

Ranking Methods in Machine Learning

A Tutorial Introduction

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In God we trust.
All others bring data.

W. Edwards Deming

Machine Learning



Computer Science

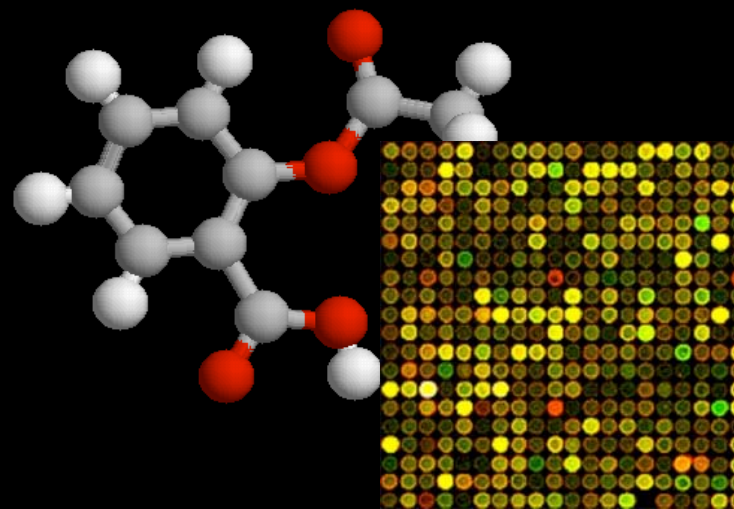


Statistics

Machine Learning



Mathematical
Optimization



Machine Learning





$A > D > B > C$



$C > A > B > D$

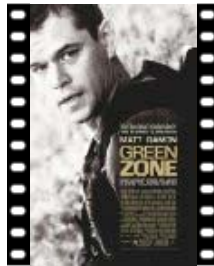
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$A > D > B > C$

$E \quad F \quad G \quad ?$





Road Map

Theory

Algorithms

Applications



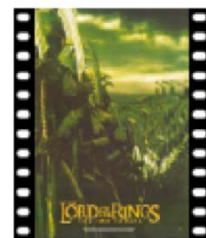
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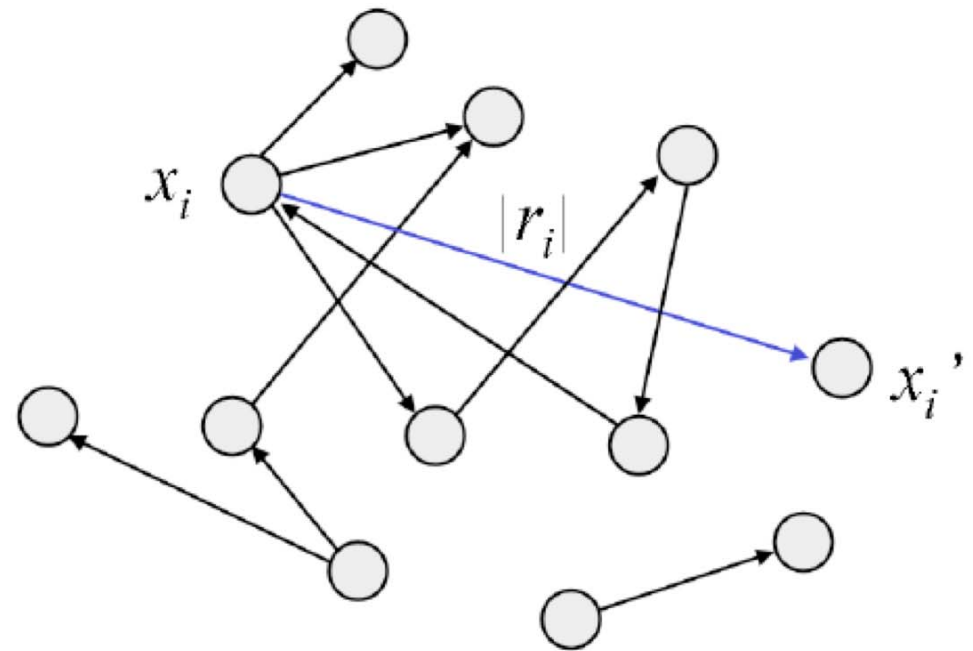
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General Ranking Problem

- ▶ Instance space X
- ▶ **Input:** Training sample $S = ((x_1, x'_1, r_1), \dots, (x_m, x'_m, r_m)) \in (X^2 \times \mathbb{R})^m$
- ▶ **Output:** Ranking function $f : X \rightarrow \mathbb{R}$





Likes



...

Dislikes



...



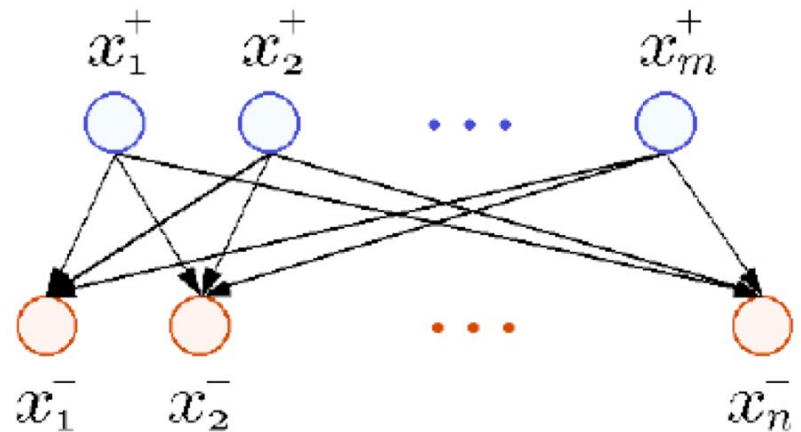
Bipartite Ranking Problem

- ▶ Instance space X
- ▶ **Input:** Training sample $S = (S_+, S_-)$:

$$S_+ = (x_1^+, \dots, x_m^+) \in X^m \quad (\text{positive examples})$$

$$S_- = (x_1^-, \dots, x_n^-) \in X^n \quad (\text{negative examples})$$

- ▶ **Output:** Ranking function $f : X \rightarrow \mathbb{R}$



Bipartite Ranking Problem

▶ Instance space X

▶ **Input:** Training sample $S = (S_+, S_-)$:

$$S_+ = (x_1^+, \dots, x_m^+) \in X^m \quad (\text{positive examples})$$

$$S_- = (x_1^-, \dots, x_n^-) \in X^n \quad (\text{negative examples})$$

▶ **Output:** Ranking function $f : X \rightarrow \mathbb{R}$

▶ Expected error: $\mathbf{er}(f) = \mathbf{P}_{(x,x') \sim \mathcal{D}_+ \times \mathcal{D}_-} [f(x) < f(x')]$

▶ Empirical error: $\widehat{\mathbf{er}}_S(f) = \frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n \mathbf{1}(f(x_i^+) < f(x_j^-))$

Generalization Bounds Review

Informally:

How does the empirical performance of a learned function generalize to its expected performance on future data?

Formally:

Let $f_S : X \rightarrow \mathbb{R}$ denote the ranking function learned from $S \in X^m \times X^n$.

- ▶ Expected error: $\mathbf{er}(f_S) = \mathbf{P}_{(x,x') \sim \mathcal{D}_+ \times \mathcal{D}_-} [f_S(x) < f_S(x')]$
- ▶ Empirical error: $\widehat{\mathbf{er}}_S(f_S) = \frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n \mathbf{1}(f_S(x_i^+) < f_S(x_j^-))$

Assume $S \sim \mathcal{D}_+^m \times \mathcal{D}_-^n$. Can we bound $\mathbf{er}(f_S)$ in terms of $\widehat{\mathbf{er}}_S(f_S)$?

Generalization Bounds Based on Uniform Convergence

Let $f_S : X \rightarrow \mathbb{R}$ denote the ranking function learned from $S \in X^m \times X^n$.

Want to bound $\text{er}(f_S)$ in terms of $\widehat{\text{er}}_S(f_S)$.

Uniform convergence approach: If $f_S \in \mathcal{F}$, then

$$\mathbf{P}_S \left[\left| \text{er}(f_S) - \widehat{\text{er}}_S(f_S) \right| \geq \epsilon \right] \leq \mathbf{P}_S \left[\sup_{f \in \mathcal{F}} \left| \text{er}(f) - \widehat{\text{er}}_S(f) \right| \geq \epsilon \right].$$

Sufficient to bound
this probability

[Vapnik & Chervonenkis, 1971]

Bounding

$$\mathbb{P}_S \left[\sup_{f \in \mathcal{F}} |\text{er}(f) - \widehat{\text{er}}_S(f)| \geq \epsilon \right]$$

Step 1: Symmetrization

Variance inequality
due to Devroye (1991)

$$\mathbb{P}_S \left[\sup_{f \in \mathcal{F}} |\text{er}(f) - \widehat{\text{er}}_S(f)| \geq \epsilon \right] \leq 2 \mathbb{P}_{S, \tilde{S}} \left[\sup_{f \in \mathcal{F}} |\widehat{\text{er}}_{\tilde{S}}(f) - \widehat{\text{er}}_S(f)| \geq \frac{\epsilon}{2} \right]$$

Step 2: Permutations and reduction to a finite class

$$\mathbb{P}_{S, \tilde{S}} \left[\sup_{f \in \mathcal{F}} |\widehat{\text{er}}_{\tilde{S}}(f) - \widehat{\text{er}}_S(f)| \geq \frac{\epsilon}{2} \right] \leq \pi_{\mathcal{F}}(2m, 2n) \cdot 2 \exp \left(\frac{-m n \epsilon^2}{8(m+n)} \right)$$

Bipartite rank-shatter coefficients
(New complexity measure for
classes of ranking functions)

McDiarmid's
inequality

Uniform Convergence Bound

Theorem. Let \mathcal{F} be a class of real-valued functions on X . Then for any $0 < \delta < 1$, we have with probability at least $1 - \delta$:

$$\sup_{f \in \mathcal{F}} |\text{er}(f) - \widehat{\text{er}}_S(f)| < \sqrt{\frac{8(m+n)}{mn} \left(\ln \pi_{\mathcal{F}}(2m, 2n) + \ln \left(\frac{4}{\delta} \right) \right)}.$$

[Agarwal et al, 2005]

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Bipartite Ranking: Basic Algorithmic Framework

Minimize a convex upper bound on the empirical ranking error, possibly with some regularization, over some class of ranking functions:

$$\min_{f \in \mathcal{F}} \left[\frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n \ell(f, x_i^+, x_j^-) + \lambda N(f) \right]$$

where

$\ell(f, x_i^+, x_j^-)$: convex upper bound on $\mathbf{1}(f(x_i^+) < f(x_j^-))$

$N(f)$: regularizer

$\lambda > 0$: regularization parameter

\mathcal{F} : class of ranking functions

Bipartite RankBoost Algorithm

$$\min_{f \in \mathcal{L}(\mathcal{F}_{\text{base}})} \left[\frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n \ell_{\text{exp}}(f, x_i^+, x_j^-) \right]$$

$$\ell_{\text{exp}}(f, x_i^+, x_j^-) = \exp \left(- \left(f(x_i^+) - f(x_j^-) \right) \right)$$

$\mathcal{L}(\mathcal{F}_{\text{base}})$ = linear combinations of functions in some
base class $\mathcal{F}_{\text{base}}$

[Freund et al, 2003]

Bipartite RankSVM Algorithm

$$\min_{f \in \mathcal{F}_K} \left[\frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n \ell_{\text{hinge}}(f, x_i^+, x_j^-) + \frac{\lambda}{2} \|f\|_K^2 \right]$$

$$\ell_{\text{hinge}}(f, x_i^+, x_j^-) = \left(1 - \left(f(x_i^+) - f(x_j^-) \right) \right)_+ \quad [u_+ = \max(u, 0)]$$

\mathcal{F}_K = reproducing kernel Hilbert space (RKHS)
with kernel function K

$$N(f) = \frac{\|f\|_K^2}{2}$$

[Herbrich et al, 2000; Joachims, 2002; Rakotomamonjy, 2004]

Bipartite RankNet Algorithm

$$\min_{f \in \mathcal{F}_{\text{neural}}} \left[\frac{1}{mn} \sum_{i=1}^m \sum_{j=1}^n \ell_{\text{logistic}}(f, x_i^+, x_j^-) \right]$$

$$\ell_{\text{logistic}}(f, x_i^+, x_j^-) = \log \left(1 + \exp \left(- \left(f(x_i^+) - f(x_j^-) \right) \right) \right)$$

$\mathcal{F}_{\text{neural}}$ = functions represented by some class of neural networks

[Burges et al, 2005]

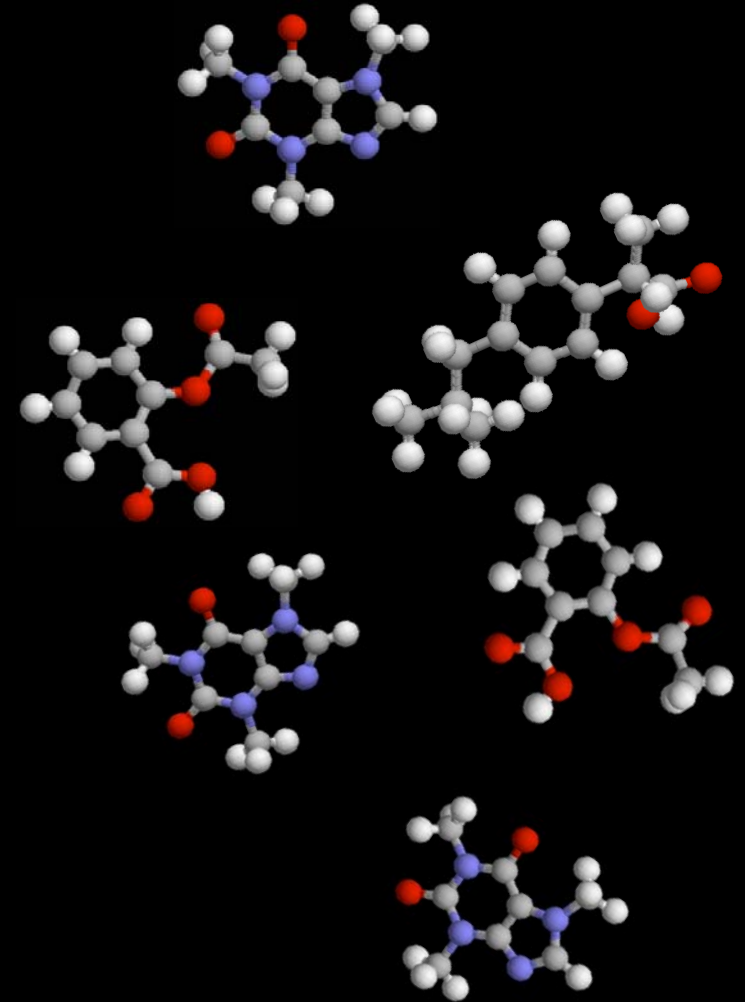
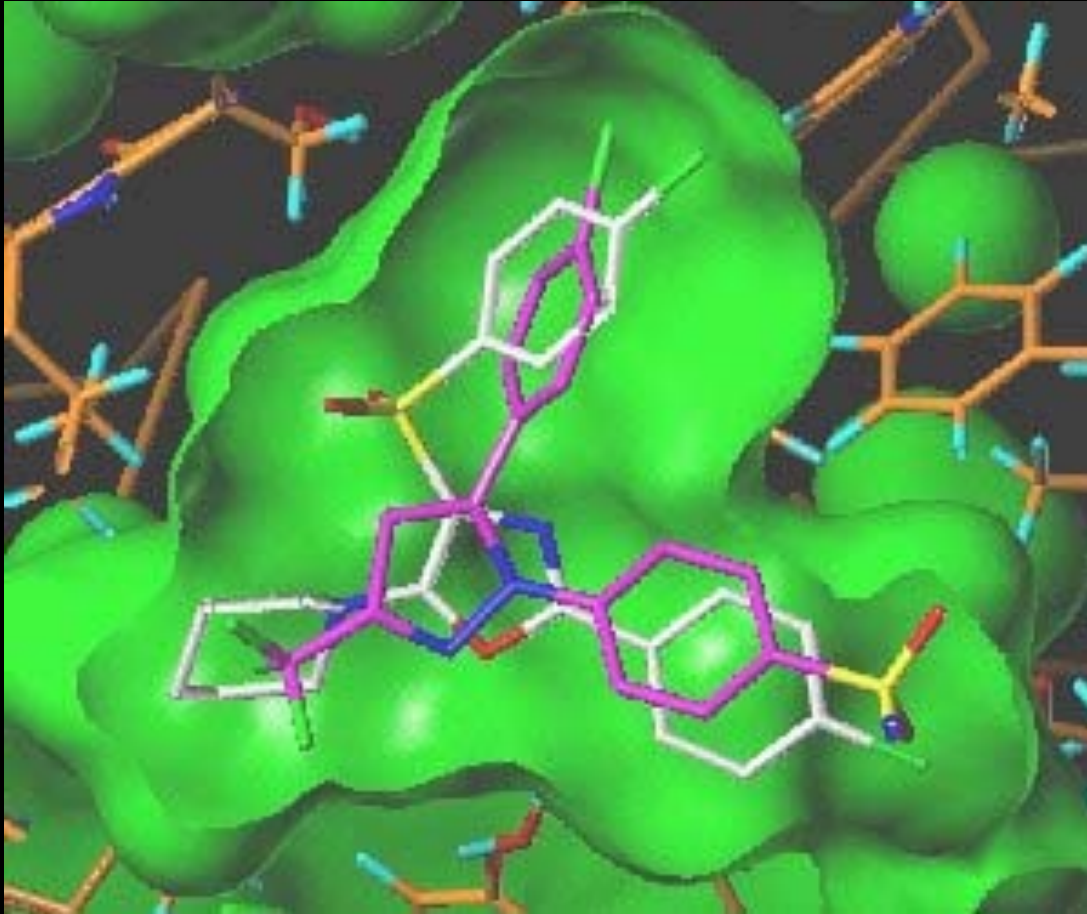
Road Map

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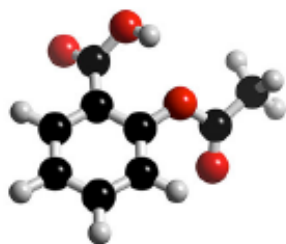
Applications

Application to Drug Discovery

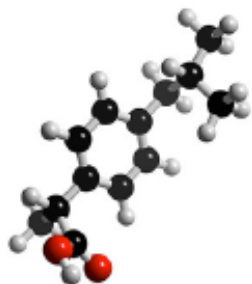


Problem: Millions of structures in a chemical library.
How do we identify the most promising ones?

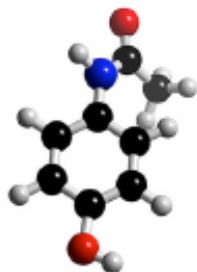
Formulation as a Ranking Problem with Real-Valued Labels



$$pIC_{50} = 5.6718$$



$$pIC_{50} = 8.2991$$



$$pIC_{50} = 4.1317$$

...

Ranking With Real-Valued Labels

- ▶ Instance space X
- ▶ Real-valued labels $Y = \mathbb{R}$
- ▶ **Input:** Training sample $S = ((x_1, y_1), \dots, (x_m, y_m)) \in (X \times \mathbb{R})^m$
- ▶ **Output:** Ranking function $f : X \rightarrow \mathbb{R}$

- ▶ Expected error:

$$\text{er}(f) = \mathbb{E}_{((x,y),(x',y')) \sim \mathcal{D} \times \mathcal{D}} [|y - y'| \mathbf{1}((y - y')(f(x) - f(x')) < 0)]$$

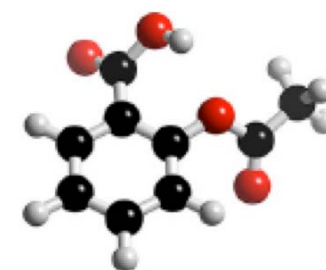
- ▶ Empirical error:

$$\widehat{\text{er}}_S(f) = \frac{1}{\binom{m}{2}} \sum_{i=1}^{m-1} \sum_{j=i+1}^m |y_i - y_j| \mathbf{1}((y_i - y_j)(f(x_i) - f(x_j)) < 0)$$

Cheminformatics Data Sets

[Sutherland et al, 2004]

Data Set	No. of Compounds	No. of Chemical (2.5D) Descriptors	pIC ₅₀ Values
DHFR inhibitors	361	70	3.3 – 9.8
COX2 inhibitors	292	74	4.0 – 9.0



DHFR Results Using RankSVM

2.5D chemical descriptors
Gaussian kernel

Training size	Ranking error	
	SVR	RankSVM
24	0.4755	0.4601
48	0.3430	0.3509
72	0.2840	0.2726
96	0.2483	0.2351
120	0.2171	0.2121
144	0.2023	0.2032
168	0.2019	0.1817
192	0.1808	0.1749
216	0.1816	0.1722
237	0.1714	0.1681

FP2 molecular fingerprints
Tanimoto kernel

Training size	Ranking error	
	SVR	RankSVM
24	0.3793	0.3546
48	0.2905	0.2896
72	0.2517	0.2421
96	0.2343	0.2201
120	0.2147	0.2052
144	0.2166	0.1988
168	0.2096	0.1966
192	0.2056	0.1962
216	0.1907	0.1787
237	0.1924	0.1798

[Agarwal et al, 2010]



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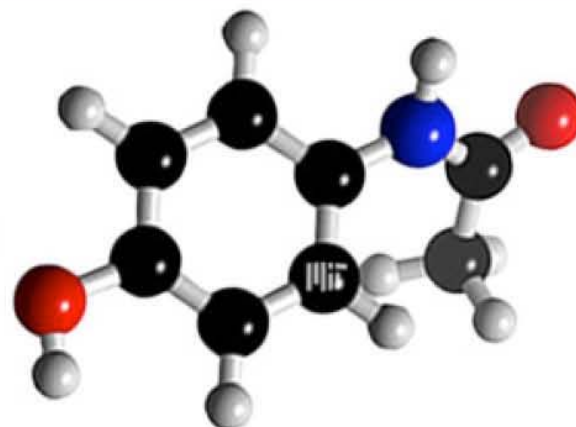
energy | cancer | diversity | global

impact

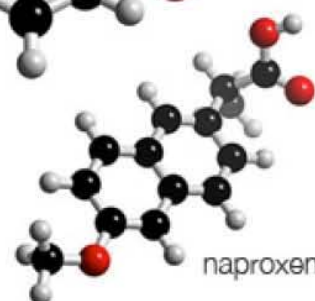
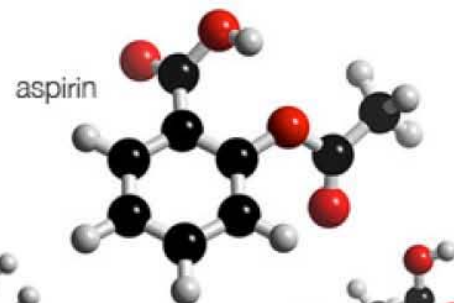
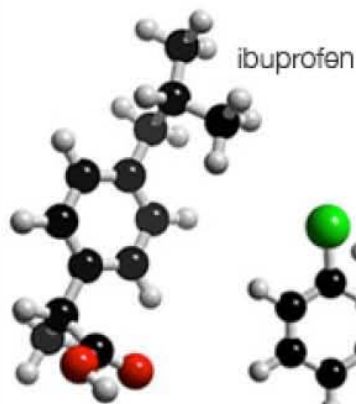
industry | public service

today's spotlight
Build a pill
Ranking algorithms could expedite drug development

If acetaminophen worked ...



... try one of these!



news

Letter to the community on MIT's financial condition

In The World: Better wound treatment for all

Toward more efficient wireless power delivery

research | campus | press

events

Memorial service for former MIT President Howard W. Johnson (today)

Artists Beyond the Desk concert (today)

Of Note: Celebrating SA+P's new Program in Art, Culture and Technology (tomorrow)

Legatum Lecture: Entrepreneur and investor Chuck Lacy (tomorrow)



Application to Bioinformatics



Searching for genetic determinants in the new millennium

N.J. Risch

Human genetics is now at a critical juncture. The molecular methods used successfully to identify the genes underlying rare mendelian syndromes are failing to find the numerous genes causing more common, familial, non-mendelian diseases . . .

Nature 405:847–856, 2000



Application to Bioinformatics



Searching for genetic determinants in the new millennium

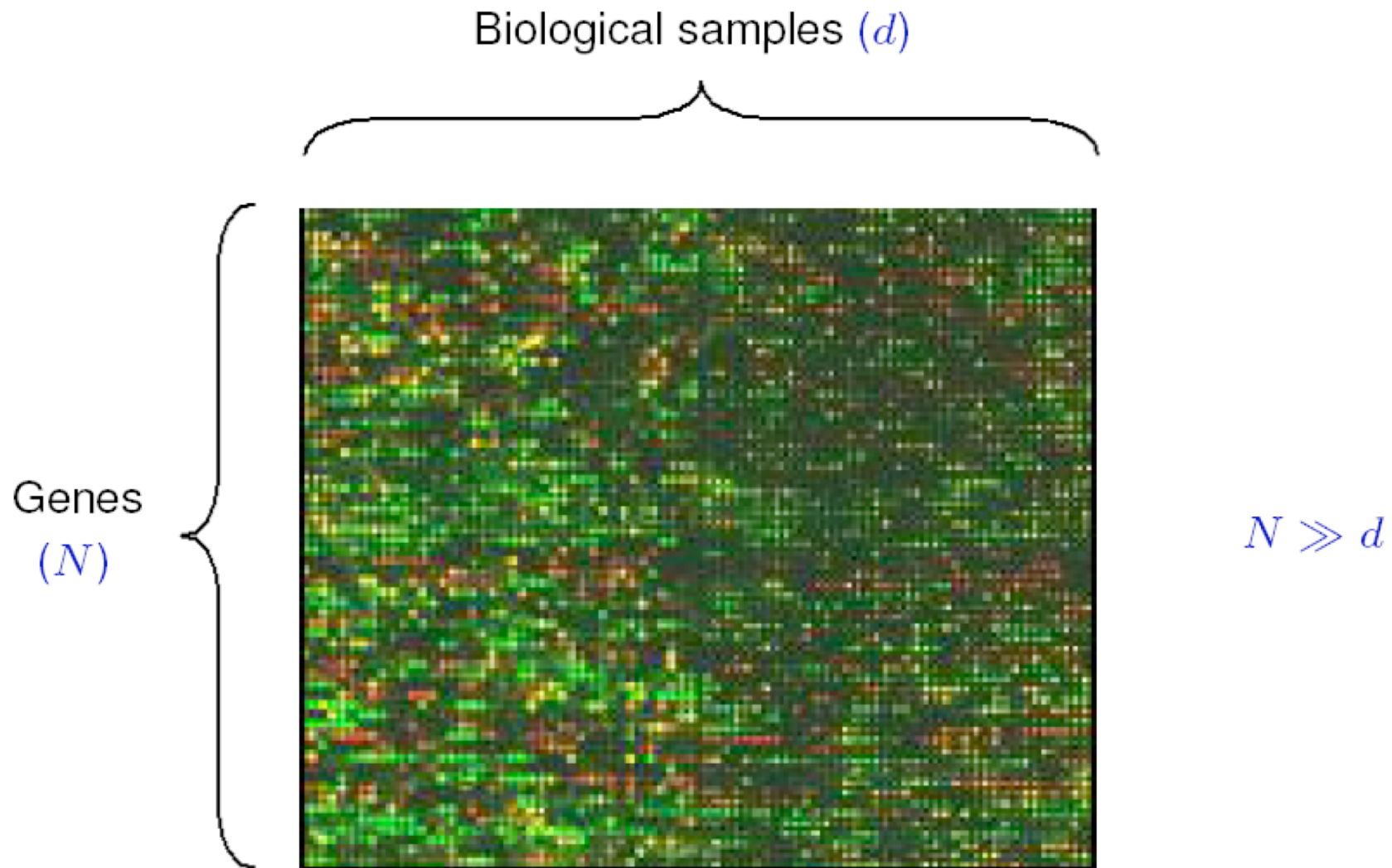
N.J. Risch

With the human genome sequence nearing completion, new opportunities are being presented for unravelling the complex genetic basis of nonmendelian disorders based on large-scale genomewide studies . . .

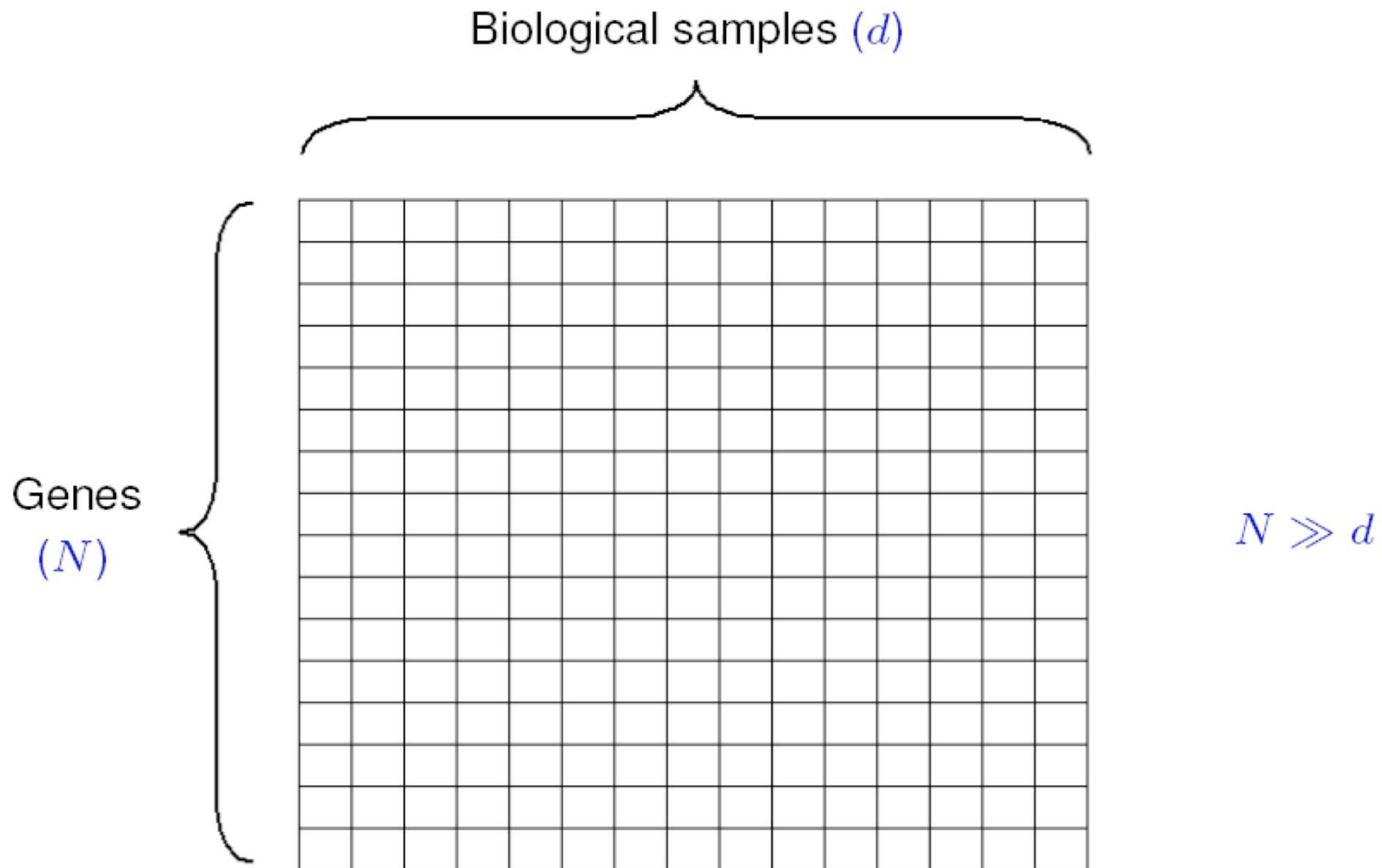


Nature 405:847–856, 2000

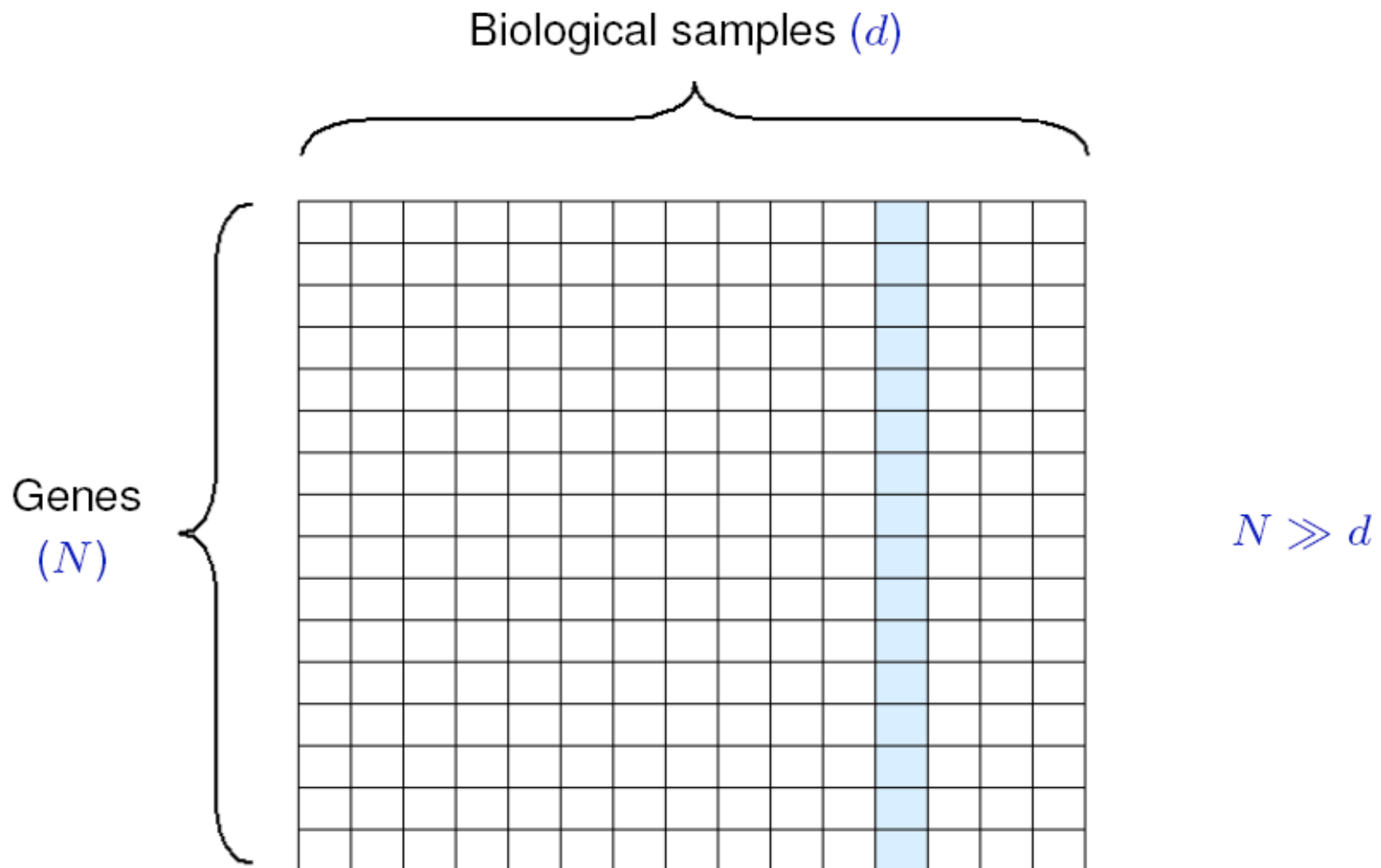
Identifying Genes Relevant to a Disease Using Microarray Gene Expression Data



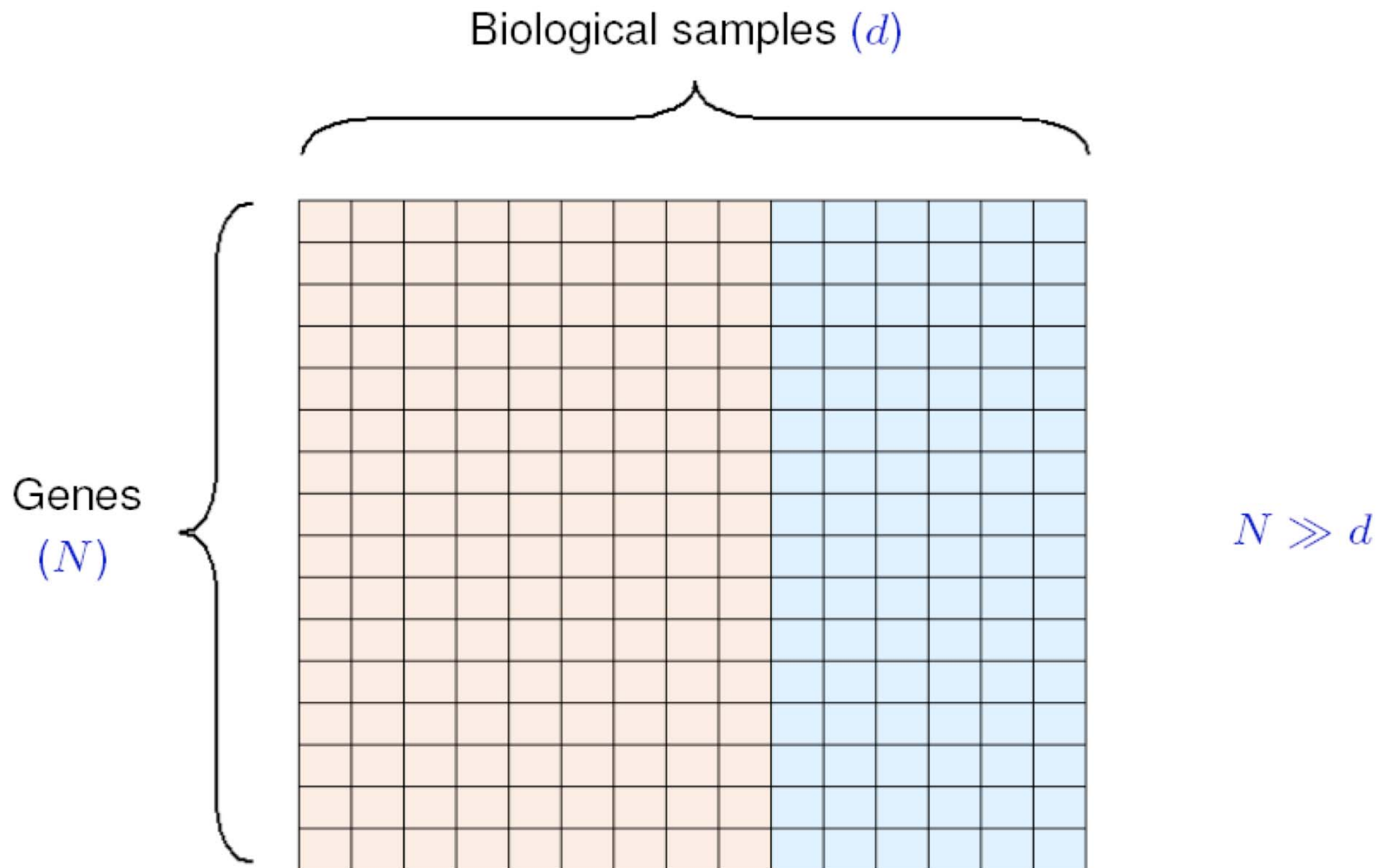
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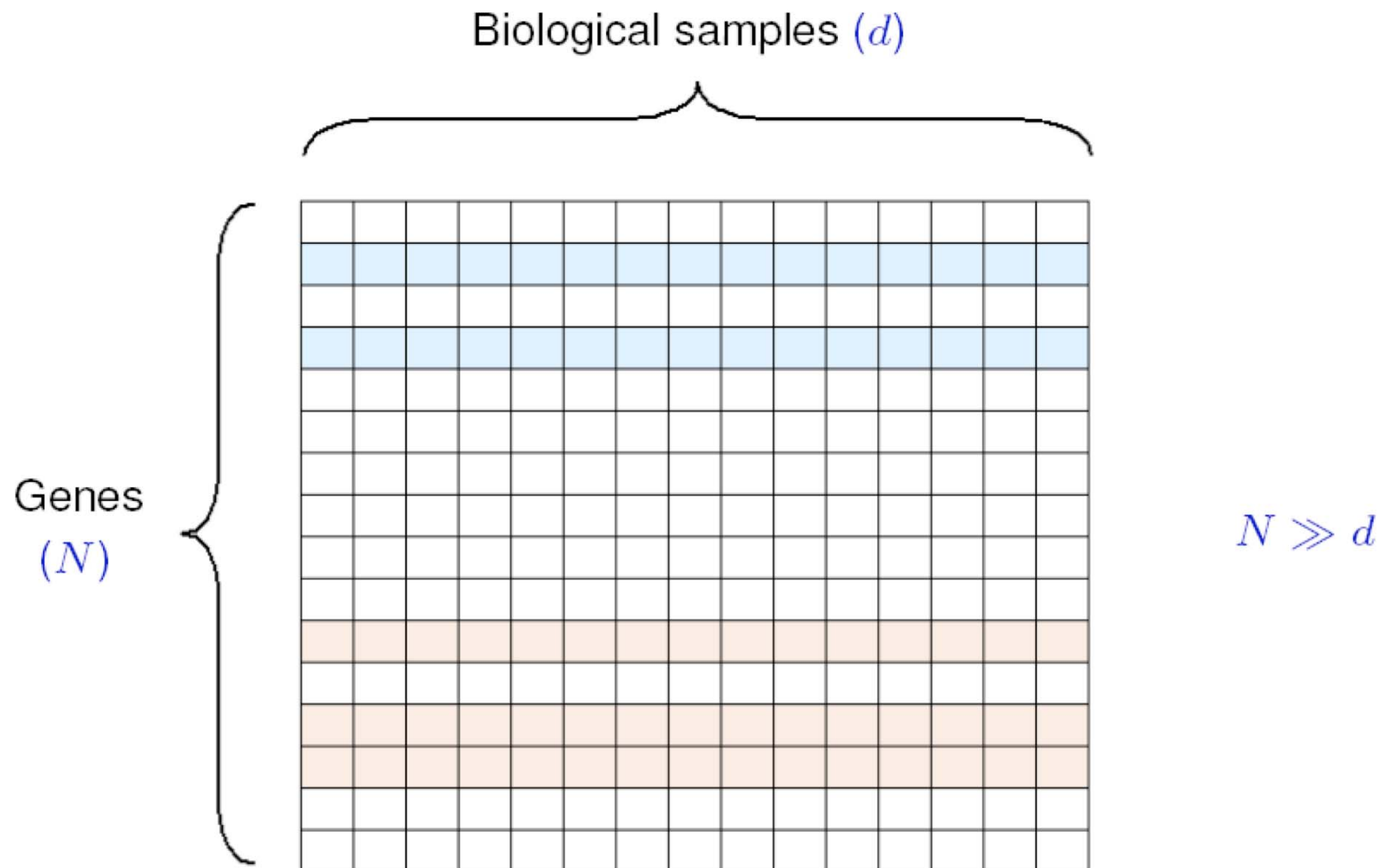
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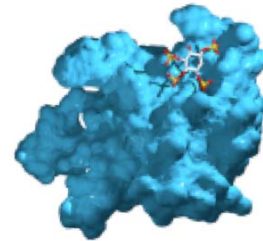
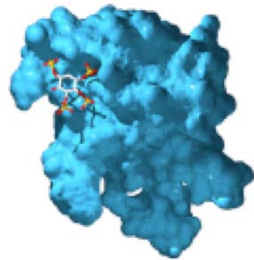


Identifying Genes Relevant to a Disease Using Microarray Gene Expression Data



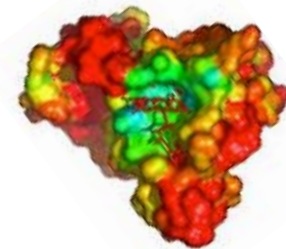
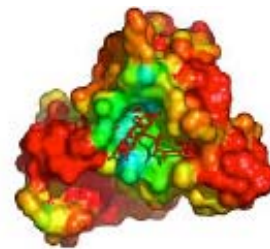
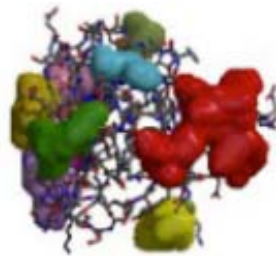
Formulation as a Bipartite Ranking Problem

Relevant



...

Not relevant

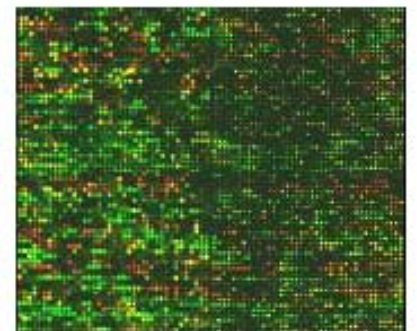


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Microarray Gene Expression Data Sets

[Golub et al, 1999; Alon et al, 1999]

Data Set	No. of Genes	No. of Tissue Samples	Notes
Leukemia	7129	72	25 AML / 47 ALL
Colon cancer	2000	62	40 tumor / 22 normal



Selection of Training Genes

Leukemia

Positive genes: Markers for AML/ALL

Myeloperoxidase
CD13
CD33
HOXA9 Homeo box A9
V-myb
CD19
CD10 (CALLA)
TCL1 (T cell leukemia)
C-myb
Deoxyhypusine synthase

Negative genes

157 genes involved in
physiological cellular functions

Colon cancer





Positive genes: Markers for colon cancer












Phospholipase A2
Keratin 6 isoform
PTP-H1
TF-III A
V-raf oncogene
MAPK kinase 1
CEA
Oncoprotein 18
PEP carboxykinase
ERK kinase 1

Negative genes

56 genes involved in
physiological cellular functions

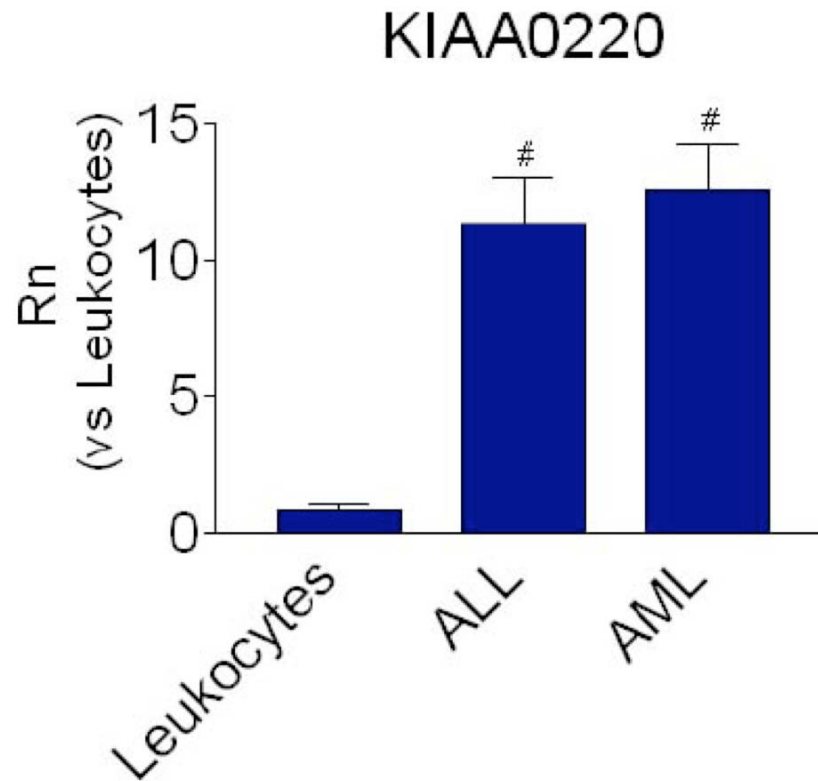
Top-Ranking Genes for Leukemia Returned by RankBoost

 Known marker;  Potential marker;
 Known therapeutic target;  Potential therapeutic target;
x No link found.

	Gene	Relevance Summary	t-Statistic Rank	Pearson Rank
1.	KIAA0220		6628	2461
2.	G-gamma globin		3578	3567
3.	Delta-globin		3663	3532
4.	Brain-expressed HHCPA78 homolog		6734	2390
5.	Myeloperoxidase		139	6573
6.	Disulfide isomerase precursor		6650	575
7.	Nucleophosmin		405	1115
8.	CD34		6732	643
9.	Elongation factor-1 β	x	4460	3413
10.	CD24		81	1
11.	60S ribosomal protein L23		1950	73
12.	5-aminolevulinic acid synthase		4750	3351

[Agarwal & Sengupta, 2009]

Biological Validation



[Agarwal et al, 2010]

Further Topics & Some Pointers

[Incomplete!]

- **Ranking performance measures that focus on accuracy at the top** [Yue et al, 2007; Clemencon & Vayatis, 2007; Cossock & Zhang, 2008; Rudin, 2009; Agarwal, 2010; also see IR ranking algorithms below]
- **Statistical consistency of ranking algorithms** [Clemencon & Vayatis, 2007; Clemencon & Vayatis, 2008; Cossock & Zhang, 2008; Duchi et al, 2010]
- **Other types of ranking problems, such as label ranking** [Crammer & Singer, 2003; Shalev-Shwartz & Singer, 2006] and **subset ranking** [Cossock & Zhang, 2008]
- **Ranking algorithms for information retrieval** [many, many recent papers; see Liu, 2009 for a survey]
- **Other applications of ranking, such as game move prediction** [Stern et al, 2007], **recommendation systems** [Stern et al, 2009], **manhole event prediction** [Rudin et al, 2010]