

- ① Architecture Team: Discuss TSP interface.
 - ② Class: Discussion on TSP. { Go over LP-relaxation } .
 - ③ Optical Mapping:
 - ④ Quiz.
 - ⑤ Optical Mapping Architecture Team.
- .

Optical Mapping:

❖ Restriction Map Model:

SMRM (Single Molecule Restriction Map)

A vector with ordered set of rational numbers on the open interval $(0, 1)$:

$$\mathcal{D}_j = (s_{1j}, s_{2j}, \dots, s_{Mj}),$$

$$0 < s_{1j} < s_{2j} < \dots < s_{Mj} < 1. \quad s_{ij} \in \mathbb{Q}$$

❖ Problem

Data: A collection of SMRM vectors:

$$\mathcal{D}_1, \mathcal{D}_2, \dots, \mathcal{D}_m$$

Desiderata: Compute a consensus vector

$$H = (h_1, h_2, \dots, h_N)$$

such that H is "consistent" with each \mathcal{D}_j .

$$H^* = \arg \min_{H, j} \text{dist}(\mathcal{D}_j, H).$$

(19)

Consensus:

$$H^* = \underset{H, j}{\arg \min} \text{dist}(D_j, H)$$

$$D_j = (s_{1j}, s_{2j}, \dots, s_{Mj;j})$$

$$\Rightarrow D_j + c = (s_{1j} + c, s_{2j} + c, \dots, s_{Mj;j} + c)$$

$$c \in [0, 1] \quad c \in \mathbb{Q} \quad -s_{1j} < c < 1 - s_{Mj;j}$$

$$\text{dist}(D_j, H) = \text{dist}(D_j + c, H)$$

$$D_j^R = (1 - s_{Mj;j}, \dots, 1 - s_{2j}, 1 - s_{1j})$$

$$\text{dist}(D_j^R, H) = \text{dist}(D_j, H).$$

Consensus:

$$H^* = \underset{H, j}{\arg \min} \left\{ \text{dist}(D_j, H), \text{dist}(D_j^R, H) \right\}$$

or

$$H^* = \underset{H, j}{\arg \min} \left\{ \text{dist}(D_j + c, H) \mid -s_{1j} < c < 1 - s_{Mj;j} \right\}$$

or

$$H^* = \underset{H, j}{\arg \min} \left\{ \text{dist}(D_j + c, H), \text{dist}(D_j^R + c, H) \right\}$$

$\underbrace{\quad \quad \quad}_{-s_{1j} < c < 1 - s_{Mj;j}}$.

Assume some distribution generating D_j 's \Rightarrow Maximum Likelihood formulation \Rightarrow

$$\langle H \rangle = \underset{H}{\arg \min} \sum_j \min \left\{ \text{dist}(D_j + c, H), \text{dist}(D_j^R + c, H) \mid -s_{1j} < c < 1 - s_{Mj;j} \right\}$$

TOY EXAMPLE PROBLEM.

(20)

(Unknown Orientation:)

Data: A set of ordered vectors with rational entries in the open interval $(0, 1)$:

$$D_1, D_2, \dots, D_\ell, D_{\ell+1}, \dots, D_m$$

A rational number $p_c \in (0, 1)$ and an integer N .

An admissible alignment of the data can be represented as

$$D'_1, D'_2, \dots, D'_\ell, D'_{\ell+1}, \dots, D'_m$$

where

$$\left. \begin{array}{l} D'_j \in \{D_j, D_j^R\} \quad (1 \leq j \leq \ell) \\ D'_j = D_j \quad (j > \ell) \end{array} \right\} \begin{array}{l} = \text{An Alignment} \\ (A_k) \end{array}$$

and

For any rational number $h_i \in [0, 1]$, define an indicator variable

$$m_{ijk} = \begin{cases} 1 & \text{if } h_i \in D'_j \\ 0 & \text{otherwise.} \end{cases}$$

Define a characteristic function

$$x_k : [0, 1] \rightarrow \{0, 1\}$$

$$: h_i \mapsto \begin{cases} 1 & \text{iff } \sum_j m_{ijk} > p_c^m. \end{cases}$$

Desiderata: Find an admissible alignment A_K such that (21)

$$|\{h \in [0,1] \mid X_K(h) = 1\}| \geq N.$$

NP-Completeness

Consider an instance of a 3-SAT problem:
With l variables:

$$x_1, x_2, \dots, x_l$$

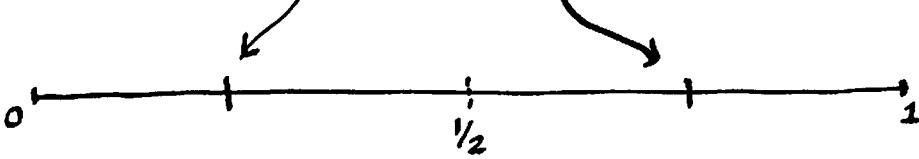
And n clauses:

$$C_1, C_2, \dots, C_n \quad (n \geq l)$$

- Assume that no clause contains a variable and its negation: x_j and \bar{x}_j (The clause is a tautology $\equiv T$)
- Restriction site associated with a ~~at~~ clause C_i .

$$f_i = \frac{i}{2(n+1)}$$

$$f_i^R = 1 - f_i = \frac{2n-i+2}{2(n+1)}$$



D_1

$f_i \leftarrow \text{if } x_j \in C_i$

\vdots

D_2

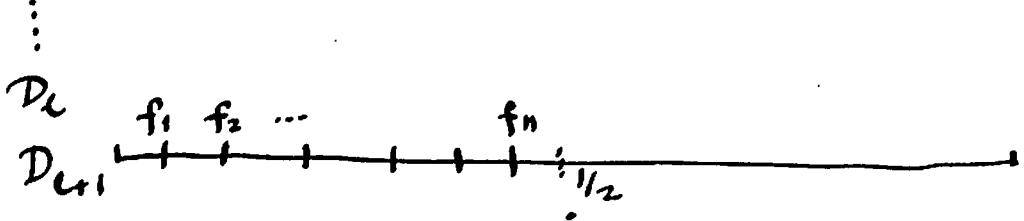
$f_i \leftarrow \text{if } \bar{x}_j \in C_i$

\vdots

$f_i^R \leftarrow \text{if } \bar{x}_j \in C_i$

D_L

f_1, f_2, \dots, f_n



$D_{m+1} \vdash \vdash \vdash \vdash \vdash \vdash$

Create a dataset $D_1, D_2, \dots, D_\ell, D_{\ell+1}, \dots, D_m$ (22)

with $m = 2\ell - 1$ as follows:

D_j has a cuts at f_i or f_i^R , only:

$$f_i \in D_j \text{ iff } x_j \in C_i$$

$$(f_i^R \in D_j \text{ iff } \bar{x}_j \in C_i)$$

$$N \equiv n, p_c = \frac{1}{2}$$

~

CNF has a satisfying assignment

⇒ Choose an admissible alignment in which

$$D'_j = \begin{cases} D_j & \text{if } x_j = \text{true} \\ D_j^R & \text{if } x_j = \text{false} \end{cases} \quad 1 \leq j \leq \ell$$

$$D'_j = D_j, \quad \ell < j \leq m.$$

∴ For every f_i , ($1 \leq i \leq n$) there are $(\ell - 1)$ matches from $D_{\ell+1}, \dots, D_m$

& at least one more from D'_1, \dots, D'_ℓ
(since each clause must be satisfied)

$$\therefore \forall_{1 \leq i \leq n} \sum_j m_{ijk} \geq \ell > \frac{2\ell - 1}{2} = p_c m.$$

$$\Rightarrow \{h \in [0,1] | x_k(h) = 1\} = \{f_1, f_2, \dots, f_n\}$$

$$\Rightarrow |h \in [0,1] | x_k(h) = 1| = n \geq N.$$

(23)

Conversely, if the CNF has no satisfying assignment, then for every admissible alignment there exists an $1 \leq i \leq n$

$$\forall_k \exists_i \quad \sum_j m_{ijk} = (l-1) < p_c n \quad \text{and} \\ |\{h \in [0,1] \mid x_n(h) = 1\}| < n. \quad \square.$$

Problem Generation:

Statistical Model:

- A model or hypothesis H .

$$= \{h_1, h_2, \dots, h_N\} \quad N \approx 40.$$

Distribution for h_i 's
Exponential gaps
or uniform gaps.

- $\Pr[D_j | H]$

$$D_j \sim H. \quad \left\{ \begin{array}{l} \text{Poisson Conditional Indep.} \\ \Pr[D_j | D_{j1}, \dots, D_{jm}, H] \\ = \Pr[D_j, H] \end{array} \right.$$

- $\Pr[\text{bad}]$, $\Pr[\text{good}] = 1 - \Pr[\text{bad}]$

$$\Pr[D_j | H] = \frac{1}{2} \sum \Pr[D_j^{(k)} | H, \text{good}] \Pr[\text{good}]$$

$$+ \frac{1}{2} \sum \Pr[D_j^{(k)} | H, \text{bad}] \Pr[\text{bad}]$$

(k) \rightarrow Alignment.

$D_j^{(k)} = D_j \text{ or } D_j^R$ with equal probability:

$D_j = \text{Good} \Rightarrow$

Choose parameters $p_c, \sigma, f.$

$h_i \in H \Rightarrow s_i \sim N(h_i, \sigma)$ with $\text{pr} = p_c.$

$s_i = \text{absent}$ with $\text{pr} = 1 - p_c.$

~~~~~.

spurious cuts  $\Rightarrow$  expand Poisson.  $e^{-\lambda_f} \frac{\lambda_f^{F_{jk}}}{F_{jk}!}$

$D_j = \text{Bad} \Rightarrow$

Poisson:  $e^{-\lambda_n} \frac{\lambda_n^{M_j}}{M_j!}$

$\text{Pr}[D_j^{(k)} | \text{good}]$

$$= \prod_{i=1}^N \left[ \Phi(p_c) \frac{e^{-(s_{ij} - h_i)^2/2\sigma^2}}{\sqrt{2\pi} \sigma_i} \right]^{m_{ijk}} (1-p_c)^{(1-m_{ijk})}$$

$$\times e^{-\lambda_f} \lambda_f^{F_{jk}}.$$

$$\text{Pr}[D_j^{(k)} | \text{bad}] = e^{-\lambda_n} \lambda_n^{M_j}$$