Tight Computational Indistinguishability Bound of Product Distributions

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Abstract

Assume that $X_0, X_1$ (respectively $Y_0, Y_1$) are $d_X$ (respectively $d_Y$) indistinguishable for circuits of a given size. It is well known that the product distributions $X_0Y_0, X_1Y_1$ are $d_X + d_Y$ indistinguishable for slightly smaller circuits. However, in probability theory where unbounded adversaries are considered through statistical distance, it is folklore knowledge that in fact $X_0Y_0$ and $X_1Y_1$ are $d_X + d_Y - d_X \cdot d_Y$ indistinguishable, and also that this bound is tight.

We formulate and prove the computational analog of this tight bound. Our proof is entirely different from the proof in the statistical case, which is non-constructive. As a corollary, we show that if $X$ and $Y$ are $d$ indistinguishable, then $k$ independent copies of $X$ and $k$ independent copies of $Y$ are almost $1 - (1 - d)^k$ indistinguishable for smaller circuits, as against $d \cdot k$ using the looser bound. Our bounds are useful in settings where only weak (i.e. non-negligible) indistinguishability is guaranteed. We demonstrate this in the context of cryptography, showing that our bounds yield simple analysis for amplification of weak oblivious transfer protocols.
1 Introduction

Computational indistinguishability is a basic concept in computational complexity and cryptography. One of the most basic bounds in this context, which is easy to see using a simple hybrid argument, is that for distributions \( X_0, X_1 \) of distance \( d_X \), and \( Y_0, Y_1 \) of distance \( d_Y \), with \( d_{XY} \) denoting the distance between \( X_0Y_0, X_1Y_1 \), we have that

\[
d_{XY} \leq d_X + d_Y,
\]

which holds both statistically and in the computational setting holds for slightly smaller circuits. However, in probability theory where statistical distance, or equivalently, indistinguishability against unbounded attackers is considered, it is folklore knowledge [Kon12, Lemma 2.2] that a better, tight bound holds:

\[
d_{XY} \leq d_X + d_Y - d_X \cdot d_Y.
\]

It is tight in the sense that for every choice of \( d_X, d_Y \), there exist distributions \( X_0, X_1 \) with distance \( d_X \) and distributions \( Y_0, Y_1 \) with distance \( d_Y \), such that \( d_{XY} = d_X + d_Y - d_X \cdot d_Y \).

The proof of this bound uses coupling [Hol12], and is thus inherently non-constructive. We provide a proof of the tight bound in the computational setting, both uniform and non-uniform, with an additive loss of \( \varepsilon \) which can be made as small as we want, by paying in increasing the running time or circuit size with relation to \( 1/\varepsilon \). To be more specific, for the non-uniform case, we (roughly) show that

**Theorem 1.1** (Informal). Let \( X_0, X_1 \) be \( d_X \) indistinguishable for size \( s_X \) circuits. (Respectively \( Y_0, Y_1, d_Y, s_Y \).) Then, for every \( k \in \mathbb{N} \), we have that \( (X_0, Y_0) \) and \( (X_1, Y_1) \) are \((d_X + d_Y - d_X \cdot d_Y + \varepsilon_k)\) indistinguishable for size \( s_k \) circuits, where

\[
\varepsilon_k \leq (d_Y)^k, \quad s_k \approx \min \{s_Y, s_X/k\}.
\]

**Corollary 1.1** (Informal). Let \( D, Q \) be distributions that are \( d \) indistinguishable for size \( s \) circuits. Then, for every \( m \in \mathbb{N} \) and \( \varepsilon \), we have that \( D^\otimes m, Q^\otimes m \) are \( (1 - (1 - d)^m + \varepsilon) \) indistinguishable for size \( s_{m,\varepsilon} \) circuits, where

\[
s_{m,\varepsilon} \approx s(1 - d)^m / \log(1/\varepsilon).
\]

And we also show similar results in the uniform setting. First we prove the isolated non-uniform analog, which we later show how to generalize to the uniform computation model. Then we show how to arrive at the corollary, that if the computational distance between \( X \) and \( Y \) is at most \( d \), then the computational distance between the \( k \)-product of \( X \) and the \( k \)-product of \( Y \) is upper bounded by almost \( 1 - (1 - d)^k \) for smaller circuits, as against \( d \cdot k \) resulted by the looser well known bound, which in particular may be larger than 1. The proof of the corollary essentially follows by (carefully) applying the bound of the isolated case again and again. It should be noted that the difference between the bounds is especially interesting when \( k \) is not very small compared to \( 1/d \). For example, if \( d = 0.5, k = 3 \), the tight bound is 0.875 while the looser bound of 1.5 \( \geq 1 \) is trivial. Finally, we show how these bounds may be used for amplification of weak oblivious transfer protocols [DKS99, Wul07], in the computational setting, providing an alternative simple analysis to the fact that the information theoretic amplification process also works computationally. Some of the techniques and statement formulations presented in this paper were inspired by Levin’s proof of the XOR Lemma [Lev87], and its presentation in [GNW95].
2 Definitions

For a distribution $D$, denote by $D^\otimes k$ the distribution of $k$ independent copies of $D$. For distributions $X_0, X_1$ over $\Omega$, a distinguisher is a boolean $A : \Omega \to \{0, 1\}$, and we let $\text{adv}_A^+(X_0, X_1) := \mathbb{E}[A(X_1) - A(X_0)]$. (The expectation is also over $A$ if it is not deterministic.) We say that distributions $X_0, X_1$ are $d$ indistinguishable for size $s$ circuits if for any such circuit $C$, we have that $\text{adv}_C^+(X_0, X_1) \leq d$. For distributions $X, Y$ we will denote by $(X, Y)$ the product distribution, given by two independent samples from $X$ and $Y$. We denote by $B(p)$ the Bernoulli distribution with parameter $p$, and more generally by $B^\ell(p)$ the distribution that is equal to $1^\ell$ with probability $p$ and otherwise 0\(^\ell\). For a string $s$, we denote by $s[i]$ the $i$th bit of $s$. We will denote by $[m]$ the set $\{1, \ldots, m\}$. We denote by $X_{1/2}$ the distribution given by $b \leftarrow \{0, 1\}, x \leftarrow X_b$. An ensemble of distributions $X = \{X_n\}$ is efficiently samplable if there exists a uniform PPT sampler that given $1^n$ outputs a sample from $X_n$.

2.1 Notation

When the same distribution is used multiple times in a single expression, e.g. $(f(D), g(D))$ for $D$, it should be interpreted that a single value $d \leftarrow D$ is sampled and given to both $f$ and $g$, rather than two independent samples.

3 The Non-Uniform Bounds and Tightness

Let us start with the non-uniform version as it is more simple and clean. The uniform version will be a generalization of the ideas presented below. Roughly speaking, we show that given a distinguisher $C$ for $(X_0, Y_0), (X_1, Y_1)$, if $C(x, \cdot)$ is not a good enough distinguisher between $Y_0, Y_1$ for all values of $x$, then we can build an amplifier for $X_0, X_1$ distinguishers. We then use this amplifier to turn the trivial distinguisher that always outputs 1 into a good enough distinguisher.

**Theorem 3.1.** Let $X_0, X_1$ be distributions over $\ell_X$ bits that are $d_X$ indistinguishable for size $s_X$ circuits. (Respectively $Y_0, Y_1, \ell_Y, d_Y, s_Y$.) Then, for every $k \in \mathbb{N}$, we have that $(X_0, Y_0)$ and $(X_1, Y_1)$ are $(d_X + d_Y - d_X \cdot d_Y + \varepsilon_k)$ indistinguishable for size $s_k$ circuits, where

$$\varepsilon_k := \frac{(d_Y)^k \cdot d_X (1 - d_Y)}{1 - (d_Y)^k} \leq (d_Y)^k, \quad s_k := \min \left\{ s_Y - \ell_X, \frac{s_X - 1}{k} - 5\ell_Y - 1 \right\}.$$

**Remark 3.1.** We note that our starting point, $k = 1$, matches the simple hybrid argument bound of $d_X + d_Y$ since $\varepsilon_1 = d_X \cdot d_Y$, and as $k$ grows larger our bound gets closer and closer to the tight bound of $d_X + d_Y - d_X \cdot d_Y$, while the circuits bound grows smaller. Also note that the bound is asymmetric with respect to the circuit size bounds. This asymmetry is important for preserving a similar circuit size when applying the isolated case over and over again. See a similar argument in [GNW95, Section 3].

**Proof.** Assume toward contradiction that for some circuit $C$ of size $s_k$, we have that

$$\text{adv}_C^+((X_0, Y_0), (X_1, Y_1)) > (d_X + d_Y - d_X \cdot d_Y + \varepsilon_k).$$

2
For every fixed \( x \), it must be that \( C(x, \cdot) \) is able to distinguish between \( Y_0 \) and \( Y_1 \) by at most \( d_Y \), otherwise we get a contradiction as the size of this circuit is \( s_k + \ell_X \leq s_Y \). Then, for every candidate distinguisher \( A \) between \( X_0 \) and \( X_1 \), we have that

\[
\text{adv}_{C}^{+} ((X_1, Y_{A(X_1)}), (X_1, Y_1)) \leq d_Y \cdot \Pr [A(X_1) = 0] \\
\text{adv}_{C}^{+} ((X_0, Y_0), (X_0, Y_{A(X_0)})) \leq d_Y \cdot \Pr [A(X_0) = 1]
\]

where \( x, y \leftarrow X_1, Y_{A(X_1)} \) is resulted by \( x \leftarrow X_1, b \leftarrow A(x), y \leftarrow Y_b \). This holds because

\[
\text{adv}_{C}^{+} ((X_1, Y_{A(X_1)}), (X_1, Y_1)) = \mathbb{E} [C(X_1, Y_1) - C(X_1, Y_{A(X_1)})] = \\
= \mathbb{E} [C(X_1, Y_1) - C(X_1, Y_0) | A(X_1) = 0] \cdot \Pr [A(X_1) = 0] + \\
+ \mathbb{E} [C(X_1, Y_1) - C(X_1, Y_1) | A(X_1) = 1] \cdot \Pr [A(X_1) = 1] = \\
= \mathbb{E}_{x \leftarrow X_1 | A(X_1) = 0} [C(x, Y_1) - C(x, Y_0)] \cdot \Pr [A(X_1) = 0] = \\
= \mathbb{E}_{x \leftarrow X_1 | A(X_1) = 0} \left[ \text{adv}_{C(x, \cdot)}^{+} (Y_0, Y_1) \right] \cdot \Pr [A(X_1) = 0] \leq d_Y \cdot \Pr [A(X_1) = 0]
\]

and using a symmetric argument for the second inequality. Using that (in general)

\[
\sum_{i \in [n]} \text{adv}_{C}^{+} (D_i, D_{i+1}) = \text{adv}_{C}^{+} (D_1, D_{n+1})
\]

we conclude that

\[
\text{adv}_{C}^{+} ((X_0, Y_0), (X_1, Y_1)) = \text{adv}_{C}^{+} ((X_0, Y_0), (X_0, Y_{A(X_0)})) + \\
+ \text{adv}_{C}^{+} ((X_0, Y_{A(X_0)}), (X_1, Y_{A(X_1)})) + \text{adv}_{C}^{+} ((X_1, Y_{A(X_1)}), (X_1, Y_1))
\]

and thus

\[
\text{adv}_{C}^{+} ((X_0, Y_{A(X_0)}), (X_1, Y_{A(X_1)})) = \text{adv}_{C}^{+} ((X_0, Y_0), (X_1, Y_1)) - \\
- \text{adv}_{C}^{+} ((X_1, Y_{A(X_1)}), (X_1, Y_1)) - \text{adv}_{C}^{+} ((X_0, Y_0), (X_0, Y_{A(X_0)})) > \\
> (d_X + d_Y - d_X \cdot d_Y + \varepsilon_k) - (d_Y \cdot \Pr [A(X_1) = 0]) - (d_Y \cdot \Pr [A(X_0) = 1]) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y (1 - \Pr [A(X_1) = 0] - \Pr [A(X_0) = 1]) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y (\Pr [A(X_1) = 1] - \Pr [A(X_0) = 1]) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y (\mathbb{E} [A(X_1)] - \mathbb{E} [A(X_0)]) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y \cdot \text{adv}_{A}^{+} (X_0, X_1).
\]

In other words, we can build a new distinguisher \( A' \) for \( X_0, X_1 \) by applying \( A \) to our input \( x \), sampling \( y \leftarrow Y_{A(x)} \) and feeding \( (x, y) \) to \( C \), and have that

\[
\text{adv}_{A'}^{+} (X_0, X_1) > (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y \cdot \text{adv}_{A}^{+} (X_0, X_1).
\]

If we start from \( A_0 \) being the trivial distinguisher that always outputs 1 and keep repeating
this process for \( k \) steps, we get that

\[
\text{adv}^*_A(X_0, X_1) > (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y \cdot \text{adv}^*_{A_{k-1}}(X_0, X_1) > \\
> (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y \cdot (d_X - d_X \cdot d_Y + \varepsilon_k) + (d_Y)^2 \cdot \text{adv}^*_{A_{k-2}}(X_0, X_1) > \\
> \cdots > (d_X - d_X \cdot d_Y + \varepsilon_k) \sum_{i=0}^{k-1} (d_Y)^i + (d_Y)^k \cdot \text{adv}^*_{A_{k}}(X_0, X_1) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) \sum_{i=0}^{k-1} (d_Y)^i = \frac{(d_X - d_X \cdot d_Y + \varepsilon_k) (1 - (d_Y)^k)}{1 - d_Y} = \\
= \frac{(d_X (1 - d_Y) + \frac{(d_Y)^k \cdot d_X (1 - d_Y)}{1 - (d_Y)^k}) (1 - (d_Y)^k)}{1 - d_Y} = \left( d_X + \frac{(d_Y)^k \cdot d_X}{1 - (d_Y)^k} \right) (1 - (d_Y)^k) = \\
= d_X (1 - (d_Y)^k) + (d_Y)^k \cdot d_X = d_X.
\]

And so, we have concluded that \( A_k \) distinguishes \( X_0 \) from \( X_1 \) with advantage better than \( d_X \). Next, for the circuit size, in order to implement \( A_k \) we start by applying \( A_{k-1} \), sample \( y_0 \leftarrow Y_0, y_1 \leftarrow Y_1 \), use a multiplexer to choose \( y \leftarrow y_b \) where \( b \) is the output gate of \( A_{k-1} \), and finally use the circuit \( C \). Instead of sampling \( y_0, y_1 \), we can simply use non-uniformity to hard-code the best samples, at the cost of \( 2\ell_Y \) gates. Implementing the multiplexer can be done using \( 3\ell_Y + 1 \) gates, with one gate computing \( \neg b \) and for every \( i \in [\ell_Y] \) another 3 gates to compute \( y[i] = (y_0[i] \land \neg b) \lor (y_1[i] \land b) \). Overall, we conclude that \( \text{size}(A_k) = \text{size}(A_{k-1}) + 5\ell_Y + 1 + s_k \) and therefore

\[
\text{size}(A_k) = \text{size}(A_0) + k \cdot (5\ell_Y + 1 + s_k) \leq 1 + k \cdot \left( 5\ell_Y + 1 + \left( \frac{s_X - 1}{k} - 5\ell_Y - 1 \right) \right) = s_X
\]

which is a contradiction to our assumption that \( d_X \) is an upper bound on the advantage of size \( s_X \) circuits distinguishing \( X_0 \) from \( X_1 \).

\[\square\]

### 3.1 The N-Fold Case

**Corollary 3.1.** Let \( D, Q \) be distributions over \( \ell \) bits that are \( d \) indistinguishable for size \( s \) circuits. Then, for every \( m \in \mathbb{N} \) and \( \varepsilon \), we have that \( D^{\otimes m}, Q^{\otimes m} \) are \( (1 - (1 - d)^m + \varepsilon) \) indistinguishable for size \( s_{m,\varepsilon} \) circuits, where

\[
s_{m,\varepsilon} = \frac{s - 1}{k_{m,\varepsilon}} - 5m\ell - 1, \quad k_{m,\varepsilon} = \left\lceil \frac{\log(\varepsilon)}{\log(1 - (1 - d)^m + \varepsilon)} \right\rceil \leq \left\lceil \frac{\log(1/d\varepsilon)}{(1 - d)^m - \varepsilon} \right\rceil.
\]

**Proof.** If \( \varepsilon \geq (1 - d)^m \) the statement is trivially true. Otherwise, we start from \( D, Q \) and use Theorem 3.1 to repeatedly add copies of \( D, Q \) for \( m - 1 \) times, using \( k_{m,\varepsilon} \) set at the statement, where each time the added copy of \( D, Q \) is treated as \( X_0, X_1 \) and \( D^{\otimes i}, Q^{\otimes i} \) are treated as \( Y_0, Y_1 \). Let \( d_i \) denote the bound on the advantage of \( i \) copies, then we have that \( d_i = d \) and \( d_i \leq d_{i-1} + d - d_{i-1} \cdot d + (d_{i-1})^{k_{m,\varepsilon}} \). We can see by induction that \( d_i \leq 1 - (1 - d)^i + \varepsilon \) for
\[ d_i \leq d_{i-1} + d - d_{i-1} \cdot d + (d_{i-1})^{km, \varepsilon} = (1 - d)d_{i-1} + d + (d_{i-1})^{km, \varepsilon} \leq (1 - d)\left(1 - (1 - d)^{i-1} + \varepsilon\right) + d + (1 - (1 - d)^{i-1} + \varepsilon)^{km, \varepsilon} = 1 - (1 - d)^i + (1 - d)\varepsilon + d + (1 - (1 - d)^{i-1} + \varepsilon)^{km, \varepsilon} \leq 1 - (1 - d)^i + (1 - d)\varepsilon + (1 - (1 - d)^m + \varepsilon)^{km, \varepsilon} \leq 1 - (1 - d)^i + \varepsilon \]

where in the last inequality we used the choice of \( km, \varepsilon \). For the circuit size, we can easily see by induction on \( i \) that \( s_{i, \varepsilon} \geq (s - 1)/km, \varepsilon - 5i\ell - 1 \), as we have that \( s_{1, \varepsilon} = s \) and

\[
\begin{align*}
s_{i, \varepsilon} & \geq \min \left\{ s_{(i-1), \varepsilon} - \ell, \frac{s - 1}{km, \varepsilon} - 5(i - 1)\ell - 1 \right\} \\
& \geq \min \left\{ \frac{s - 1}{km, \varepsilon} - 5(i - 1)\ell - 1 - \ell, \frac{s - 1}{km, \varepsilon} - 5(i - 1)\ell - 1 \right\} \geq \frac{s - 1}{km, \varepsilon} - 5i\ell - 1.
\end{align*}
\]

\( \square \)

### 3.2 Tightness

This is somewhat folklore knowledge, that we explicitly state for the sake of completeness. We show that for every choice of \( d_X, d_Y, s_X, s_Y, \ell_X, \ell_Y \) there exist two pairs of distributions \( X_0, X_1 \) and \( Y_0, Y_1 \), such that \( X_0, X_1 \) are over \( \ell_X \) bits and cannot be distinguished with advantage better than \( d_X \) by size \( s_X \) circuits (resp. for \( Y_0, Y_1 \) with \( \ell_Y, d_Y, s_Y \)), yet \( (X_0, Y_0) \) and \( (X_1, Y_1) \) can be distinguished with advantage \( d_X + d_Y - d_X \cdot d_Y \) using a size 1 circuit. For the n-fold case, we show that for every choice of \( d, s, \ell \) there exist distributions \( X, Y \) over \( \ell \) bits with distance at most \( d \) against \( s \)-sized circuits, such that \( X^\otimes k, Y^\otimes k \) can be distinguished with advantage \( 1 - (1 - d)^k \) using a circuit of size \( 2k - 1 \). We will use statistical distance in these examples, noting that the statistical distance between distributions is equal to the maximal advantage of unbounded adversaries distinguishing between them, and that the statistical distance from a constant variable is equal to the probability to differ from it.

For the isolated case, we let \( X_0 \equiv 0^{\ell_X}, X_1 := B^{\ell_X}(d_X), Y_0 \equiv 0^{\ell_Y}, Y_1 := B^{\ell_Y}(d_Y) \). We have that size \( s_X \) circuits can distinguish between \( X_0, X_1 \) with advantage at most \( d_X \) (resp. for \( Y_0, Y_1 \) with \( s_Y, d_Y \)) as this is the statistical distance between them. Also, it is easy to verify that the simple size 1 circuit which given \( (x, y) \) computes \( x[1] \lor y[1] \) distinguishes between \( (X_0, Y_0) \) and \( (X_1, Y_1) \) with advantage \( 1 - (1 - d_X)(1 - d_Y) = d_X + d_Y - d_X \cdot d_Y \).

For the n-fold case, let \( X \equiv 0^\ell, Y := B^\ell(d) \), then size \( s \) circuits can distinguish \( X \) from \( Y \) with advantage at most \( d \). Yet the circuit of size \( 2k - 1 \) which given \( (z_1, \ldots, z_k) \) computes \( \lor_i z_i[1] \) (using a full binary tree of OR gates) distinguishes between \( X^\otimes k \) and \( Y^\otimes k \) with advantage \( 1 - (1 - d)^k \).

### 4 The Uniform Variant

We used non-uniformity two times in the proof of Theorem 3.1. The second time, which is easier to deal with, is in the circuit size analysis where we hard-coded the best samples of
y_0, y_1 to each iteration of A_i. Instead, in the uniform version, we will use uniform samplers of Y_0, Y_1. The first use of non-uniformity was when we assumed that C(x, \cdot) is at most a d_Y-distinguisher between Y_0 and Y_1, for every fixed x. More specifically, we used this to get that

$$\text{adv}_C^+ \left( (X_1, Y_{A(X_1)}), (X_1, Y_1) \right) \leq d_Y \cdot \Pr \left[ A(X_1) = 0 \right].$$

For the uniform case, we will relax this condition to x not being easy to hard-code, in the following sense:

$$\Pr_{x \leftarrow X_{1/2}} \left[ \text{adv}_{C(x, \cdot)}^+ (Y_0, Y_1) > d_Y + \varepsilon_k \right] \leq \varepsilon_k$$

where X_{1/2} is given by b \leftarrow \{0, 1\}, x \leftarrow X_b. If this condition doesn’t hold then we can efficiently compute a good x, except for negligible probability, assuming that efficient uniform samplers for X_0, X_1, Y_0, Y_1 exist. Otherwise, we will see that

$$\text{adv}_C^+ \left( (X_1, Y_{A(X_1)}), (X_1, Y_1) \right) \leq d_Y \cdot \Pr \left[ A(X_1) = 0 \right] + 3\varepsilon_k$$

and so almost the same argument from the non-uniform case works, except that now we lose another small additive term. Let us state and prove this more formally:

**Lemma 4.1.** Let X_0 = \{X_{0,n}\}, X_1 = \{X_{1,n}\}, Y_0 = \{Y_{0,n}\}, Y_1 = \{Y_{1,n}\} be ensembles of efficiently samplable distributions, and d_X(n), d_Y(n) be efficiently computable functions between 0 and 1. Then, for every k \in \mathbb{N} and time t(n) Turing machine M distinguishing (X_0, Y_0) from (X_1, Y_1) infinitely often with advantage at least (d_X + d_Y - d_X \cdot d_Y + 7\varepsilon_k) for

$$
\varepsilon_k := \frac{(d_Y)^k \cdot d_X (1 - d_Y)}{1 - (d_Y)^k} \leq (d_Y)^k,
$$

we have that either M efficiently yields a distinguisher for Y_0, Y_1 through a hard-coding of x, in the sense that for infinitely many n’s

$$\Pr_{x \leftarrow X_{1/2}} \left[ \text{adv}_{M(1^n, x, \cdot)}^+ (Y_0, Y_1) > d_Y + \varepsilon_k \right] > \varepsilon_k,$$

or there exists a time t \cdot \text{poly}(nk) infinitely often distinguisher between X_0, X_1 with advantage at least d_X.

**Proof.** For the sake of notational ease, we will drop the asymptotic notation and replace M(1^n) with C. Assume that for all but finitely many n’s,

$$\Pr_{x \leftarrow X_{1/2}} \left[ \text{adv}_{C(x, \cdot)}^+ (Y_0, Y_1) > d_Y + \varepsilon_k \right] \leq \varepsilon_k.$$

Then, for every candidate distinguisher A between X_0 and X_1, for all but finitely many n’s, we have that

$$\text{adv}_C^+ \left( (X_1, Y_{A(X_1)}), (X_1, Y_1) \right) \leq d_Y \cdot \Pr \left[ A(X_1) = 0 \right] + 3\varepsilon_k$$

$$\text{adv}_C^+ \left( (X_0, Y_0), (X_0, Y_{A(X_0)}) \right) \leq d_Y \cdot \Pr \left[ A(X_0) = 1 \right] + 3\varepsilon_k$$
where \(x, y \leftarrow X_1, Y_{A(X_1)}\) is resulted by \(x \leftarrow X_1, b \leftarrow A(x)\), \(y \leftarrow Y_b\). To see this, we first note that

\[
\varepsilon_k \geq \Pr_{x \leftarrow X_{1/2}} \left[ \text{adv}_c^+(Y_0, Y_1) > d_Y + \varepsilon_k \right] \\
\geq \frac{1}{2} \Pr[A(X_1) = 0] \Pr_{x \leftarrow X_{1/2}, A(X_1) = 0} \left[ \text{adv}_c^+(Y_0, Y_1) > d_Y + \varepsilon_k \right]
\]

which implies that

\[
\mathbb{E}_{x \leftarrow X_1 | A(X_1) = 0} \left[ \text{adv}_c^+(Y_0, Y_1) \right] \leq d_Y + \varepsilon_k + \frac{2\varepsilon_k}{\Pr[A(X_1) = 0]} \leq d_Y + \frac{3\varepsilon_k}{\Pr[A(X_1) = 0]}.
\]

Plugging it into the last inequality in the following, we get

\[
\text{adv}_c^+\left((X_1, Y_{A(X_1)}), (X_1, Y_1)\right) = \mathbb{E}\left[C(X_1, Y_1) - C(X_1, Y_{A(X_1)})\right] = \\
= \mathbb{E}\left[C(X_1, Y_1) - C(X_1, Y_0)\right] A(X_1) = 0 \cdot \Pr[A(X_1) = 0] + \\
+ \mathbb{E}\left[C(X_1, Y_1) - C(X_1, Y_1)\right] A(X_1) = 1 \cdot \Pr[A(X_1) = 1] = \\
= \mathbb{E}_{x \leftarrow X_1 | A(X_1) = 0} \left[C(x, Y_1) - C(x, Y_0)\right] \cdot \Pr[A(X_1) = 0] = \\
= \mathbb{E}_{x \leftarrow X_1 | A(X_1) = 0} \left[ \text{adv}_c^+(Y_0, Y_1) \right] \cdot \Pr[A(X_1) = 0] \leq d_Y \cdot \Pr[A(X_1) = 0] + 3\varepsilon_k
\]

and use a symmetric argument for the second upper bound. Using that (in general)

\[
\sum_{i \in [n]} \text{adv}_c^+(D_i, D_{i+1}) = \text{adv}_c^+(D_1, D_{n+1})
\]

we conclude that

\[
\text{adv}_c^+\left((X_0, Y_0), (X_1, Y_1)\right) = \text{adv}_c^+\left((X_0, Y_0), (X_0, Y_{A(X_0)})\right) + \\
+ \text{adv}_c^+\left((X_0, Y_{A(X_0)}), (X_1, Y_{A(X_1)})\right) + \text{adv}_c^+\left((X_1, Y_{A(X_1)}), (X_1, Y_1)\right)
\]

and thus

\[
\text{adv}_c^+\left((X_0, Y_{A(X_0)}), (X_1, Y_{A(X_1)})\right) = \text{adv}_c^+\left((X_0, Y_0), (X_1, Y_1)\right) - \\
- \text{adv}_c^+\left((X_1, Y_{A(X_1)}), (X_1, Y_1)\right) - \text{adv}_c^+\left((X_0, Y_0), (X_0, Y_{A(X_0)})\right) > \\
> (d_X + d_Y - d_X \cdot d_Y + 7\varepsilon_k) - (d_Y \cdot \Pr[A(X_1) = 0] + 3\varepsilon_k) - (d_Y \cdot \Pr[A(X_0) = 1] + 3\varepsilon_k) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y (1 - \Pr[A(X_1) = 0] - \Pr[A(X_0) = 1]) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y (\Pr[A(X_1) = 1] - \Pr[A(X_0) = 1]) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y (\mathbb{E}[A(X_1)] - \mathbb{E}[A(X_0)]) = \\
= (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y \cdot \text{adv}_A^+(X_0, X_1).
\]

In other words, we can build a new distinguisher \(A'\) for \(X_0, X_1\) by applying \(A\) to our input \(x\), sampling \(y \leftarrow Y_{A(x)}\) and feeding \((x, y)\) to \(C\), and have that

\[
\text{adv}_{A'}^+(X_0, X_1) > (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y \cdot \text{adv}_A^+(X_0, X_1).
\]
If we start from $A_0$ being the trivial distinguisher that always outputs 1 and keep repeating this process for $k$ steps, we get that

$$\text{adv}_{A_k}^+(X_0, X_1) > (d_X - d_X \cdot d_Y + \varepsilon_k) + d_Y \cdot \text{adv}_{A_{k-1}}^+(X_0, X_1) >$$
$$> (d_X - d_X \cdot d_Y + \varepsilon_k) + (d_Y - d_X \cdot d_Y + \varepsilon_k) + (d_Y)^2 \cdot \text{adv}_{A_{k-2}}^+(X_0, X_1) >$$
$$> \cdots > (d_X - d_X \cdot d_Y + \varepsilon_k) \sum_{i=0}^{k-1} (d_Y)^i + (d_Y)^k \cdot \text{adv}_{A_0}^+(X_0, X_1) =$$
$$= (d_X - d_X \cdot d_Y + \varepsilon_k) \sum_{i=0}^{k-1} (d_Y)^i = \frac{(d_X - d_X \cdot d_Y + \varepsilon_k) (1 - (d_Y)^k)}{1 - d_Y} =$$
$$= \frac{\left( d_X (1 - d_Y) + \frac{(d_Y)^k d_X (1 - d_Y)}{1 - (d_Y)^k} \right) (1 - (d_Y)^k)}{1 - d_Y} = \left( d_X + \frac{(d_Y)^k d_X}{1 - (d_Y)^k} \right) (1 - (d_Y)^k) =$$
$$= d_X (1 - (d_Y)^k) + (d_Y)^k \cdot d_X = d_X.$$

And so, we have concluded that $A_k$ distinguishes $X_0$ from $X_1$ with advantage better than $d_X$. In order to implement $A_k$ we need to run $C$, sample $Y_0, Y_1$ and use a multiplexer, for $k$ times, so we conclude that $\text{time}(A_k) = t \cdot \text{poly}(n, k)$.

**Remark 4.1.** In particular, we can use this lemma to show that if $X_0, X_1$ are $d_X$ ind. and $Y_0, Y_1$ are $d_Y$ ind. then $(X_0, Y_0)$ and $(X_1, Y_1)$ are $d_X + d_Y - d_X \cdot d_Y + 7 \varepsilon_k$ ind. for Turing machines with running time of

$$t = \min \{ t_X / \text{poly}(n, k), t_Y / \text{poly}(n, 1, \varepsilon_k) \},$$

which may be good enough for a constant number of uses, but does not work well beyond that, as every use costs us a division of the time bound by a polynomial. This is why we cannot prove the $n$-fold case immediately by repeatedly applying Lemma 4.1. The key idea is that we do not need to keep resampling and testing over and over again, but instead, once we find a good enough $x$ in the $i$'th coordinate, we fix it for the rest of the process, or if the hard-coding of the $i$'th coordinate does not succeed, the above lemma states we can distinguish there.

**Theorem 4.1.** Let $X = \{X_n\}, Y = \{Y_n\}$ be ensembles of efficiently samplable distributions that are $d(n)$ indistinguishable for time $t(n)$ Turing machines. Then, for every $m = m(n)$, we have that $X^{\otimes m}$ and $Y^{\otimes m}$ are $(1 - (1 - d)^m + 7m \varepsilon)$ indistinguishable for time $t_{m, \varepsilon}$ Turing machines, where

$$t_{m, \varepsilon} = t / \text{poly}(n, m, k_{m, \varepsilon}, 1/\varepsilon), \quad k_{m, \varepsilon} = \left\lceil \frac{\log(\varepsilon)}{\log(1 - (1 - d)^m + 7m \varepsilon)} \right\rceil \leq \left\lceil \frac{\log(1/\varepsilon)}{(1 - d)^m - 7m \varepsilon} \right\rceil.$$

**Proof.** For $i = 0, 1, \ldots, m - 1$, we try to hard-code the $m - i$'th coordinate using $\text{poly}(n, 1/\varepsilon)$ samples, and getting a distinguisher for $X^{\otimes m-i}, Y^{\otimes m-i}$ with advantage of at least $1 - (1 - d)^{m-i} + 7(m - i) \varepsilon$ except for negligible probability (the probability that the estimate was good but not truthful to the expectation) until for some $i$ we fail to find a good value to hard-code (if we reached $i = m - 1$ and succeeded then we are done). Once we fail, we apply
We are given (except for negligible probability) good samples for the coordinates in $m$ then fix this good $X$ such circuit failed, then one can build a distinguisher for them, and we are done.

Let us be more explicit about how we sample and hard-code the $m - i$th coordinate: We are given (except for negligible probability) good samples for the coordinates in $m - i + 1, \ldots, m$ and hard-code them into $A$, getting a $1 - (1 - d)^{m-i} + 7(m - i)\varepsilon$ distinguisher for $X^{\otimes m-i}, Y^{\otimes m-i}$, which we view as the product of $X^{\otimes m-i}, Y^{\otimes m-i}$ with $X, Y$. We first note that our choice of $k$ guarantees that $\varepsilon_k \leq \varepsilon$ for all $1 - (1 - d)^{m-i} + 7(m - i)\varepsilon$. We start by trying to work under the “hard-coding” assumption that

$$\Pr_{z \leftarrow X/Y} \left[ \text{adv}^+_A(X^{\otimes m-i-1}, Y^{\otimes m-i-1}) > 1 - (1 - d)^{m-i-1} + 7(m - i - 1)\varepsilon + \varepsilon \right] > \varepsilon$$

and generate a distinguisher for $X^{\otimes m-i-1}, Y^{\otimes m-i-1}$ as follows: Keep sampling $z \leftarrow X/Y$ and estimating $\text{adv}^+_A(X^{\otimes m-i-1}, Y^{\otimes m-i-1})$ using $r$ samples from $X^{\otimes m-i-1}/Y^{\otimes m-i-1}$, until we succeed in finding $z$ with an estimate of at least $1 - (1 - d)^{m-i-1} + 7(m - i - 1)\varepsilon + 0.5\varepsilon$, then fix this good $z$ in this coordinate and move forward, or stop after $q$ tries if no such $z$ has been found. Using Hoeffding’s inequality, for every $z$, the probability that the estimate’s error is greater than $\varepsilon/2$ is at most $2e^{-r(\varepsilon/2)^2/2}$. If all estimates were $\varepsilon/2$ accurate and a good $z$ has been drawn, the process succeeds in finding a $z$ with advantage of at least $1 - (1 - d)^{m-i-1} + 7(m - i - 1)\varepsilon$ and we can move on, so our probability to fail at that, under the above assumption, is at most

$$q \cdot 2e^{-r\varepsilon^2/32} + (1 - \varepsilon)^q \leq 2e^{\log(q/2) - r\varepsilon^2/32} + e^{-q\varepsilon} \leq \text{neg}(n)$$

by choosing, say,

$$q = n/\varepsilon = \text{poly}(n, 1/\varepsilon), \quad r = 64n/\varepsilon^3 > (\log(q/2) + n) \cdot 32/\varepsilon^2 = \text{poly}(n, 1/\varepsilon).$$

Hence paying with a time complexity of $t_{m,\varepsilon} \cdot \text{poly}(n, 1/\varepsilon)$ for every coordinate.

If we could not find a good $z$, we use Lemma 4.1: If we can distinguish $X^{\otimes m-i}, Y^{\otimes m-i}$ with advantage

$$(1 - d) \left( 1 - (1 - d)^{m-i-1} + 7(m - i - 1)\varepsilon \right) + d + 7\varepsilon =$$

$$= 1 - (1 - d)^{m-i} + (1 - d)7(m - i - 1)\varepsilon + 7\varepsilon \leq$$

$$\leq 1 - (1 - d)^{m-i} + 7(m - i)\varepsilon \leq \text{adv}^+_A(X^{\otimes m-i}, Y^{\otimes m-i})$$

and the assumption about finding a good $z$ to hard-code for $X^{\otimes m-i-1}, Y^{\otimes m-i-1}$ does not hold, then we can build a $d$-distinguisher for $X, Y$ in time $t_{m,\varepsilon} \cdot \text{poly}(n, k)$. The probability that at some point in the process we failed to hard-code a good $z$ at the $m - i$th coordinate even though the assumption held is $m(n) \cdot \text{neg}(n) = \text{neg}(n)$. \qed

We remark this proof is easily generalized to the case where not all pairs in the product are identical, that is, for $\otimes X_i$ and $\otimes Y_i$, with a distance bound of $(1 - \prod_i (1 - d_i) + 7m\varepsilon)$.  

9
5 Applications

As an application, we consider the amplification of weak oblivious transfer protocols. We briefly explain how our bounds, paired with Yao’s XOR lemma, yield a natural generalization in the computational setting of the amplification process presented in [DKS99, Subsection 4.3]. We stress that it is already known that the same amplification process also works computationally [Wul07], yet we find the following approach more straightforward. For the sake of simplicity, we consider the amplification of error-less $(p, q)$-weak semi-honest 1-2 OT: The receiver with bit $c$ is trying to learn $b_c$, where $b_0, b_1$ is the database of the sender. We say the protocol is $(p, q)$ weak if the view of the sender when $c = 0$ is $p$-indistinguishable from its view when $c = 1$ (equivalently, $c$ is at most $p$-correlated to the view of the sender), and the view of the receiver when $b_c = 0$ is $q$-indistinguishable from its view when $b_c = 1$. We have an operation called S-Reduce that amplifies indistinguishability against the sender but worsens indistinguishability against the receiver, and an operation called R-Reduce that amplifies indistinguishability against the receiver but worsens indistinguishability against the sender. Our goal is to use them repeatedly one after the other in order to amplify both parameters. It is already shown in [DKS99, Lemma 4] exactly how this is done, so our focus will be on showing that almost the same analysis of the S-Reduce and R-Reduce operations also holds computationally. Let us start with the security of the receiver: In S-Reduce where $c = \bigoplus_i c_i$ we can use Yao’s XOR Lemma to show that $p$ is reduced to $p^k + \varepsilon$, and in R-Reduce where $c_i = c$ we use our bound to show that $p$ is increased to at most $1 - (1 - p)^k + \varepsilon$. For the sender, roughly speaking: In S-Reduce, we have a product of $k$ execution pairs that are $q$-indistinguishable each (whether "the other bit" is 0 or 1), and we can use our bound to get that they are $1 - (1 - q)^k + \varepsilon$ indistinguishable. In R-Reduce, "the other bit" is equal to the XOR of $k$ bits that are at most $q$-correlated to the execution, independently, so we use the XOR lemma to conclude that "the other bit" is at most $q^k + \varepsilon$ correlated to the execution. We conclude that essentially, the same analysis from the information theoretic setting works, up to an additive $\varepsilon$ for each use. Let $p(n)$ be a bound on the total number of calls to the original protocol in the information-theoretic transformation, then all advantages throughout the process are $1/p(n)$-bounded away from 1, otherwise we wouldn’t be able to reduce them to negligible. If we choose $\varepsilon' = \varepsilon/p(n)$ then for every advantage $d$ through the process we have $d + \varepsilon' \leq \varepsilon + (1 - \varepsilon)d$, so we can imagine, for a simple analysis, as if every call to either S-Reduce or R-Reduce incurs a chance of $\varepsilon$ at failing and revealing everything, and otherwise works exactly like the information-theoretic world. Since the number of calls is polynomial, the total probability of failing is at most poly($n$) · $\varepsilon$ and we can make it as (polynomially) small as we want. There is one issue however - the running time. In the information-theoretic process we make log($n$) calls to S-Reduce and R-Reduce, and each such call, when using Yao’s XOR Lemma or the bounds in this paper, decreases the bound on the running time by a division in a polynomial. Therefore, we need the assumption that our weak OT is secure against $n^{O(\log n)}$ adversaries. We believe this stronger requirement may be removed by a more careful analysis (perhaps we again waste too much time on resampling and testing unnecessary values, see Remark 4.1), but without a proof.
References


