

Lecture 23: Colorings with Low Discrepancy

Bansal's algorithm (2010). This computes the desired coloring through a sequence of semicolorings, where a semicoloring is an arbitrary mapping $c: V \rightarrow [-1, 1]$.

The algorithm starts with the semicoloring $x_0 = 0$. Then it produces a sequence

$x_0, x_1, x_2, \dots, x_l \in [-1, 1]^n$ of semicolorings. What l should be will be determined later.

The t -th step. x_t is obtained from x_{t-1} as follows.

Let $A_t = \{j \in V : (x_{t-1})_j \neq \pm 1\}$ is the set of coordinates that are still active in the t -th step. We will use semidefinite programming to compute a coloring of A_t by unit vectors (rather than ± 1).

More explicitly, we compute unit vectors $u_{t,j}$ where $j \in A_t$ so that

$$\left\| \sum_{j \in A_t \cap S_i} u_{t,j} \right\|^2 \leq D^2$$

for all i , with $D \geq 0$ as small as possible.

For convenience, we also set $u_{t,j} = 0$ for $j \notin A_t$.

Next, we generate a random vector r_t from the n -dimensional standard normal distribution, i.e., the coordinates of r_t are independent $N(0, 1)$ random variables.

Then we set $(\Delta_t)_j = \tau \cdot r_t^T u_{t,j}$ for $j = 1, \dots, n$.

Here τ is a small parameter.

We will update x_{t-1} to x_t as follows: (2)

- A tentative value of x_t is $\tilde{x}_t = x_{t-1} + \Delta_t$.
- But we need to truncate each coordinate to the interval $[-1, 1]$. So we update \tilde{x}_t as:

$$(x_t)_j = \begin{cases} +1 & \text{if } (\tilde{x}_t)_j \geq 1 \\ -1 & \text{if } (\tilde{x}_t)_j \leq -1 \\ (\tilde{x}_t)_j & \text{otherwise,} \end{cases}$$

for $j = 1, \dots, n$.

The length l of the sequence x_0, x_1, \dots, x_l of semicorings will be set to $\frac{C_1 \cdot \log n}{\tau^2}$ for a suitable constant C_1 .

As we will see in due time $\tau = \frac{1}{C_0 n \sqrt{\log n}}$ for another suitable constant C_0 .

This concludes the description of Bansal's algorithm.

Roughly speaking, why does the algorithm work? The algorithm can be regarded as a random walk in the cube $[-1, 1]^n$. The projection of the random walk to a given coordinate axis behaves like a 1-dimensional random walk with increments having the normal distribution $\tau \cdot N(0, 1) = N(0, \tau^2)$. Such a walk will typically cross the boundary of the interval $[-1, 1]$ in about $\frac{1}{\tau^2}$ steps. After $\frac{\log n}{\tau^2}$ steps, it is very likely to have crossed the boundary. This means that x_l is very likely to be a ± 1 vector.

Regarding the imbalance of any S_i , it starts out

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at 0 under the semicoloring x_0 and in the t -th step, it is changed by $\sum_{j \in S_i} \tau \cdot r_t^T u_{t,j} = \tau r_t^T v_{t,i}$

where $v_{t,i} = \sum_{j \in S_i} u_{t,j}$. But the $u_{t,j}$ were selected with the goal of making $\|v_{t,i}\|$ small. So the imbalance of each S_i grows slowly during the algorithm.

We will now make these arguments formally.

Claim 1. The algorithm produces a coloring with probability $\geq 1 - 1/n$.

Proof. Let x_0, x_1, \dots, x_l be the sequence of semicolorings generated by the algorithm. Let $j \in \{1, \dots, n\}$ be a fixed index. We will call the sequence $(x_0)_j, (x_1)_j, \dots, (x_l)_j$ the j -th coordinate random walk.

We will say that the j -th coordinate random walk terminates if $(x_l)_j \in \{\pm 1\}$. To prove Claim 1, it is enough to show that, for every j , the probability that the j -th coordinate random walk does not terminate is $\leq 1/n^2$. Then w.p. $\geq 1 - \frac{1}{n}$, all of the coordinate random walks terminate (by the union bound).

To simplify notation, let us fix j and write $X_t = (x_t)_j - (x_{t-1})_j$, for $t = 1, \dots, l$.

We want to claim that X_1, X_2, \dots, X_l are independent random variables but this is not true.

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This is because once $(x_t)_j \in \{\pm 1\}$, this value stays fixed. Let us define $t_0 \leq l$ to be the last step of the j -th coordinate walk for which $(x_{t_0})_j \in (-1, 1)$.

For $t \leq t_0$, we have $X_t = (\Delta_t)_j = \Gamma \cdot r_t^T u_{t,j}$ for some unit vector $u_{t,j}$. Recall that r_t is n -dimensional Gaussian, independent of $u_{t,j}$. Recall that r_t is spherically symmetric — so $r_t^T u_{t,j}$ has the ^{1-dimensional} standard normal distribution $N(0, 1)$ and $\boxed{\Gamma \cdot r_t^T u_{t,j}}$ has the 1-dimensional normal distribution $N(0, \Gamma^2)$.

We want to claim that X_1, \dots, X_{t_0} are independent random variables. Again, one has to be careful since t_0 itself is not independent of X_1, X_2, \dots , so even the formulation of such a claim may not be clear.

So we will formulate our claim of independence in the following way. Let Z'_1, \dots, Z'_l be a new sequence of independent random variables, each with the $N(0, \sigma^2)$ distribution. We define another sequence of random variables Z_1, Z_2, \dots, Z_l as follows:

$$Z_t = \begin{cases} X_t & \text{for } t \leq t_0 \\ Z'_t & \text{for } t > t_0. \end{cases}$$

We claim that Z_1, Z_2, \dots, Z_l are independent.

The justification is that if we fix the values of r_1, \dots, r_l in the algorithm and also the values of the auxiliary variables Z'_1, \dots, Z'_{t-1} arbitrarily, the values of Z_1, \dots, Z_{t-1} are determined uniquely, while Z_t has the $N(0, \sigma^2)$ distribution.

So if the j -th coordinate walk does not terminate, then all the partial sums $Z_1 + Z_2 + \dots + Z_t$ belong to $(-1, 1)$, $t = 1, 2, \dots, l$.

Lemma. Let Z_1, \dots, Z_l be independent random variables, each with the $N(0, \tau^2)$ distribution. Then the probability that all of the partial sums $\sum_{i=1}^t Z_i$, $t = 1, 2, \dots, l$ are in $(-1, 1)$ is at most $e^{-k_1 \lfloor \tau^2 l \rfloor}$, for a suitable constant k_1 .

So Claim 1 follows from the above lemma (since $l = \frac{C_1 \cdot \log n}{\tau^2}$). Note that the above lemma implies that the probability the j -th coordinate random walk does not terminate $\leq e^{-k_1 [C_1 \log n]} \leq \frac{1}{n^2}$ for $C_1 = \frac{3}{k_1}$ and n sufficiently large.

Proof of the lemma. Let $k = \tau^2$. Let us assume for convenience that $k \in \mathbb{Z}$. Let us partition the sequence Z_1, Z_2, \dots into contiguous blocks of length k and let s_j be the sum of the j -th block. Formally, $s_j = \sum_{i=(j-1)k+1}^{jk} Z_i$. The number of blocks is $\lfloor \frac{l}{k} \rfloor$.

Fact. If X, Y are independent $N(0, 1)$ random variables and $a, b \in \mathbb{R}$, then $ax + by$ has the 1-dimensional normal distribution $N(0, a^2 + b^2)$.

So each s_j has the standard normal distribution $N(0, k\tau^2) = N(0, 1)$. Thus $\Pr[|s_j| \geq 2] \geq c_0$ for a suitable c_0 (in fact, $c_0 \approx 0.0455$).

If $\sum_{i=1}^t z_i \in (-1, 1)$ for all $t = 1, 2, \dots, l$, then necessarily $|\delta_j| < 2$ for all j . The δ_j are independent, thus the probability that $|\delta_j| < 2$ for all j is $\leq (1 - c_0)^{\lfloor L/l \rfloor} = e^{-k_1 L^{r^2 l}}$. Thus the lemma is proved. \blacksquare

We will next show the following claim.

Claim 2. The discrepancy of the (semi)coloring x_l is $O(H \cdot \log(mn))$, where H is the hereditary vector discrepancy of F .

(Note that if x_l is not a coloring (this happens w.p. $\leq 1/n$), we restart the algorithm from scratch. Thus x_l is always a coloring.) as returned by the algo.

The vector discrepancy of F (denoted by $\text{vecdisc}(F)$) is $\min D \geq 0$ s.t. $\left\| \sum_{j \in S_i} u_j \right\|^2 \leq D^2$ for $i = 1, 2, \dots, m$ and $\|u_j\|^2 = 1$ for $j = 1, \dots, n$. The hereditary vector discrepancy of F is $\max_{A \subseteq V} \text{vecdisc}(F|_A)$.

In order to prove Claim 2, we will prove

$$\Pr[\text{disc}(F, x_l) > D_{\max}] \leq 1/n,$$

where $D_{\max} = O(H \cdot \log(mn))$.

Let us fix a set $S_i \in F$ and let $D_i = \sum_{j \in S_i} (x_l)_j$ be its imbalance in x_l . We will show $\Pr[|D_i| > D_{\max}] \leq \frac{1}{mn}$ for every i . Then Claim 2 will follow from the union bound.