{0, 1\}^n$, with $|Z_i| \leq d+1$, we define the distribution X_{Z_1,\ldots,Z_l} to be X conditioned on the event that for every $1 \leq i \leq l$, the set of distinct values in X^i is exactly Z_i . We can view X as a convex combination of the distributions X_{Z_1,\ldots,Z_l} . Note that these distributions are simply concatenations of independent *d*-multinomial distributions on $(\{0,1\}^n)^{d+1}$. Call a distribution X_{Z_1,\ldots,Z_l} 'good' if for at least m values $i \in [l], |Z_i| = d + 1$, i.e., the *i*'th block contains d+1 distinct elements. It follows from Lemma 4.7 that the mass of 'good' distributions in the convex combination representing X, is at least $1 - \gamma$. using Claim 4.3, for a good distribution X_{Z_1,\ldots,Z_l} , $E(X_{Z_1,\ldots,Z_l})$ is completely uniform. Thus, E(X) is γ -close to uniform.

Using our conversion from general *d*-multinomial distributions to δ -bounded *d*-multinomial distributions, we get an extractor for general *d*-multinomial distributions.

Corollary 4.8 (Extractors for d-multinomial distributions). Fix any integers n, m, d and k and any $0 < \gamma < 1$. There is a γ -extractor $E : (\{0, 1\}^n)^k \rightarrow \{0, 1\}^m$ for the class of s-valued d-multinomial distributions whenever $s \ge m \cdot 8(d+1) + \frac{16 \cdot (2d)^5}{\gamma} \cdot \log k$. E is computable in time $O(nk \cdot d^2)$

Proof. Let F be the algorithm from Lemma 4.5 for $\delta = \frac{\gamma}{16 \cdot (2d)^5}$ Given $x \in (\{0, 1\}^n)^k$, our extractor E works by first applying F on x. We then possibly add at most d n-bit strings to F(x) to make the number of *n*-bit strings it contains a multiple of d + 1(at each step, we add the lexicographically first string that does not yet appear in F(x)). We then compute E'(F(x)), where E' is the extractor for δ -bounded dmultinomial distributions from Theorem 4.6. Let Xbe an s-valued d-multinomial distribution. Lemma 4.5 guarantees that F(X) is a convex combination of s'-valued δ -bounded d-multinomial distribution for $s' \ge s - (1/\delta) \cdot \log k \ge m \cdot 8(d+1)$. The possible additions make the components of F(X) δ -bounded 2d-multinomial distributions. As $s' \ge m \cdot 8(d+1) >$ $m \cdot 4(2d+1)$ it now follows from Theorem 4.6 that E(X) = E'(F(X)) is γ -close to uniform.

Using Corollary 4.8, the proof of Theorem 1.5 is similar to the one of Theorem 1.2.

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