Like a Needle in a Haystack: Rates and Algorithms for Group Testing

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Joint work with Oliver Johnson, Matthew Aldridge and Karen Gunderson

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Warm-up examples

- Molecular biology (DNA screening)
- Recommender systems



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- Molecular biology (DNA screening)
- Recommender systems
- Spectrum sensing
- High-throughput screening techniques
- Network tomography
- Cryptography and cyber security

Problem

Find a specific, rare sequence of nucleotides among many DNA samples



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 \sim

 $\mathcal{N}\mathcal{N}\mathcal{N}$

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Problem

Suggest songs to a user based on his past preferences.



Problem Suggest songs to a user based on his past preferences.

User selects song



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COMMON FEATURES



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• Search for *sparse* property



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- Property can be tested on groups



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GROUP TESTING Search methods to recover a sparse subset of items from a population that share a feature which can be detected on groups



















Anatomy of an algorithm





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 bits per test



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 R_{A} measures how much we learn with each test, on average. $R_{A}^{*}(\beta)$ supremum of rates for algorithm A. (Yes, it's bounded).

Why do we like rates?



Universal upper bound for noiseless GT

Theorem (Aldridge, B., Johnson, 2013)

The probability of success of any algorithm A can be upper-bounded as

$$\mathbb{P}(success) \leq rac{2^{T_{A}}}{\binom{N}{K}}$$
 .



Two important consequences





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Rates and Algorithms for Group Testir
Two important consequences

$$\mathbb{P}(\text{success}) \leq \frac{2^{T_{A}}}{\binom{N}{K}}$$

• Successful algorithms use $T \ge \log_2 {\binom{N}{K}}$ tests optimal algorithms use $T = c \log_2 {\binom{N}{K}}$ tests



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• Successful algorithms have $R^*_{\mathtt{A}}(\beta) \leq 1$

















Can we get to R = 1?







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Rates and Algorithms for Group Testing

Can we get to R = 1?



Can we get to R = 1?



Can we get to R = 1? population



Positive, halve















Can we get to R = 1?





- Hwang, 1972 (HGBS)
- Achieves $R_{\text{HGBS}} = C = 1$
- Careful choice of sample size is key



• Maybe



- Maybe
- What we did:



- Maybe
- What we did:
 - Introduced and studied DD (Definite Defectives) and SSS (Smallest Satisfying Set)
 - ...hence Bernoulli sampling, $x_{ij} \sim \text{Bern}(p)$, $p = \frac{1}{K+1}$
 - Showed limitation of Bernoulli-based algorithms







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$$\left\{ \begin{array}{l} T_{\text{DD}} \leq \max\left\{\beta, 1 - \beta\right\} \text{e}K \ln N \\ R_{\text{DD}}^*(\beta) \geq \frac{1}{e \ln 2} \min\left\{1, \frac{\beta}{1 - \beta}\right\} \end{array} \right\}$$

What $R^*_{DD}(\beta)$ looks like



Computing $R^*_{DD}(\beta)$: core of the proof – construction


• $\mathcal{PD} =$

{items not in negative tests}



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For every defective $i \in \mathcal{K}$: $L_i = \#$ tests with i and no item from \mathcal{PD}



















$$\sum_{g=0}^{N-K} \binom{N-K}{g} (1-p)^{gm_0} (1-(1-p)^{m_0})^{N-K-g}$$
$$\times \mathbb{P}(\text{success} \mid M_0 = m_0, G = g)$$
$$\geq \max\{0, 1-K \exp(\Theta(T, m_0))\}$$



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$$M_0 \in (\mathbb{E}M_0 - \varepsilon, \mathbb{E}M_0 + \varepsilon)$$
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- $M_0 \in (\mathbb{E}M_0 \varepsilon, \mathbb{E}M_0 + \varepsilon)$ w.h.p.
- $\Theta(T, m_0) \leq -(\max\{\beta, 1-\beta\}) \ln N$ for M_0 close to $\mathbb{E}M_0$





Group testing as LP:



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minimize
$$\mathbf{1}^{\top} \mathbf{z}$$

subject to $\mathbf{x}_t \cdot \mathbf{z} = 0$ for t with $y_t = 0$
 $\mathbf{x}_t \cdot \mathbf{z} \ge 1$ for t with $y_t = 1$
 $(\mathbf{1}^{\top} \mathbf{z} \ge K)$
 $\mathbf{z} \in \{0, 1\}^N$



Group testing as LP:

minimize subject to $\mathbf{1}^{\mathsf{T}} \mathbf{z}$ $\mathbf{x}_{t} \cdot \mathbf{z} = 0 \quad \text{for } t \text{ with } y_{t} = 0$ $\mathbf{x}_{t} \cdot \mathbf{z} \ge 1 \quad \text{for } t \text{ with } y_{t} = 1$ $(\mathbf{1}^{\mathsf{T}} \mathbf{z} \ge K)$ $\mathbf{z} \in \{0, 1\}^{N}$ • SSS brute-force searches for a satisfying ${\cal K}$



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$$R_{\rm SSS}^*(\beta) \ge \frac{1}{\ln 2} \max_{\alpha \in [\ln 2, 1]} \min\left\{ 2\alpha e^{-\alpha} \frac{\beta}{2-\beta}, -\ln\left(1-2e^{-\alpha}+2e^{-2\alpha}\right) \right\} .$$

What $R^*_{\rm SSS}(\beta)$ looks like



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Definition

X is K-separable if, for any K-subsets $\mathcal{L}, \mathcal{M} \subset \{1, \dots, N\}$, it is

$$\bigvee_{i\in\mathcal{M}}\mathbf{x}^{(i)}\neq\bigvee_{j\in\mathcal{L}}\mathbf{x}^{(j)}$$



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Notice that $\mathbf{y} = \bigvee_{\mathcal{K}} \mathbf{x}^{(i)}$.

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X is ε -almost K-separable if there are at most $\varepsilon \binom{N}{K}$ K-subsets that break separability.



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ldea: What if we relax separability?

Definition

X is ε -almost K-separable if there are at most $\varepsilon \binom{N}{K}$ K-subsets that break separability.

- Almost-separable matrices exist
- Need $T = O(K \log N)$ tests to get one
- A Bernoulli test design is almost-separable w.h.p. (via concentration)



Theorem (Aldridge, B., Johnson, 2014)

Consider SSS using T_{SSS} tests. Then,

$$\mathbb{P}(\text{success}) \to 1 \Rightarrow T_{\text{SSS}} > \frac{(1-\beta) \text{e} \ln 2}{\beta} \log_2 \binom{N}{K} ,$$

and, if the necessary condition is violated,

$$T_{\rm SSS} \le \frac{(1-\beta) e \ln 2}{\beta} \log_2 \binom{N}{K} \Rightarrow \mathbb{P}(\text{success}) \le \frac{2}{3}$$



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Adaptivity gap



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• Group testing



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 - Collective detection of sparse properties
 - Information-poor tests



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- SSS: bound the performance of *any* non-adaptive algorithm based on Bernoulli sampling
- DD: order-optimal, rate-optimal (for dense problems)
- Future work: noise models, applications, non-identical GT, non-independent GT, collateral open questions...