

## Lecture Space-Bounded Derandomization

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## 1 Space-Bounded Derandomization

We now discuss derandomization of space-bounded algorithms. Here non-trivial results can be shown *without making any unproven assumptions*, in contrast to what is currently known for derandomizing time-bounded algorithms. We show first that<sup>1</sup>  $\mathcal{BPL} \subseteq \text{SPACE}(\log^2 n)$  and then improve the analysis and show that<sup>2</sup>  $\mathcal{BPL} \subseteq \text{TIME}(\text{poly}(n), \log^2 n) \subseteq \mathcal{SC}$ . (Note: we already know

$$\mathcal{RL} \subseteq \mathcal{NL} \subseteq \text{SPACE}(\log^2 n)$$

but this does not by itself imply  $\mathcal{BPL} \subseteq \text{SPACE}(\log^2 n)$ .)

With regard to the first result, we actually prove something more general:

**Theorem 1** *Any randomized algorithm (with two-sided error) that uses space  $S = \Omega(\log n)$  and  $R$  random bits can be converted to one that uses space  $\mathcal{O}(S \log R)$  and  $\mathcal{O}(S \log R)$  random bits.*

Since any algorithm using space  $S$  uses time at most  $2^S$  (by our convention regarding probabilistic machines) and hence at most this many random bits, the following is an immediate corollary:

**Corollary 2** *For  $S = \Omega(\log n)$  it holds that  $\text{BPPSPACE}(S) \subseteq \text{SPACE}(S^2)$ .*

**Proof** Let  $L \in \text{BPPSPACE}(S)$ . Theorem 1 shows that  $L$  can be decided by a probabilistic machine with two-sided error using  $\mathcal{O}(S^2)$  space and  $\mathcal{O}(S^2)$  random bits. Enumerating over all random bits and taking majority, we obtain a deterministic algorithm that uses  $\mathcal{O}(S^2)$  space. ■

## 2 $\mathcal{BPL} \subseteq \text{SPACE}(\log^2 n)$

We now prove Theorem 1. Let  $M$  be a probabilistic machine running in space  $S$  (and time  $2^S$ ), using  $R$  random bits, and deciding a language  $L$  with two-sided error. (Note that  $S, R$  are functions of the input length  $n$ , and the theorem requires  $S = \Omega(\log n)$ .) We will assume without loss of generality that  $M$  always uses exactly  $R$  random bits on all inputs. Fixing an input  $x$  and letting  $\ell$  be some parameter, we will view the computation of  $M_x$  as a random walk on a multi-graph in the following way: the nodes of the graph correspond to all  $N \stackrel{\text{def}}{=} 2^{\mathcal{O}(S)}$  possible configurations of  $M_x$ , and there is an edge from  $a$  to  $b$  labeled by the string  $r \in \{0, 1\}^\ell$  if and only if  $M_x$  moves from configuration  $a$  to configuration  $b$  after reading  $r$  as its next  $\ell$  random bits. Computation of  $M_x$  is then equivalent to a random walk of length  $R/\ell$  on this graph, beginning from the node corresponding to the initial configuration of  $M_x$ . If  $x \in L$  then the probability that this random

<sup>1</sup> $\mathcal{BPL}$  is the two-sided-error version of  $\mathcal{RL}$ .

<sup>2</sup> $\mathcal{SC}$  stands for “Steve’s class”, and captures computation that *simultaneously* uses polynomial time and polylogarithmic space.

walk ends up in an accepting state is at least  $2/3$ , while if  $x \notin L$  then the probability that this random walk ends up in an accepting state is at most  $1/3$ .

It will be convenient to represent this process using an  $N \times N$  transition matrix  $Q_x$ , where the entry in column  $i$ , row  $j$  is the probability that  $M_x$  moves from configuration  $i$  to configuration  $j$  after reading  $\ell$  random bits. Vectors of length  $N$  whose entries are non-negative and sum to 1 correspond to probability distributions over the configurations of  $M_x$  in the natural way. If we let  $\mathbf{s}$  denote the probability distribution that places probability 1 on the initial configuration of  $M_x$  (and 0 elsewhere), then  $Q_x^{R/\ell} \cdot \mathbf{s}$  corresponds to the probability distribution over the final configuration of  $M_x$ ; thus:

$$\begin{aligned} x \in L &\Rightarrow \sum_{i \in \text{accept}} \left( Q_x^{R/\ell} \cdot \mathbf{s} \right)_i \geq 3/4 \\ x \notin L &\Rightarrow \sum_{i \in \text{accept}} \left( Q_x^{R/\ell} \cdot \mathbf{s} \right)_i \leq 1/4. \end{aligned}$$

The statistical difference between two vectors/probability distributions  $\mathbf{s}, \mathbf{s}'$  is

$$\text{SD}(\mathbf{s}, \mathbf{s}') \stackrel{\text{def}}{=} \frac{1}{2} \cdot \|\mathbf{s} - \mathbf{s}'\| = \frac{1}{2} \cdot \sum_i |\mathbf{s}_i - \mathbf{s}'_i|.$$

If  $Q, Q'$  are two transition matrices — meaning that all entries are non-negative, and the entries in each column sum to 1 — then we abuse notation and define

$$\text{SD}(Q, Q') \stackrel{\text{def}}{=} \max_{\mathbf{s}} \{\text{SD}(Q\mathbf{s}, Q'\mathbf{s})\},$$

where the maximum is taken over all  $\mathbf{s}$  that correspond to probability distributions. Note that if  $Q, Q'$  are  $N \times N$  transition matrices and  $\max_{i,j} \{|Q_{i,j} - Q'_{i,j}|\} \leq \varepsilon$ , then  $\text{SD}(Q, Q') \leq N\varepsilon/2$ .

## 2.1 A Useful Lemma

The pseudorandom generator we construct will use a family  $H$  of pairwise-independent functions as a building block.

**Definition 1**  $H = \{h_k : \{0, 1\}^\ell \rightarrow \{0, 1\}^\ell\}$  is a family of pairwise-independent functions if for all distinct  $x_1, x_2 \in \{0, 1\}^\ell$  and any  $y_1, y_2 \in \{0, 1\}^\ell$  we have:

$$\Pr_{h \in H} [h(x_1) = y_1 \wedge h(x_2) = y_2] = 2^{-2\ell}.$$

It is easy to construct a pairwise-independent family  $H$  whose functions map  $\ell$ -bit strings to  $\ell$ -bit strings and such that (1)  $|H| = 2^{2\ell}$  (and so choosing a random member of  $H$  is equivalent to choosing a random  $2\ell$ -bit string) and (2) functions in  $H$  can be evaluated in  $\mathcal{O}(\ell)$  space.

For  $S \subseteq \{0, 1\}^\ell$ , define  $\rho(S) \stackrel{\text{def}}{=} |S|/2^\ell$ . We define a useful property and then show that a function chosen from a pairwise-independent family satisfies the property with high probability.

**Definition 2** Let  $A, B \subseteq \{0, 1\}^\ell$ ,  $h : \{0, 1\}^\ell \rightarrow \{0, 1\}^\ell$ , and  $\varepsilon > 0$ . We say  $h$  is  $(\varepsilon, A, B)$ -good if:

$$\left| \Pr_{x \in \{0, 1\}^\ell} \left[ x \in A \wedge h(x) \in B \right] - \rho(A) \cdot \rho(B) \right| \leq \varepsilon.$$

Note that this is equivalent to saying that  $h$  is  $(\varepsilon, A, B)$ -good if

$$\left| \Pr_{x \in A} [h(x) \in B] - \rho(B) \right| \leq \varepsilon / \rho(A).$$

**Lemma 3** *Let  $A, B \subseteq \{0, 1\}^\ell$ ,  $H$  be a family of pairwise-independent functions, and  $\varepsilon > 0$ . Then:*

$$\Pr_{h \in H} [h \text{ is not } (\varepsilon, A, B)\text{-good}] \leq \frac{\rho(A)\rho(B)}{2^\ell \varepsilon^2}.$$

**Proof** The proof is fairly straightforward. Consider the quantity

$$\begin{aligned} \mu &\stackrel{\text{def}}{=} \text{Exp}_{h \in H} \left[ \left( \rho(B) - \Pr_{x \in A} [h(x) \in B] \right)^2 \right] \\ &= \text{Exp}_{h \in H} \left[ \rho(B)^2 + \Pr_{x_1 \in A} [h(x_1) \in B] \cdot \Pr_{x_2 \in A} [h(x_2) \in B] - 2\rho(B) \cdot \Pr_{x_1 \in A} [h(x_1) \in B] \right] \\ &= \rho(B)^2 + \text{Exp}_{x_1, x_2 \in A; h \in H} [\delta_{h(x_1) \in B} \cdot \delta_{h(x_2) \in B} - 2\rho(B) \cdot \delta_{h(x_1) \in B}], \end{aligned}$$

where  $\delta_{h(x) \in B}$  is an indicator random variable which is equal to 1 if  $h(x) \in B$  and 0 otherwise. Since  $H$  is pairwise independent, it follows that:

- For any  $x_1$  we have  $\text{Exp}_{h \in H} [\delta_{h(x_1) \in B}] = \Pr_{h \in H} [h(x_1) \in B] = \rho(B)$ .
- For any  $x_1 = x_2$  we have  $\text{Exp}_{h \in H} [\delta_{h(x_1) \in B} \cdot \delta_{h(x_2) \in B}] = \text{Exp}_{h \in H} [\delta_{h(x_1) \in B}] = \rho(B)$ .
- For any  $x_1 \neq x_2$  we have  $\text{Exp}_{h \in H} [\delta_{h(x_1) \in B} \cdot \delta_{h(x_2) \in B}] = \Pr_{h \in H} [h(x_1) \in B \wedge h(x_2) \in B] = \rho(B)^2$ .

Using the above, we obtain

$$\mu = \rho(B)^2 + \frac{\rho(B)}{|A|} + \frac{\rho(B)^2(|A| - 1)}{|A|} - 2\rho(B)^2 = \frac{\rho(B) - \rho(B)^2}{|A|} = \frac{\rho(B)(1 - \rho(B))}{|A|}.$$

Using Markov's inequality,

$$\begin{aligned} \Pr_{h \in H} [h \text{ is not } (\varepsilon, A, B)\text{-good}] &= \Pr_{h \in H} \left[ \left( \Pr_{x \in A} [h(x) \in B] - \rho(B) \right)^2 > (\varepsilon / \rho(A))^2 \right] \\ &\leq \frac{\mu \cdot \rho(A)^2}{\varepsilon^2} = \frac{\rho(B)(1 - \rho(B))\rho(A)}{2^\ell \varepsilon^2} \leq \frac{\rho(B)\rho(A)}{2^\ell \varepsilon^2}. \end{aligned}$$

■

## 2.2 The Pseudorandom Generator and Its Analysis

### 2.2.1 The Basic Step

We first show how to reduce the number of random bits by roughly half. Let  $H$  denote a pairwise-independent family of functions, and fix an input  $x$ . Let  $Q$  denote the transition matrix corresponding to transitions in  $M_x$  after reading  $\ell$  random bits; that is, the  $(i, j)$ th entry of  $Q$  is the

probability that  $M_x$ , starting in configuration  $i$ , moves to configuration  $j$  after reading  $\ell$  random bits. So  $Q^2$  is a transition matrix denoting the probability that  $M_x$ , starting in configuration  $i$ , moves to configuration  $j$  after reading  $2\ell$  random bits. Fixing  $h \in H$ , let  $Q_h$  be a transition matrix where the  $(i, j)$ th entry in  $Q_h$  is the probability that  $M_x$ , starting in configuration  $i$ , moves to configuration  $j$  after reading the  $2\ell$  “random bits”  $r \parallel h(r)$  (where  $r \in \{0, 1\}^\ell$  is chosen uniformly at random). Put differently,  $Q^2$  corresponds to taking two uniform and independent steps of a random walk, whereas  $Q_h$  corresponds to taking two steps of a random walk where the first step (given by  $r$ ) is random and the second step (namely,  $h(r)$ ) is a deterministic function of the first. We now show that these two transition matrices are “very close”. Specifically:

**Definition 3** Let  $Q, Q_h, \ell$  be as defined above, and  $\varepsilon \geq 0$ . We say  $h \in H$  is  $\varepsilon$ -good for  $Q$  if

$$\text{SD}(Q_h, Q^2) \leq \varepsilon/2.$$

**Lemma 4** Let  $H$  be a pairwise-independent function family, and let  $Q$  be an  $N \times N$  transition matrix where transitions correspond to reading  $\ell$  random bits. For any  $\varepsilon > 0$  we have:

$$\Pr_{h \in H} [h \text{ is not } \varepsilon\text{-good for } Q] \leq \frac{N^6}{\varepsilon^2 2^{2\ell}}.$$

**Proof** For  $i, j \in [N]$  (corresponding to configurations in  $M_x$ ), define

$$B_{i,j} \stackrel{\text{def}}{=} \{x \in \{0, 1\}^\ell \mid x \text{ takes } Q \text{ from } i \text{ to } j\}.$$

For fixed  $i, j, k$ , we know from Lemma 3 that the probability that  $h$  is not  $(\varepsilon/N^2, B_{i,j}, B_{j,k})$ -good is at most  $N^4 \rho(B_{i,j}) / \varepsilon^2 2^\ell$ . Applying a union bound over all  $N^3$  triples  $i, j, k \in [N]$ , and noting that for any  $i$  we have  $\sum_j \rho(B_{i,j}) = 1$ , we have that  $h$  is  $(\varepsilon/N^2, B_{i,j}, B_{j,k})$ -good for all  $i, j, k$  except with probability at most  $N^6 / \varepsilon^2 2^\ell$ .

We show that whenever  $h$  is  $(\varepsilon/N^2, B_{i,j}, B_{j,k})$ -good for all  $i, j, k$ , then  $h$  is  $\varepsilon$ -good for  $Q$ . Consider the  $(i, k)$ th entry in  $Q_h$ ; this is given by:  $\sum_{j \in [N]} \Pr[r \in B_{i,j} \wedge h(r) \in B_{j,k}]$ . On the other hand, the  $(i, k)$ th entry in  $Q^2$  is:  $\sum_{j \in [N]} \rho(B_{i,j}) \cdot \rho(B_{j,k})$ . Since  $h$  is  $(\varepsilon/N^2, B_{i,j}, B_{j,k})$ -good for every  $i, j, k$ , the absolute value of their difference is

$$\begin{aligned} & \left| \sum_{j \in [N]} \left( \Pr[r \in B_{i,j} \wedge h(r) \in B_{j,k}] - \rho(B_{i,j}) \cdot \rho(B_{j,k}) \right) \right| \\ & \leq \sum_{j \in [N]} \left| \Pr[r \in B_{i,j} \wedge h(r) \in B_{j,k}] - \rho(B_{i,j}) \cdot \rho(B_{j,k}) \right| \\ & \leq \sum_{j \in [N]} \varepsilon / N^2 = \varepsilon / N. \end{aligned}$$

It follows that  $\text{SD}(Q_h, Q^2) \leq \varepsilon/2$  as desired. ■

The lemma above gives us a pseudorandom generator that reduces the required randomness by (roughly) half. Specifically, define a pseudorandom generator  $G_1 : \{0, 1\}^{2\ell + R/2} \rightarrow \{0, 1\}^R$  via:

$$G_1(r_1, \dots, r_{R/2\ell}; h) = r_1 \parallel h(r_1) \parallel \dots \parallel r_{R/2\ell} \parallel h(r_{R/2\ell}), \quad (1)$$

where  $h \in H$  (so  $|h| = 2\ell$ ) and  $r_i \in \{0, 1\}^\ell$ . Assume  $h$  is  $\varepsilon$ -good for  $Q$ . Running  $M_x$  using the output of  $G_1(h, \dots)$  as the “random tape” generates the probability distribution

$$\overbrace{Q_h \cdots Q_h}^{R/2\ell} \cdot \mathbf{s}$$

for the final configuration, where  $\mathbf{s}$  denotes the initial configuration of  $M_x$  (i.e.,  $\mathbf{s}$  is the probability distribution that places probability 1 on the initial configuration of  $M_x$ , and 0 elsewhere). Running  $M_x$  on a truly random tape generates the probability distribution

$$\overbrace{Q^2 \cdots Q^2}^{R/2\ell} \cdot \mathbf{s}$$

for the final configuration. Letting  $k = R/2\ell$  we have

$$\begin{aligned} 2 \cdot \text{SD} \left( \overbrace{Q_h \cdots Q_h}^k \cdot \mathbf{s}, \overbrace{Q^2 \cdots Q^2}^k \cdot \mathbf{s} \right) &= \left\| \left( \overbrace{Q_h \cdots Q_h}^k - \overbrace{Q^2 \cdots Q^2}^k \right) \cdot \mathbf{s} \right\| \\ &= \left\| \sum_{i=0}^{k-1} \left( \overbrace{Q_h \cdots Q_h}^{k-i} \overbrace{Q^2 \cdots Q^2}^i - \overbrace{Q_h \cdots Q_h}^{k-i-1} \overbrace{Q^2 \cdots Q^2}^{i+1} \right) \cdot \mathbf{s} \right\| \\ &\leq \sum_{i=0}^{k-1} \left\| \left( \overbrace{Q_h \cdots Q_h}^{k-i} \overbrace{Q^2 \cdots Q^2}^i - \overbrace{Q_h \cdots Q_h}^{k-i-1} \overbrace{Q^2 \cdots Q^2}^{i+1} \right) \cdot \mathbf{s} \right\| \\ &= \sum_{i=0}^{k-1} \left\| \overbrace{Q_h \cdots Q_h}^{k-i-1} \cdot (Q_h - Q^2) \cdot \overbrace{Q^2 \cdots Q^2}^i \cdot \mathbf{s} \right\| \\ &\leq k \cdot \varepsilon. \end{aligned}$$

This means that the behavior of  $M_x$  when run using the output of the pseudorandom generator is very close to the behavior of  $M_x$  when run using a truly random tape: in particular, if  $x \notin L$  then  $M_x$  in the former case accepts with probability at most

$$\Pr[\text{accepts} \wedge h \text{ is } \varepsilon\text{-good for } Q] + \Pr[h \text{ is not } \varepsilon\text{-good for } Q] \leq (1/4 + k\varepsilon/2) + N^6/\varepsilon^2 2^\ell;$$

similarly, if  $x \in L$  then  $M_x$  in the former case accepts with probability at least  $3/4 - k\varepsilon/2 - N^6/\varepsilon^2 2^\ell$ . Summarizing (and slightly generalizing):

**Corollary 5** *Let  $H$  be a pairwise-independent function family, let  $Q$  be an  $N \times N$  transition matrix where transitions correspond to reading  $\ell$  random bits, let  $k > 0$  be an integer, and let  $\varepsilon > 0$ . Then except with probability at most  $N^6/\varepsilon^2 2^\ell$  over choice of  $h \in H$  we have:*

$$\text{SD} \left( \overbrace{Q_h \cdots Q_h}^k, \overbrace{Q^2 \cdots Q^2}^k \right) \leq k\varepsilon/2.$$

### 2.2.2 Recursing

Fixing  $h_1 \in H$ , note that  $Q_{h_1}$  is a transition matrix and so we can apply Corollary 5 to it as well. Moreover, if  $Q$  uses  $R$  random bits then  $Q_{h_1}$  uses  $R/2$  random bits (treating  $h_1$  as fixed). Continuing in this way for  $I \stackrel{\text{def}}{=} \log(R/2\ell) + 1 = \log(R/\ell)$  iterations, we obtain a transition matrix  $Q_{h_1, \dots, h_I}$ . Say *all  $h_i$  are  $\varepsilon$ -good* if  $h_1$  is  $\varepsilon$ -good for  $Q$ , and for each  $i > 1$  it holds that  $h_i$  is  $\varepsilon$ -good for  $Q_{h_1, \dots, h_{i-1}}$ . By Corollary 5 we have:

- All  $h_i$  are  $\varepsilon$ -good except with probability at most  $N^6 I / \varepsilon^2 2^\ell$ .
- If all  $h_i$  are  $\varepsilon$ -good then

$$\text{SD}(Q_{h_1, \dots, h_I}, \overbrace{Q^2 \cdots Q^2}^{R/2\ell}) \leq \frac{\varepsilon}{2} \cdot \sum_{i=1}^I \frac{R}{2^i \ell} = \frac{\varepsilon}{2} \cdot \left( \frac{R}{\ell} - 1 \right).$$

Equivalently, we obtain a pseudorandom generator

$$G_I(r; h_1, \dots, h_I) \stackrel{\text{def}}{=} G_{I-1}(r; h_1, \dots, h_{I-1}) \| G_{I-1}(h_I(r); h_1, \dots, h_{I-1}),$$

where  $G_1$  is as in Equation (1).

### 2.2.3 Putting it All Together

We now easily obtain the desired derandomization. Recall  $N = 2^{\mathcal{O}(s)}$ . Set  $\varepsilon = 2^{-S}/10$ , and set  $\ell = \Theta(S)$  so that  $\frac{N^6 S}{\varepsilon^2 2^\ell} \leq 1/20$ . Then the number of random bits used (as input to  $G_I$  from the previous section) is  $\mathcal{O}(\ell \cdot \log(R/\ell) + \ell) = \mathcal{O}(S \log R)$  and the space used is bounded by that as well (using the fact that each  $h \in H$  can be evaluated using space  $\mathcal{O}(\ell) = \mathcal{O}(S)$ ). All  $h_i$  are good except with probability at most  $N^6 \log(R/\ell) / \varepsilon^2 2^\ell \leq N^6 S / \varepsilon^2 2^\ell \leq 1/20$ ; assuming all  $h_i$  are good, the statistical difference between an execution of the original algorithm and the algorithm run with a pseudorandom tape is bounded by  $2^{-S}/20 \cdot R \leq 1/20$ . Theorem 1 follows easily.

## 3 $BPL \subseteq SC$

A deterministic algorithm using space  $\mathcal{O}(\log^2 n)$  might potentially run for  $2^{\mathcal{O}(\log^2 n)}$  steps; in fact, as described, the algorithm from the proof of Corollary 2 uses this much time. For the particular pseudorandom generator we have described, however, it is possible to do better. The key observation is that instead of just choosing the  $h_1, \dots, h_I$  at random and simply hoping that they are all  $\varepsilon$ -good, we will instead deterministically search for  $h_1, \dots, h_I$  which *are* each  $\varepsilon$ -good. This can be done in polynomial time (when  $S = \mathcal{O}(\log n)$ ) because: (1) for a given transition matrix  $Q_{h_1, \dots, h_{i-1}}$  and candidate  $h_i$ , it is possible to determine in polynomial time and polylogarithmic space whether  $h_i$  is  $\varepsilon$ -good for  $Q_{h_1, \dots, h_{i-1}}$  (this relies on the fact that the number of configurations  $N$  is polynomial in  $n$ ); (2) there are only a polynomial number of possibilities for each  $h_i$  (since  $\ell = \Theta(S) = \mathcal{O}(\log n)$ ).

Once we have found the good  $\{h_i\}$ , we then cycle through all possible choices of the seed  $r \in \{0, 1\}^\ell$  and take majority (as before). Since there are a polynomial number of possible seeds, the algorithm as a whole runs in polynomial time.

(For completeness, we discuss the case of general  $S = \Omega(\log n)$  assuming  $R = 2^S$ . Checking whether a particular  $h_i$  is  $\varepsilon$ -good requires time  $2^{\mathcal{O}(S)}$ . There are  $2^{\mathcal{O}(S)}$  functions to search through at each stage, and  $\mathcal{O}(S)$  stages altogether. Finally, once we obtain the good  $\{h_i\}$  we must then enumerate through  $2^{\mathcal{O}(S)}$  seeds. The end result is that  $\text{BSPACE}(S) \subseteq \text{TIMESPC}(2^{\mathcal{O}(S)}, S^2)$ .)

## Bibliographic Notes

The results described here are due to [2, 3], both of which are very readable. See also [1, Lecture 16] for a slightly different presentation.

## References

- [1] O. Goldreich. Introduction to Complexity Theory (July 31, 1999).
- [2] N. Nisan. Pseudorandom Generators for Space-Bounded Computation. *STOC* '90.
- [3] N. Nisan.  $RL \subseteq SC$ . *Computational Complexity* 4: 1–11, 1994. (Preliminary version in *STOC* '92.)