

Context sensitive information: Model validation by information theory

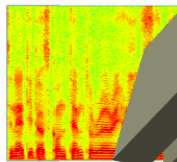
Joachim M. Buhmann

Computer Science Department, ETH Zurich

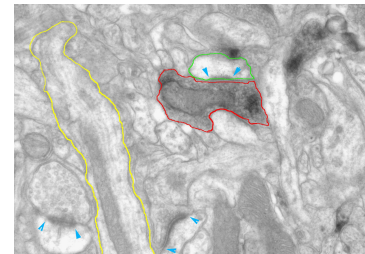


ETH Machine Learning Laboratory

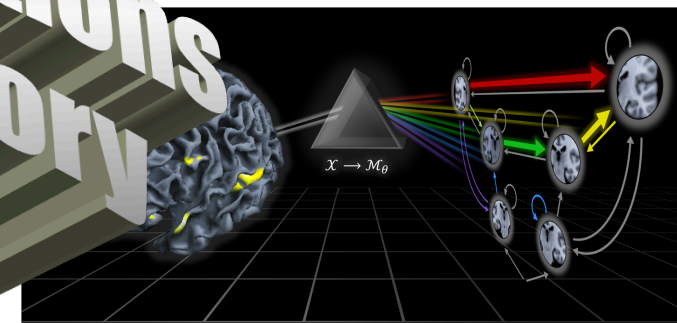
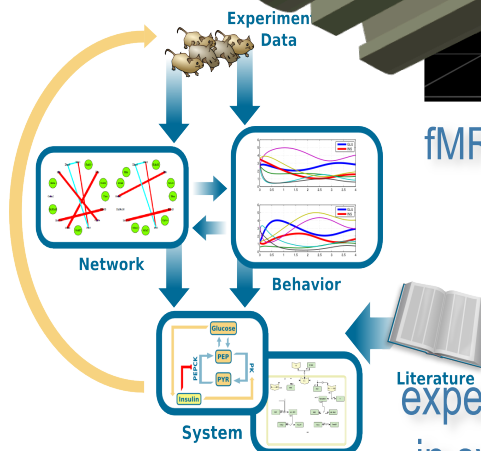
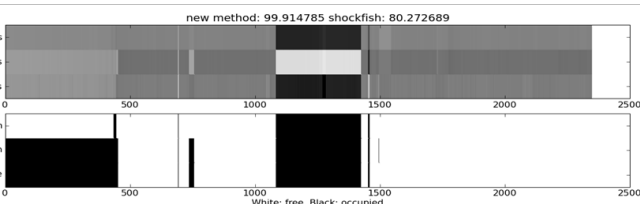
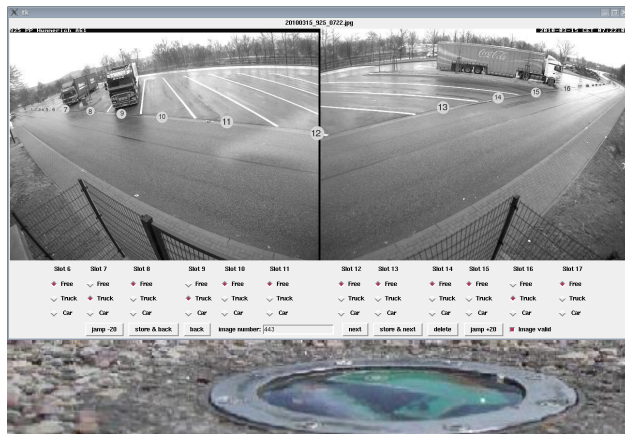
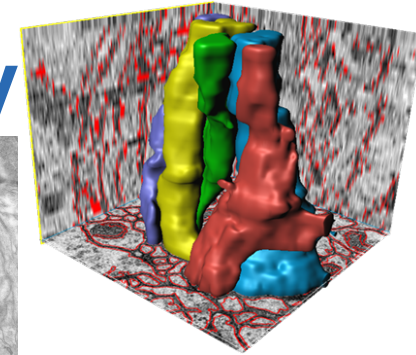
adaptive signal processing



Applications & Theory



neuron reconstruction



fMRI analysis for e.g. disease diagnosis

Literature
experimental design
in systems biology

The data deluge of the Zetta Byte (2^{70}) Age

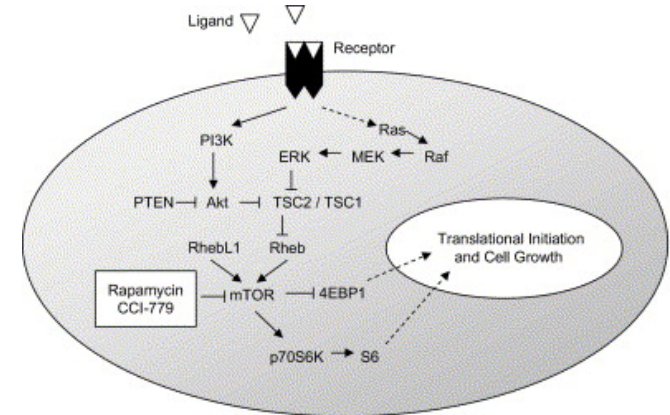
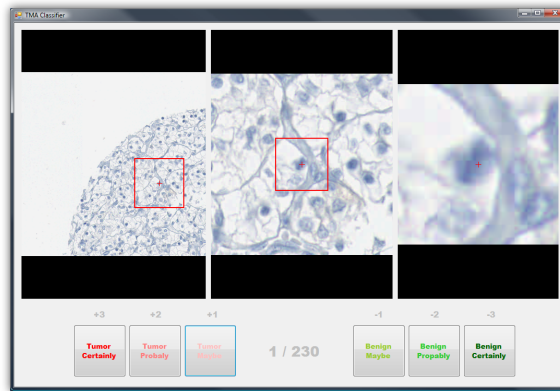
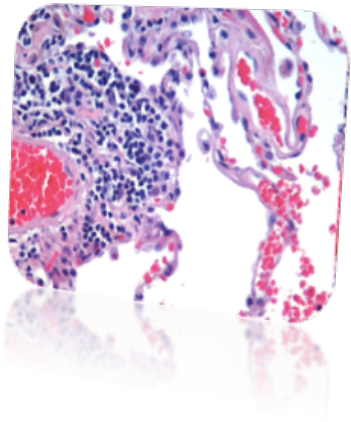
My scientific beliefs:

Mathematics as a language to describe reality has been invented for *prediction*

Model validation is more important than model design when algorithms generate models

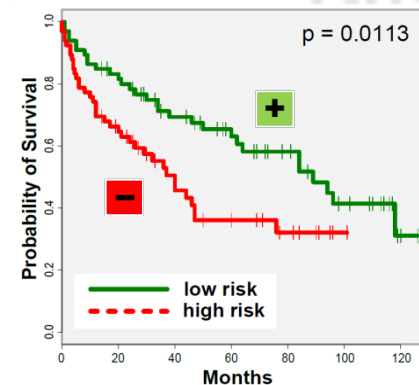
DATA

Modern information society and the Information Technology value chain



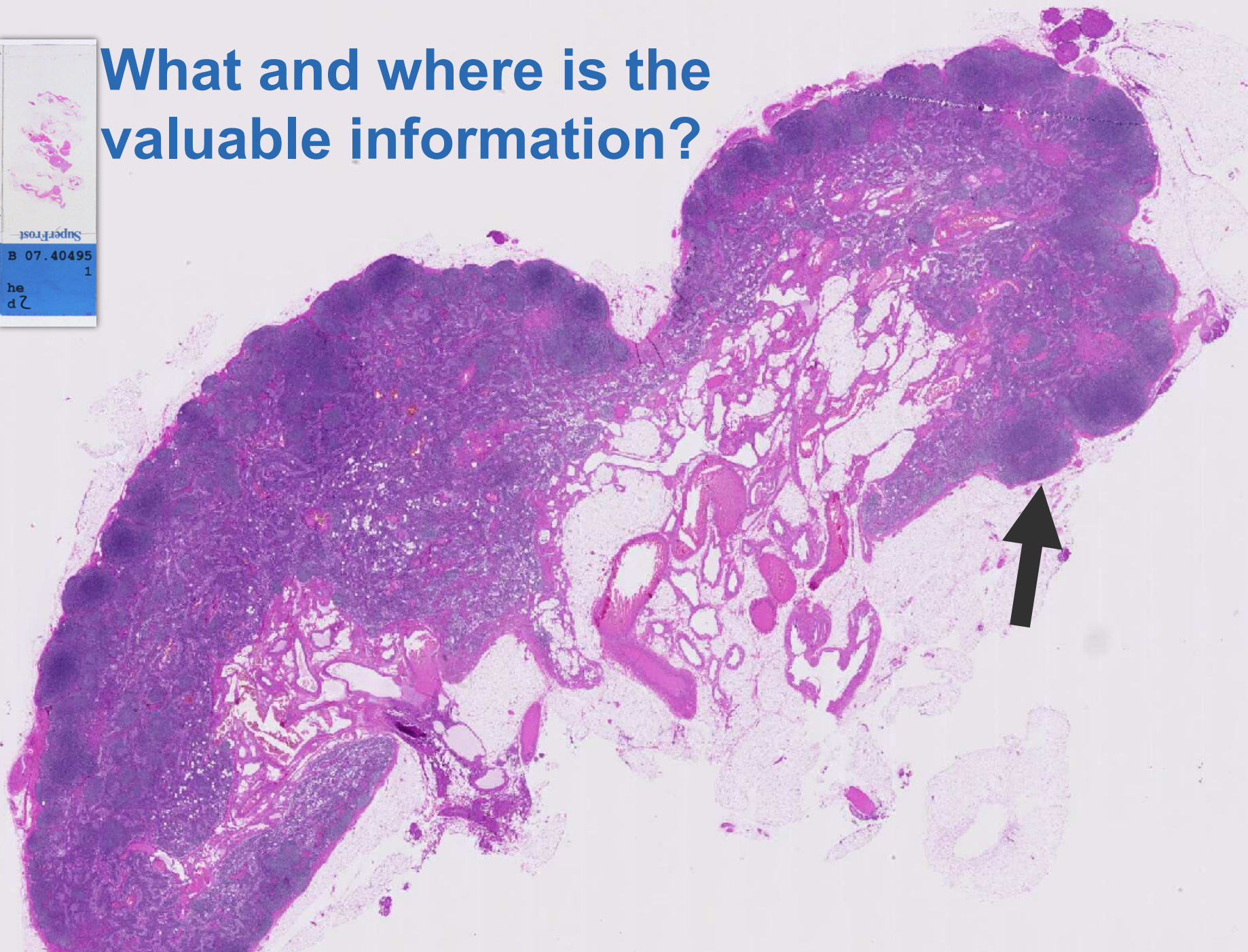
Activation of the mTOR Signaling Pathway in Renal Clear Cell Carcinoma. Robb et al., J Urology 177:346 (2007)

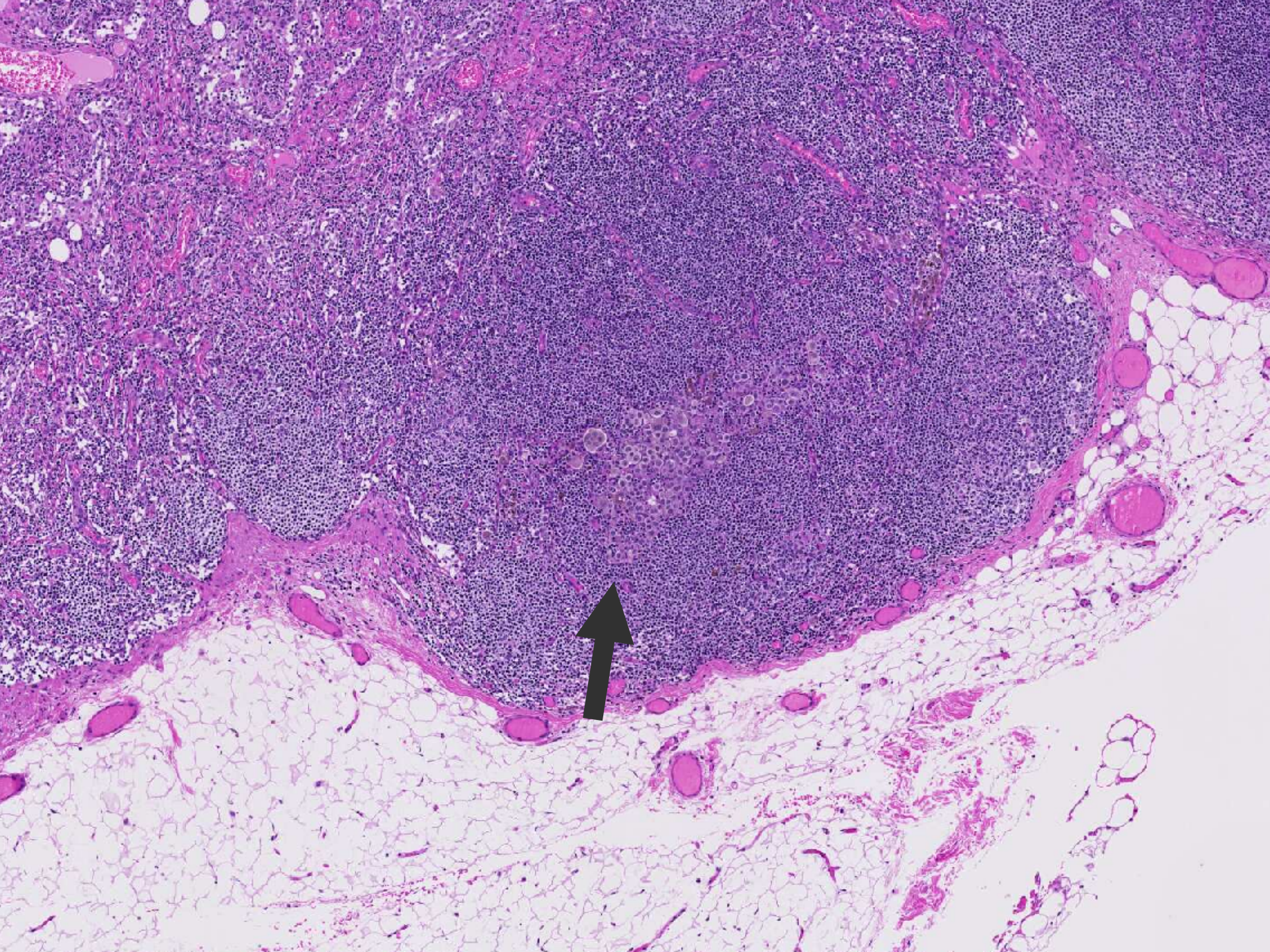
Data → **Information** → **Knowledge**

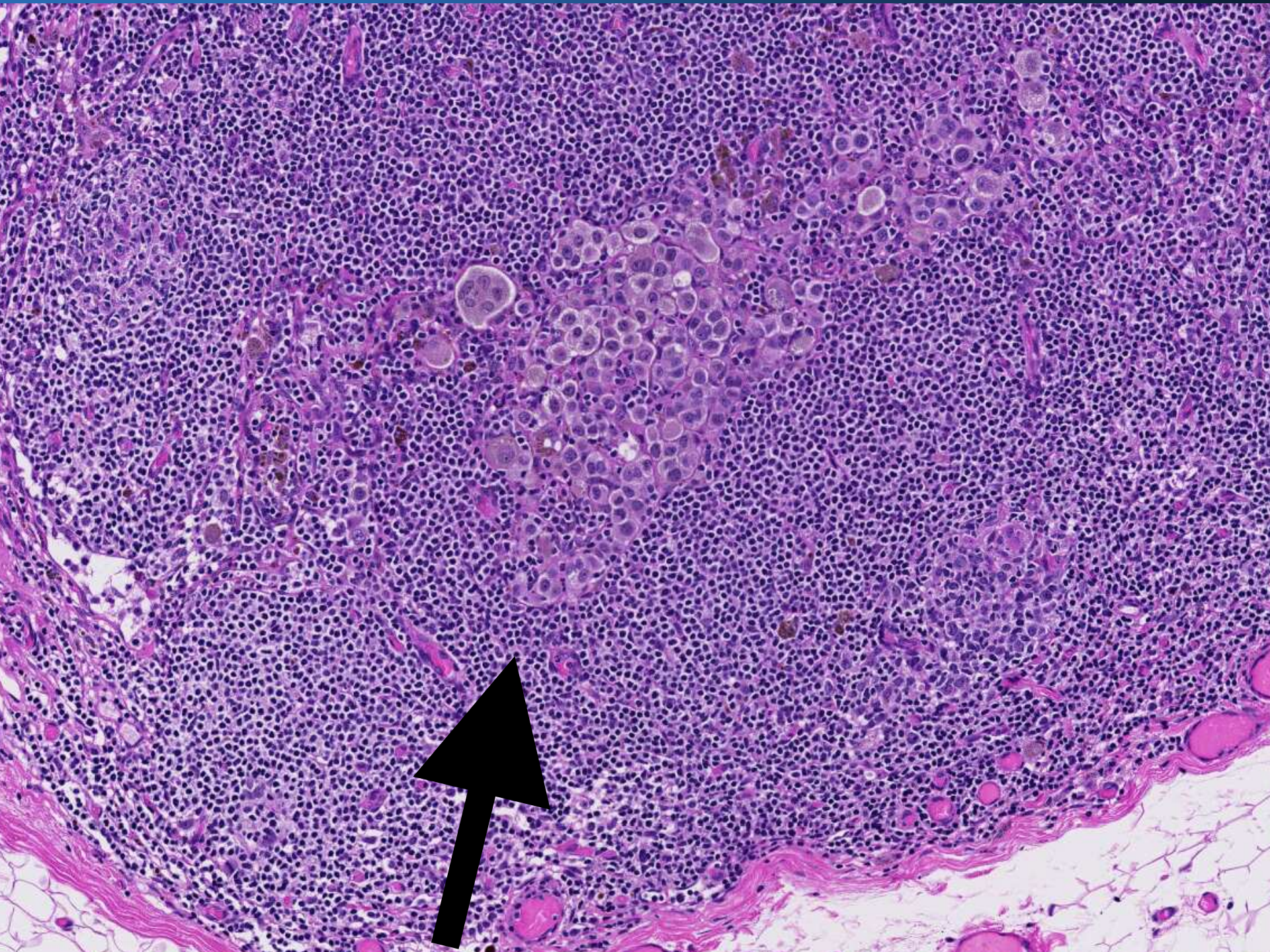


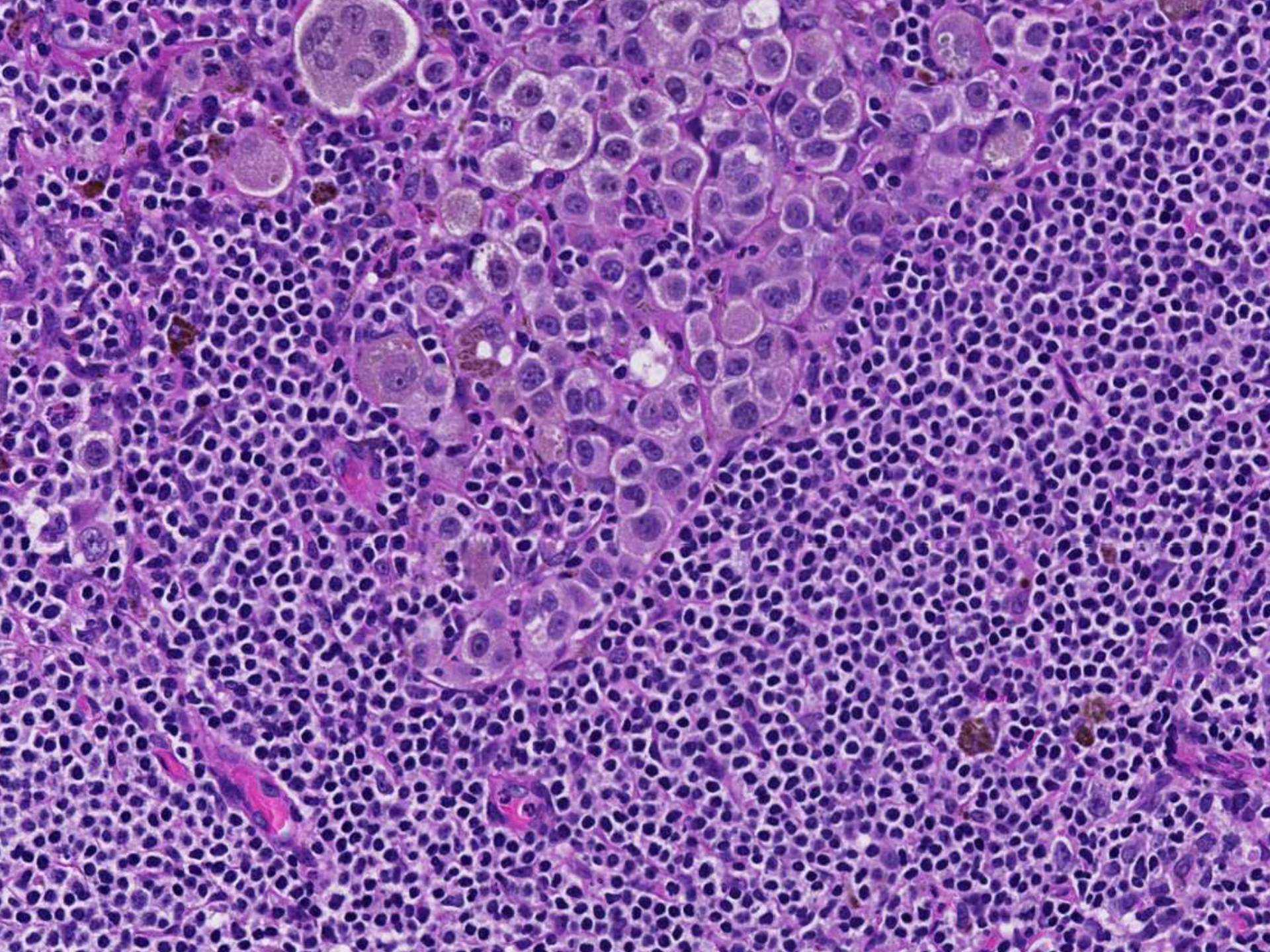
Value

What and where is the
valuable information?

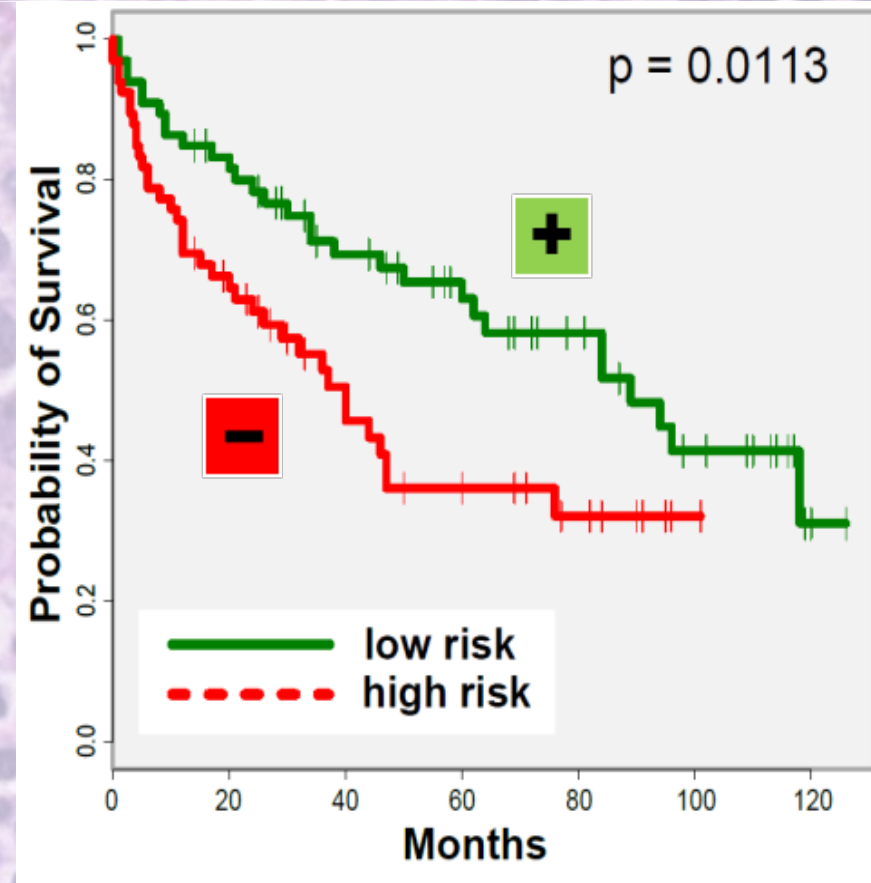
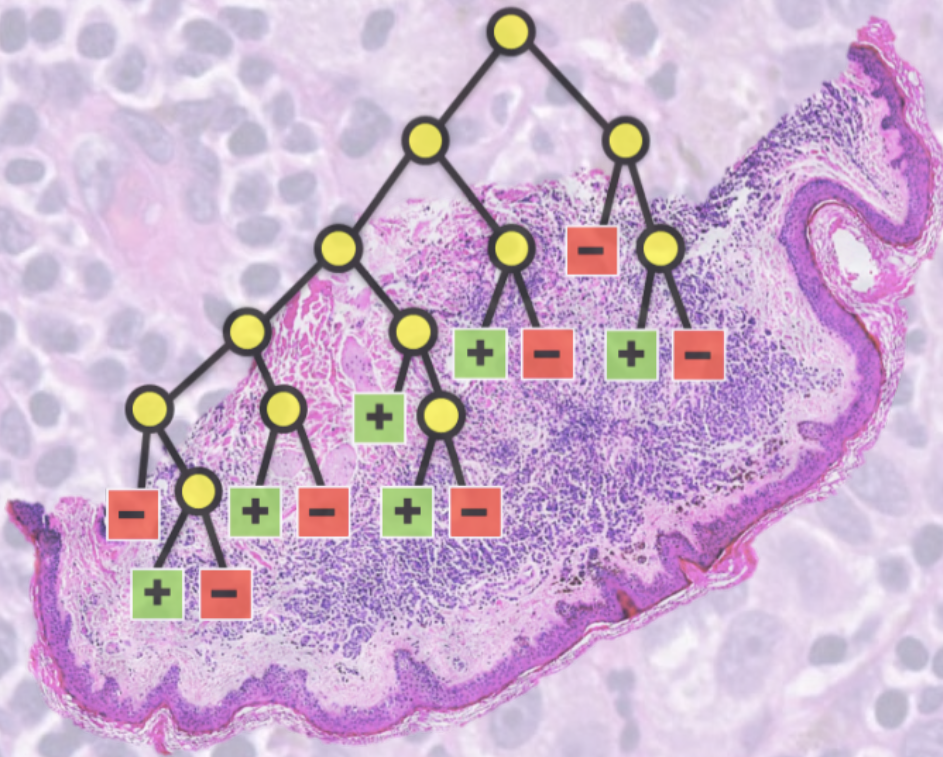






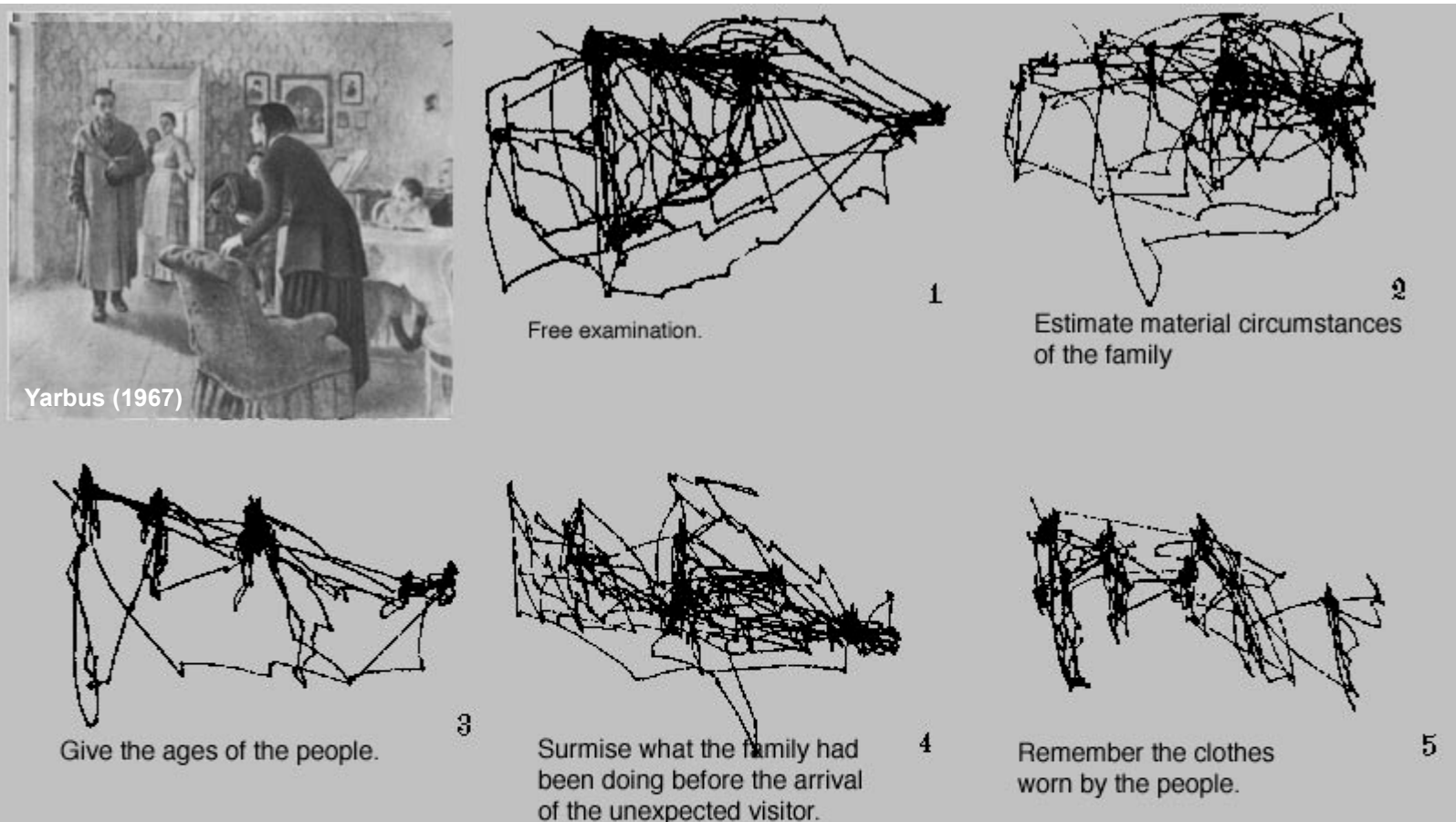


Survival risk: Where is this information?



Relevant information in data – where can we find it ?

The relevance of **information** is determined by the task !



Yarbus (1967)

1 Free examination.

2 Estimate material circumstances of the family

3 Give the ages of the people.

4 Surmise what the family had been doing before the arrival of the unexpected visitor.

5 Remember the clothes worn by the people.

Outline

- Relation between **information theory** and **machine learning**
- ***Context sensitive information***
- Coding by approximate optimization: ***Error free communication enables model selection!***
- Examples from cluster model selection
 - Validating spectral clustering
 - Approximate sorting

Scientific strategy for model validation

1. *Information processing under uncertainty*

(noisy inputs) yields indistinguishable solutions!

=> ***approximation sets***

2. **Approximation sets** quantize the solution space.

Consequently, **quantization creates symbols.**

3. **Symbols** define a **code** with an error rate.

=> **Optimality criterion:** Find the maximally informative symbols with zero error rate!

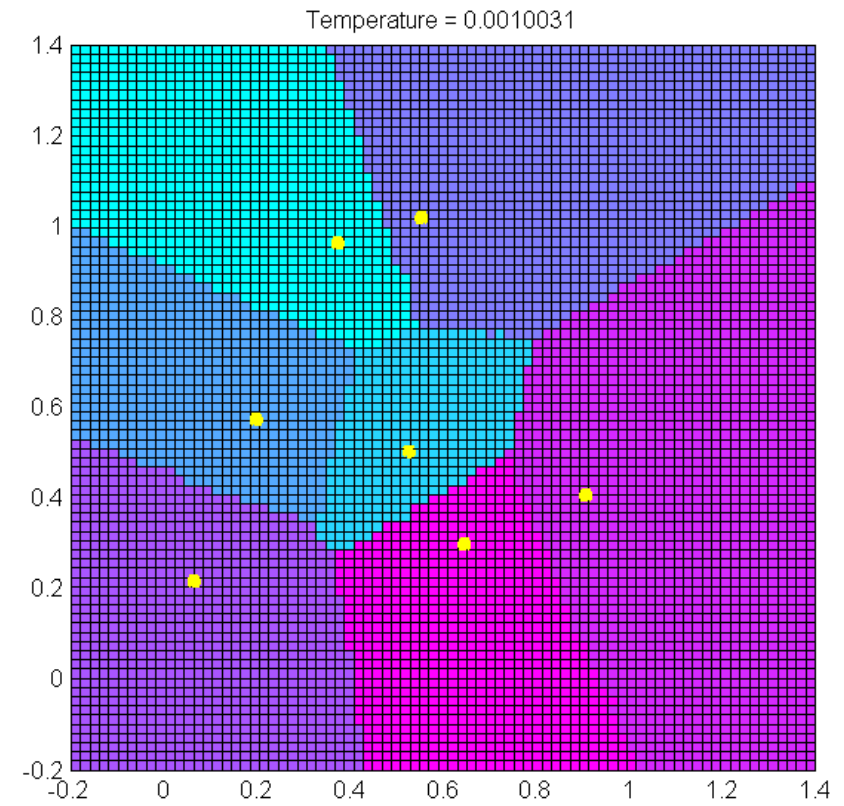
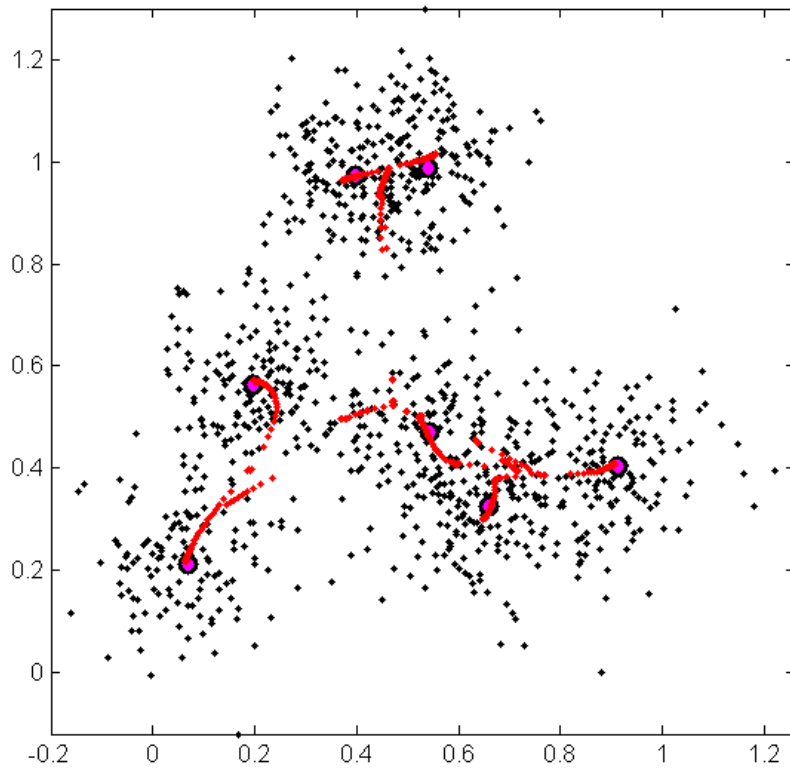
Context and task sensitive information!

- What do we look for?

Solutions out of a hypothesis class

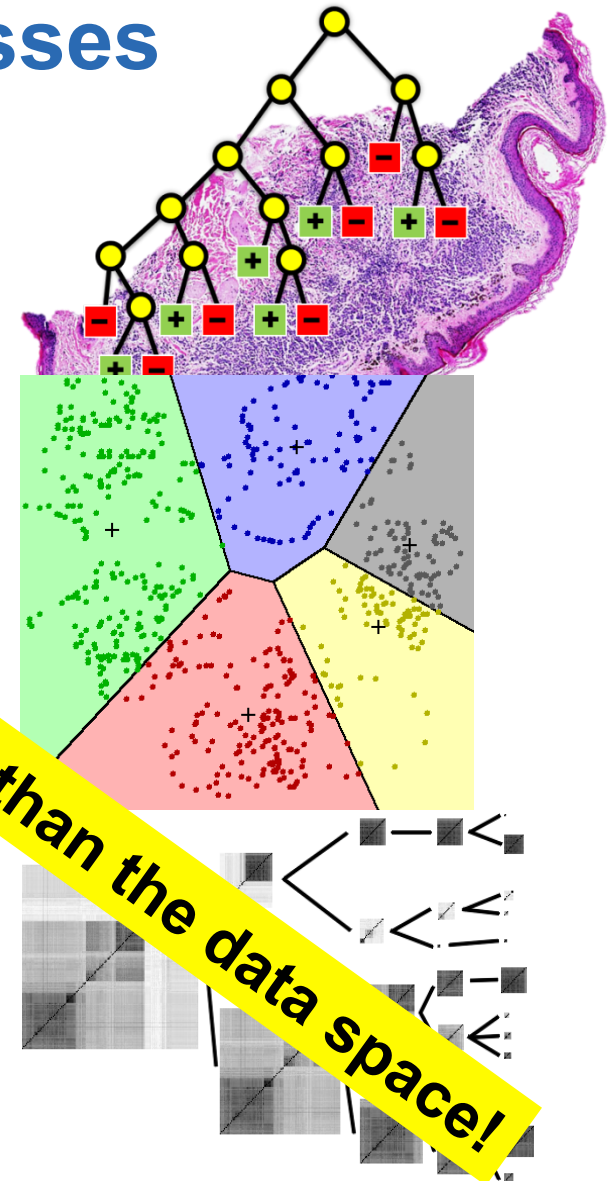
- *Select models* (cost functions) to rank hypotheses how well they interpret/explain data.
 - ***Validate*** the selected model(s)
- ⇒ ***Robust*** and ***expressive*** models are preferred over brittle ones! (***stability vs. informativeness***)
- ⇒ **# of stable bits = context sensitive information**

K-means hypotheses of increasing quality



Examples of hypothesis classes

- **Classifications**
- **Partitions or clusters:**
compactness/connectivity costs
- **Trees or dendrograms:**
partitions with ultrametricity;
Tree depth? # leaves?



Information Theory & Machine Learning

IT-Concepts

- Code book $\{c_1, \dots, c_n\} =$
hypothesis class

0000000	0100101	1001110	1100110
0001111	0101010	1001110	1100110
0010110	0110011	1010101	1100110
0011001	0111100	1011010	1111010

- Noisy **channel**

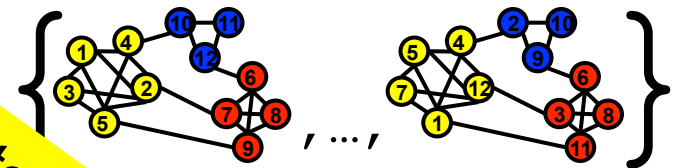
1100110 $\rightarrow$ 0100110

- Decoder: minimize
Hamming distance

$$|\underline{0}100110 - 1100110| < \\ |\underline{0}100110 - 1010101|$$

Pattern Recognition elements

- Set of approximation sets
(hypothesis class)



optimization problem



- Decoding by **approximate**
optimization of test instance

Approximate Optimization

- Define weights for hypotheses with low costs

$$w_{\beta}(c, \mathbf{X}) = \exp(-\beta R(c, \mathbf{X})) \quad c \in \mathcal{C}(\mathbf{X})$$

- Total weight of low cost hypotheses (partition fct.)

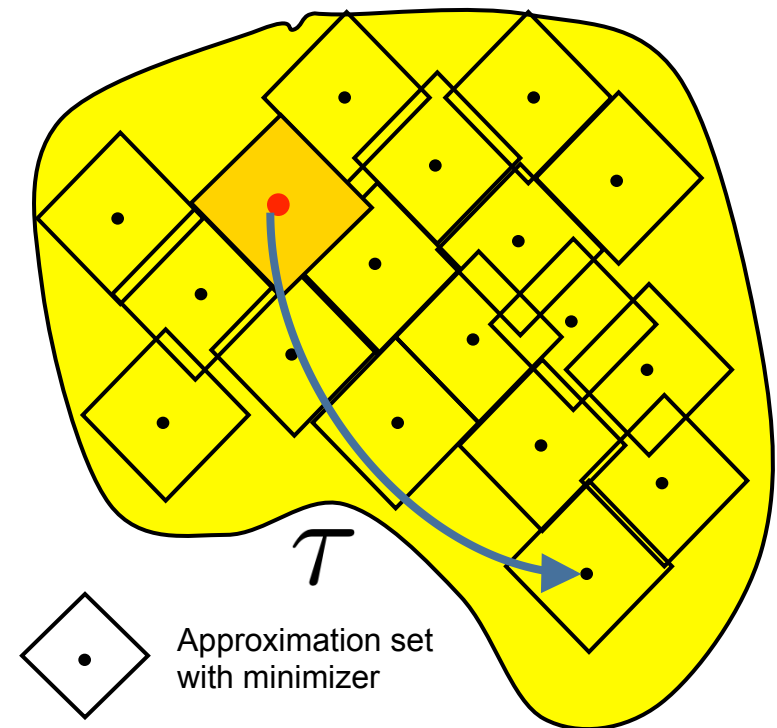
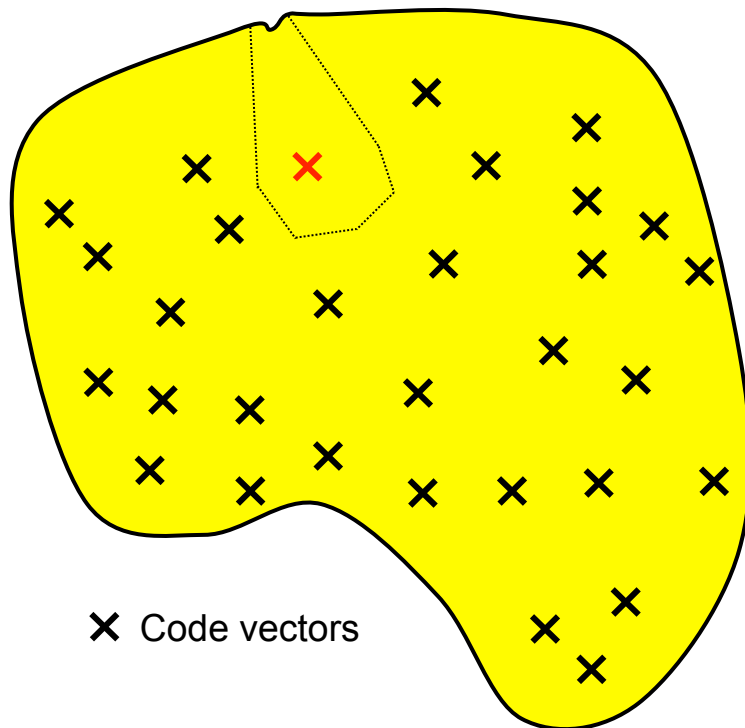
$$Z_{\beta}(\mathbf{X}) = \sum_{c \in \mathcal{C}(\mathbf{X})} \exp(-\beta R(c, \mathbf{X}))$$

- Special case: **Approximation set**

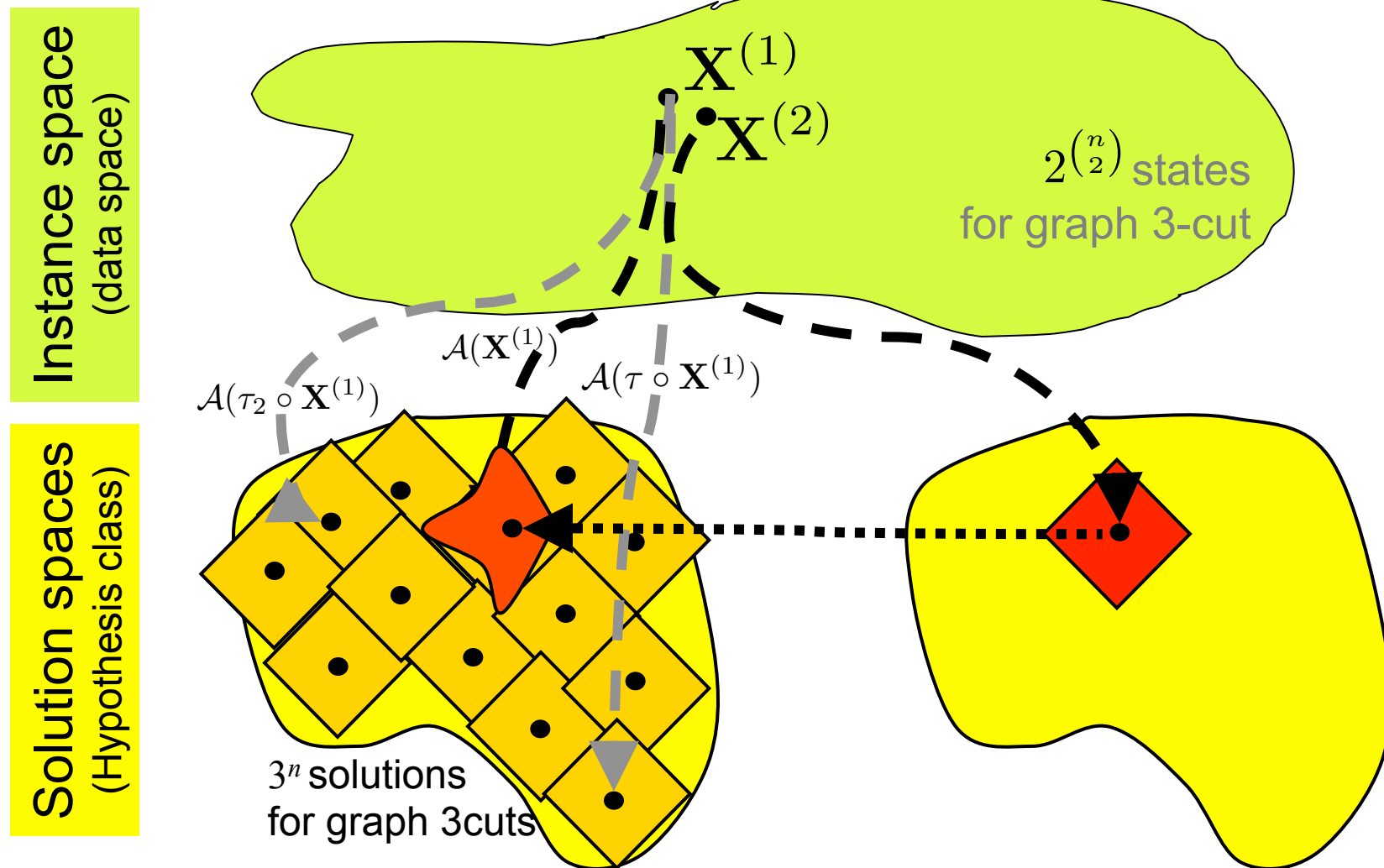
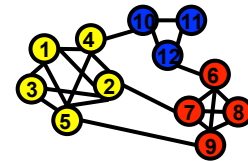
$$w_{\beta}(c, \mathbf{X}) = \begin{cases} 1, & R(c, \mathbf{X}) \leq R(c^{\perp}, \mathbf{X}) + 1/\beta; \\ 0, & \text{otherwise.} \end{cases}$$

Code problems define approximation sets

- IT: **Space of strings** is partitioned by code vectors
- ML: **Hypothesis class** is partitioned by code problems

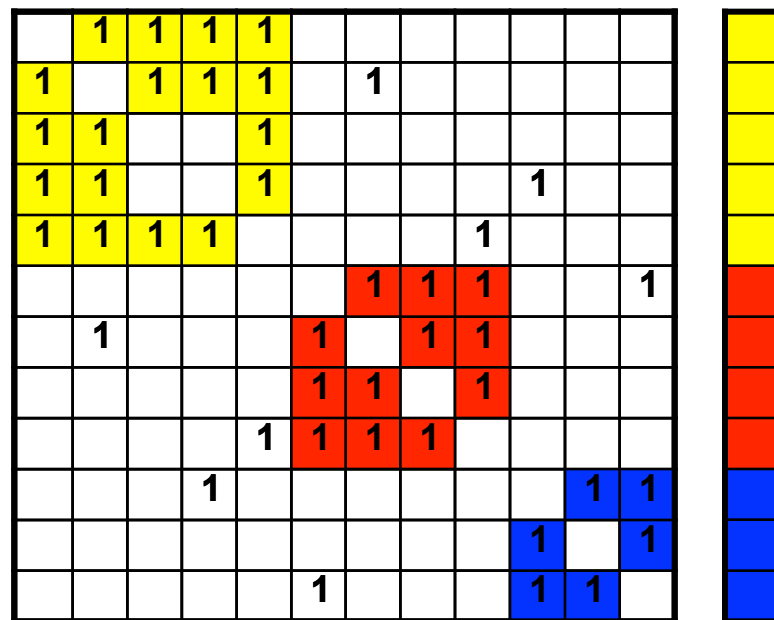
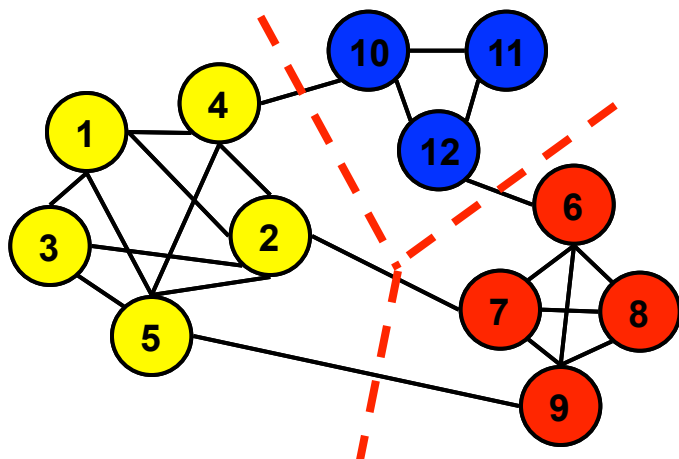


Generalization in optimization

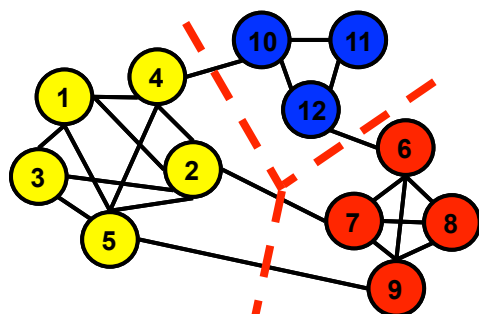
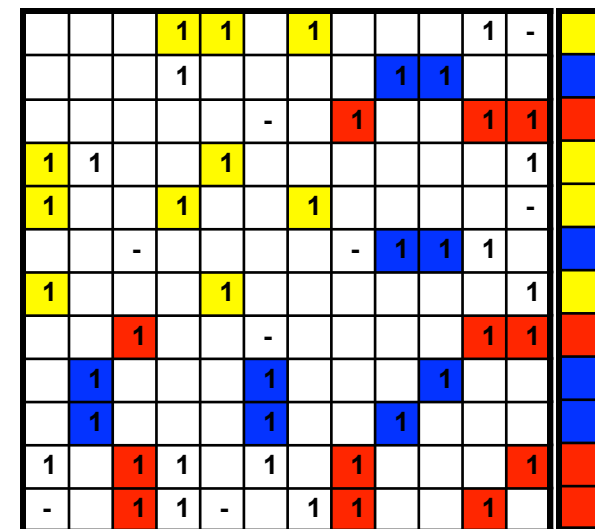
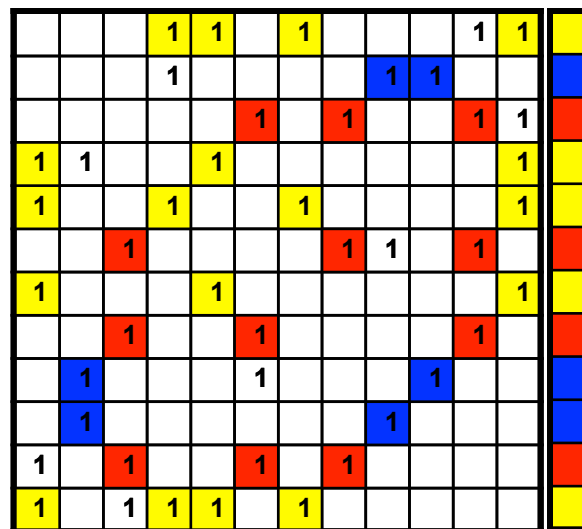
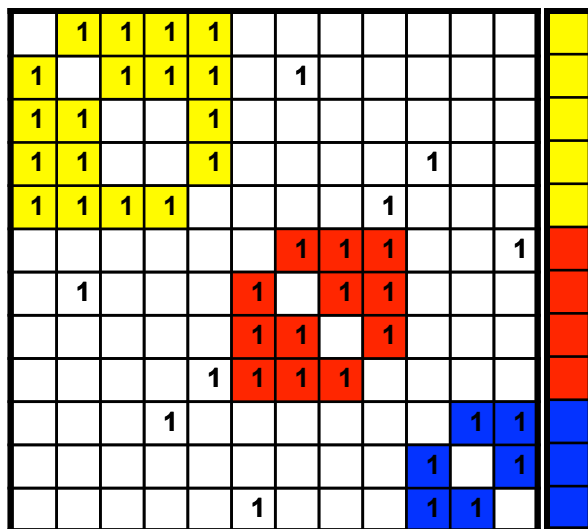
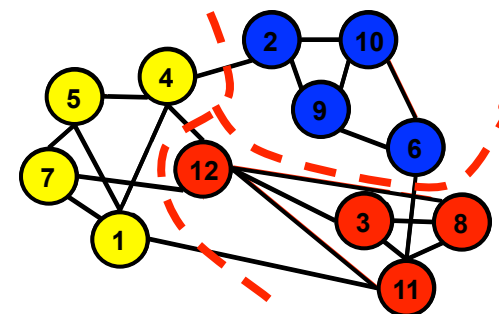
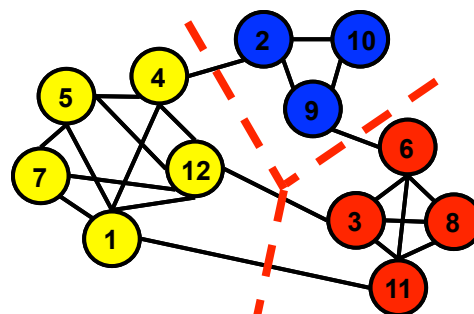


Graph Cut - Clustering in three groups

- **Graph representation:** **vertices** denote **objects**
edges express (dis)similarities
- **Hypothesis class:** all **3-cuts** of a graph



graph cut *code* problems


$$\dots$$


Coding by Code Problems

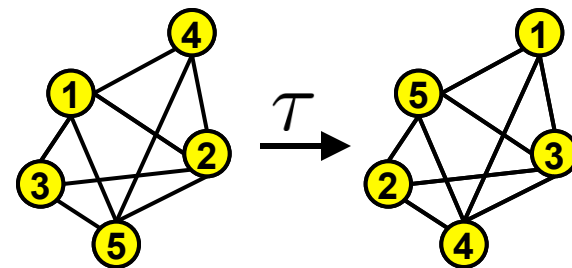
- **Idea:** define a **code** by transforming a given optimization problem with a **transformation set**

$$\mathcal{T} = \left\{ \tau : R(c, \mathbf{X}) = R(\tau \circ c, \tau_{\mathbf{X}} \circ \mathbf{X}) \right\}$$

⇒ Codebook $\mathbb{T} = \{ \tau_i \in \mathcal{T} : 1 \leq i \leq 2^{n\rho} \}$

Combinatorial optimization:

permutation of vertices in a graph



- **Identifiable transformations $\mathcal{T}$ are the messages!**

Coding with Graph Cut approximation sets

define a set of
code problems

problem generator PG

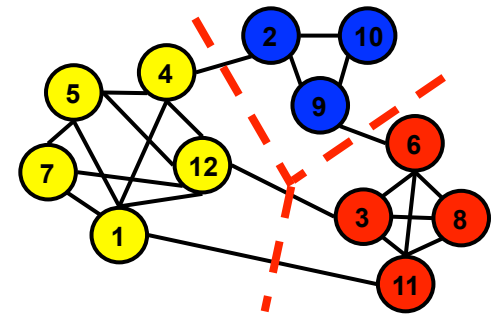
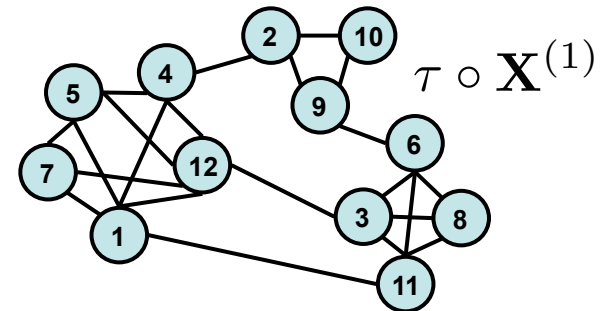
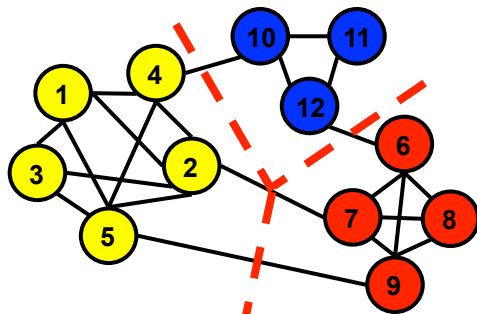
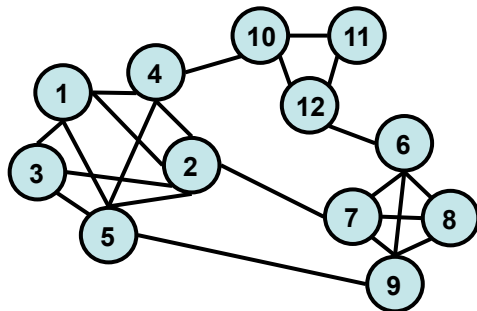
$R(\cdot, \mathbf{X}^{(1)})$

$R(\cdot, \mathbf{X}^{(1)})$

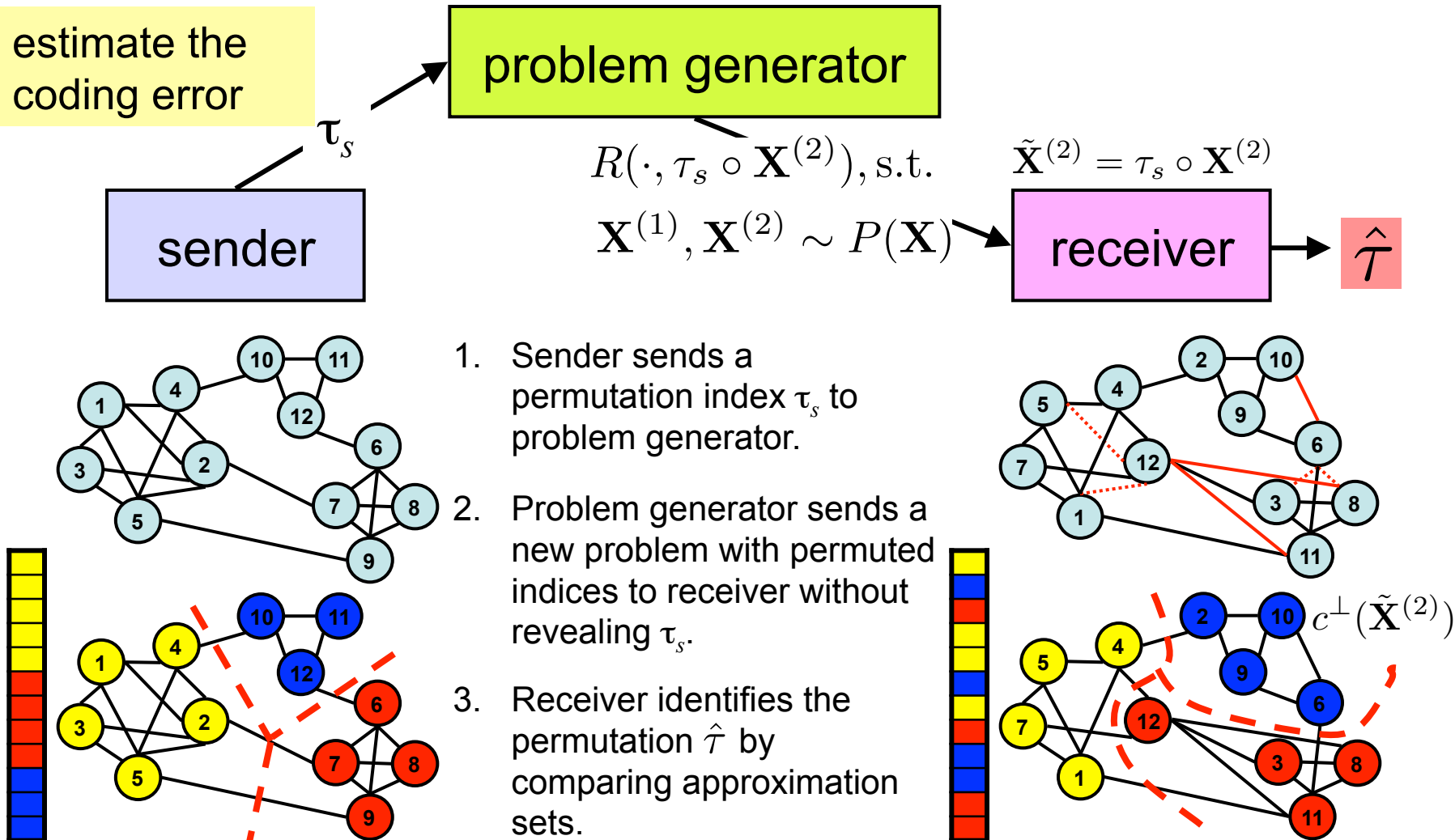
sender

receiver

$\{\tau_1, \dots, \tau_{2^{n\rho}}\}$



Communication by approximation sets



Communication Process

- Receiver **compares sets of hypothesis weights**
- **Decoding by maximizing weight overlap**

$$\hat{\tau} := \arg \max_{\tau} \# (\text{yellow diamond}_{\tau} \cap \text{red star})$$

- **Error event:** $\hat{\tau} \neq \tau_s$
- Calculate $\lim_{n \rightarrow \infty} P(\hat{\tau} \neq \tau_s | \tau_s, \mathbf{X}^{(1)}, \mathbf{X}^{(2)}) = 0$

Error Probability

- Estimate conditional error for data $\mathbf{X}^{(1)}, \mathbf{X}^{(2)}$ given a set of random transformations $\tau \in \mathcal{T}$

$$\begin{aligned}
 P\left(\hat{\tau} \neq \tau_s | \tau_s, \mathbf{X}^{(1,2)}\right) &= P\left(\max_{j \neq s} Z_{\beta,j}^{(1\&2)} > Z_{\beta,s}^{(1\&2)} | \tau_s, \mathbf{X}^{(1,2)}\right) \\
 &\stackrel{\text{Union bound}}{\leq} \sum_{j \neq s} P\left(Z_{\beta,j}^{(1\&2)} > Z_{\beta,s}^{(1\&2)} | \tau_s, \mathbf{X}^{(1,2)}\right) \\
 &\leq 2^{n\rho} P\left(Z_{\neq s}^{(1\&2)} > Z_s^{(1\&2)} | \tau_s, \mathbf{X}^{(1,2)}\right) \\
 &\stackrel{\text{Markov ineq.}}{\leq} 2^{n\rho} \frac{\mathbb{E}_{\tau \neq s} Z_{\neq s}^{(1\&2)}}{Z_s^{(1\&2)}} \leq 2^{n\rho} \frac{Z_{\beta}^{(1)} Z_{\beta}^{(2)}}{|\mathcal{T}| Z_s^{(1\&2)}}
 \end{aligned}$$

Asymptotical error-free communication

$\lim_{n \rightarrow \infty} P(\hat{\tau} \neq \tau_s | \tau_s, \mathbf{X}^{(1)}, \mathbf{X}^{(2)}) = 0$ **is possible if ...**

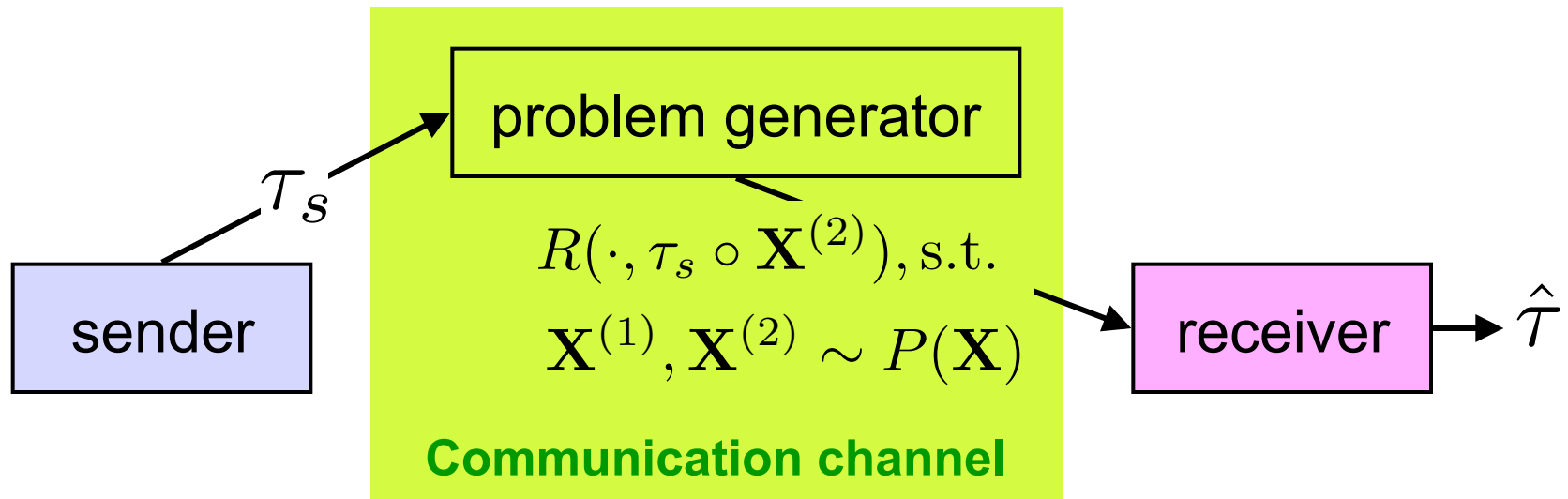
- ... rate ρ is bounded by mutual information $\hat{\mathcal{I}}_\beta(\tau_s, \hat{\tau})$

$$\begin{aligned} \rho < \hat{\mathcal{I}}_\beta(\tau_s, \hat{\tau}) &\equiv \frac{1}{n} \log_2 \frac{|\mathcal{T}| Z_\beta^{(1\&2)}}{Z_\beta^{(1)} Z_\beta^{(2)}} \\ &= \frac{1}{n} \left(\log_2 \frac{|\mathcal{T}|}{Z_\beta^{(1)}} + \log_2 \frac{|\mathcal{C}^{(2)}|}{Z_\beta^{(2)}} - \log_2 \frac{|\mathcal{C}^{(2)}|}{Z_\beta^{(1\&2)}} \right) \end{aligned}$$

Bound calculation involves
partition functions for individual
and joint costs

$$\begin{aligned} Z_\beta^{(\nu)} &= \sum_{c \in \mathcal{C}(\mathbf{X}^{(\nu)})} \exp(-\beta R(c, \mathbf{X}^{(\nu)})), \quad \nu = 1, 2 \\ Z_\beta^{(1\&2)} &= \sum_{c \in \mathcal{C}(\mathbf{X}^{(1)})} \exp\left(-\beta(R(c, \mathbf{X}^{(1)}) + R(c, \mathbf{X}^{(2)}))\right) \end{aligned}$$

Model Selection by Maximization of Approximation Capacity



- Optimize the communication channel w.r.t. precision γ (β) and cost function $R(.,.)$
- Maximize Mutual Information

ASC for binary channel consistent with Shannon information theory

- Hypothesis class: set of binary strings $\xi^{(1)}, \xi^{(2)} \in \{-1, 1\}^n$

- Communication:

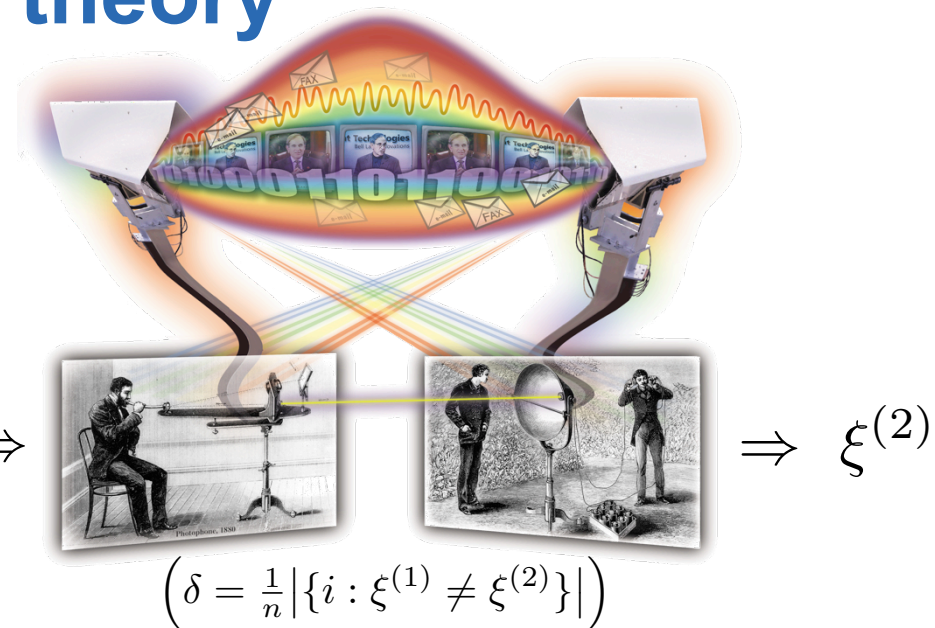
- Costs of string s : Hamming distance

$$R(s, \xi^{(1)}) = \sum_{i=1}^n \mathbb{I}_{\{s_i \neq \xi_i^{(1)}\}}$$

- Mutual information:

$$\mathcal{I}_\beta = \ln 2 + (1 - \delta) \ln \cosh \beta - \ln(\cosh \beta + 1)$$

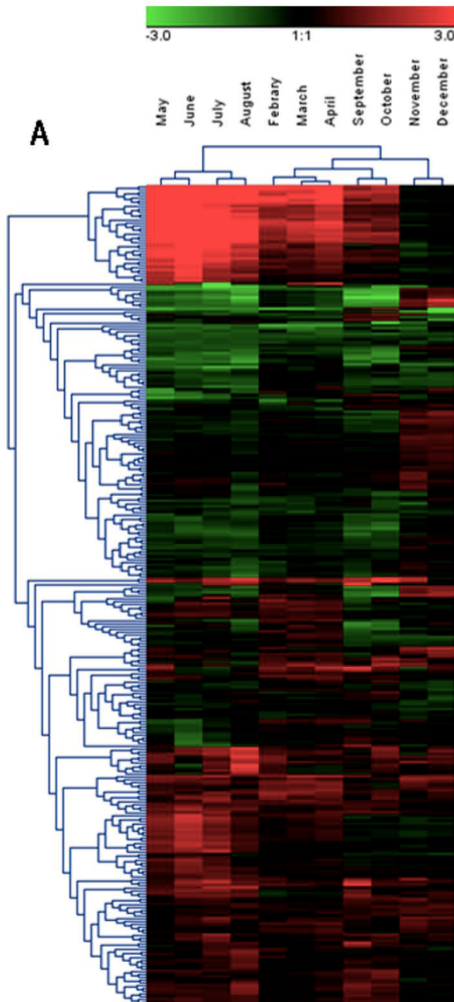
$$\text{for } (*) \quad \frac{d\mathcal{I}_\beta}{d\beta} = 0 \quad \stackrel{(*)}{=} \quad \ln 2 + (1 - \delta) \ln(1 - \delta) + \delta \ln \delta$$



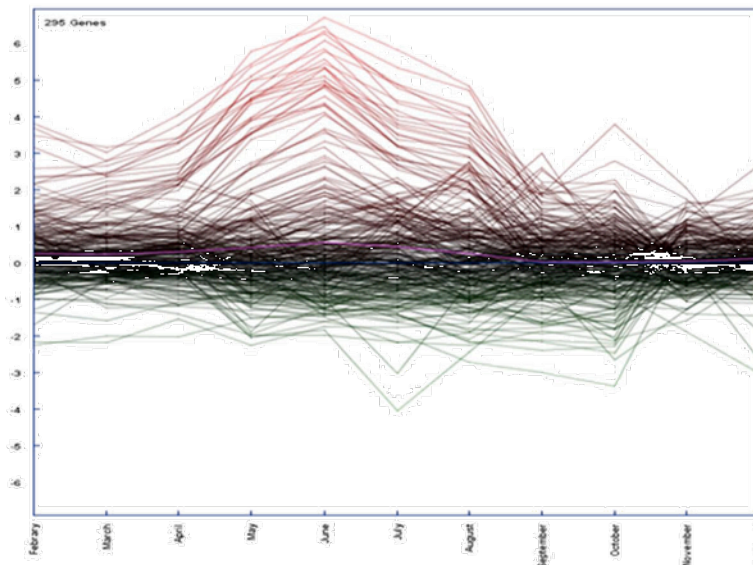
Channel capacity of BSC

Validation of relational clustering

Joint work with
M. H. Chehreghani & A.G. Busetto

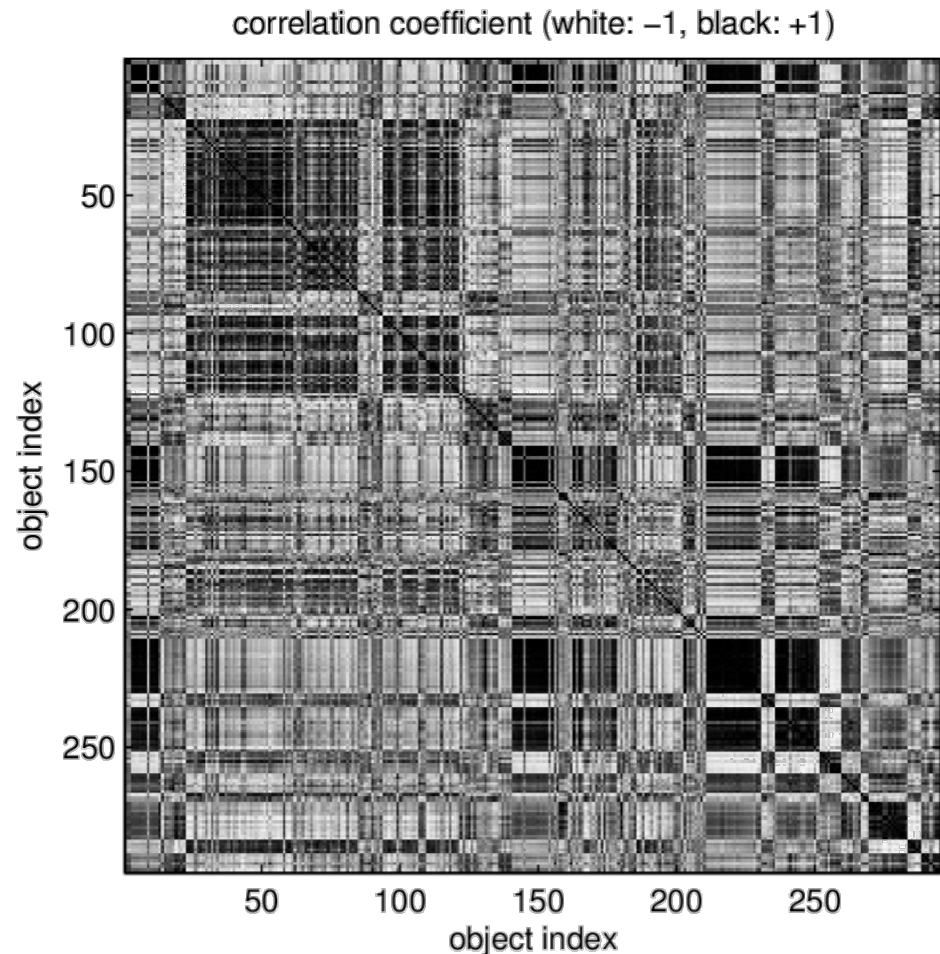


Gene Expression Rhythms (A) in the mussel *mytilus galloprovincialis* (Lam.) across an Annual Cycle



Correlations between gene expressions

- **Gene expression time series** are compared by Pearson correlation coefficient
- Relational clustering by *normalized Cut* (Ncut), *correlation clustering* (CC) or *pairwise clustering* (PC)



Validating Relational clustering

- Cost function for pairwise clustering

$$R^{pc}(c, \mathbf{X}) = - \sum_{k=1}^K \mathcal{G}_k \sum_{(i,j) \in \mathcal{E}_{kk}} \frac{X_{ij}}{|\mathcal{E}_{kk}|}$$

- Normalized cut

$$R^{nc}(c; W) = \sum_{v \leq k} \left(\frac{\text{cut}(\mathcal{G}_v, \mathcal{V} \setminus \mathcal{G}_v)}{\text{assoc}(\mathcal{G}_v, \mathcal{V})} \right)$$

- Correlation clustering

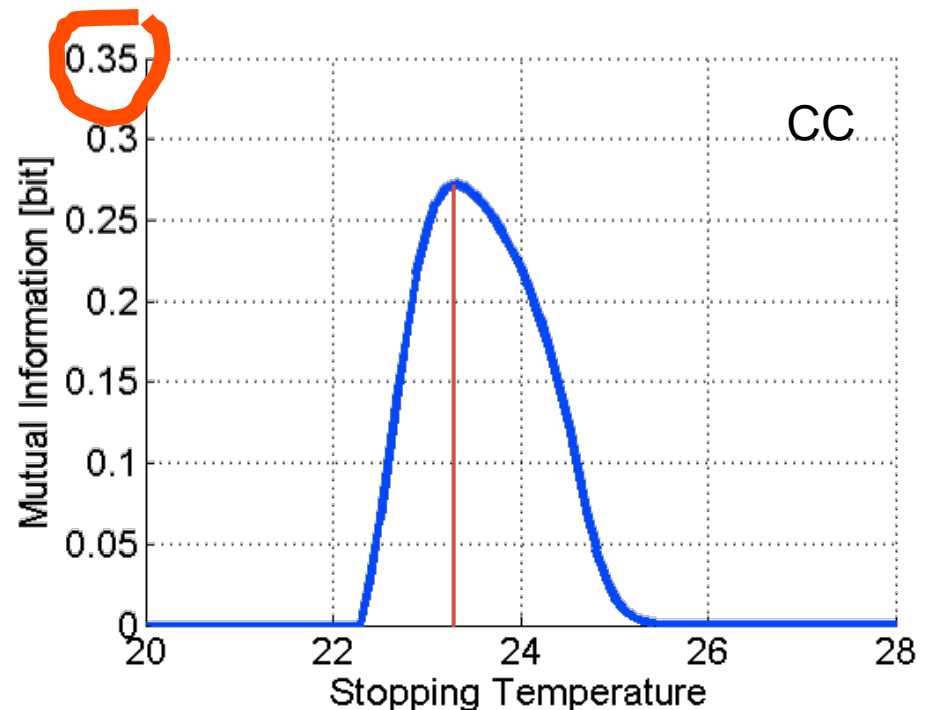
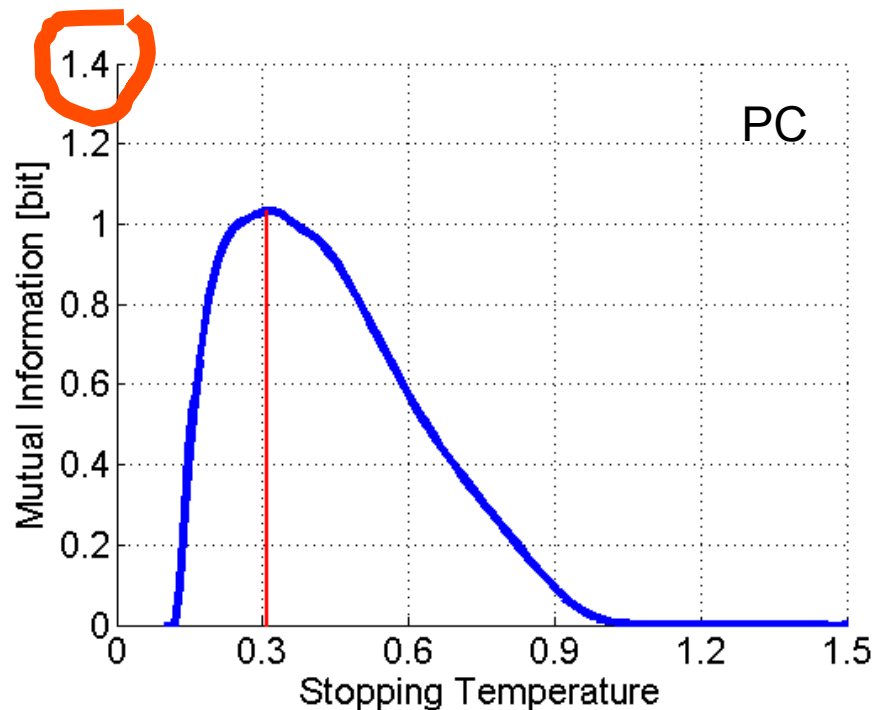
$$R^{cc}(c, \mathbf{X}) = \frac{1}{2} \sum_{k,l=1}^K \sum_{(i,j) \in \mathcal{E}_{kl}} \left((-1)^{\delta_{kl}} X_{ij} + |X_{ij}| \right)$$

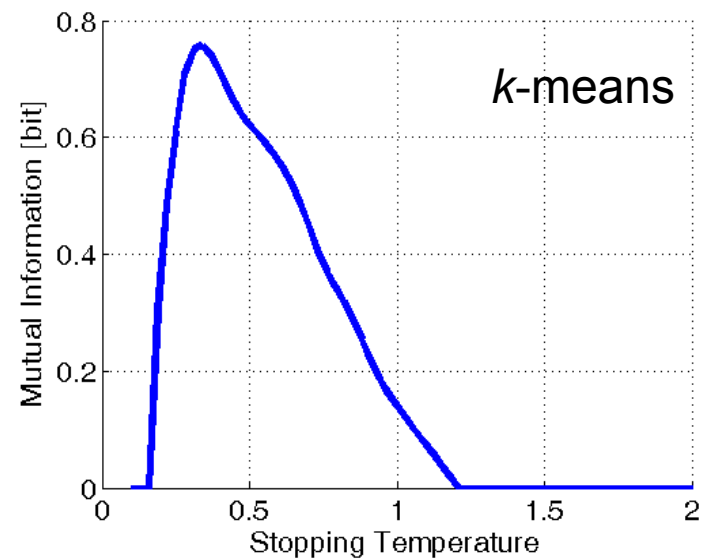
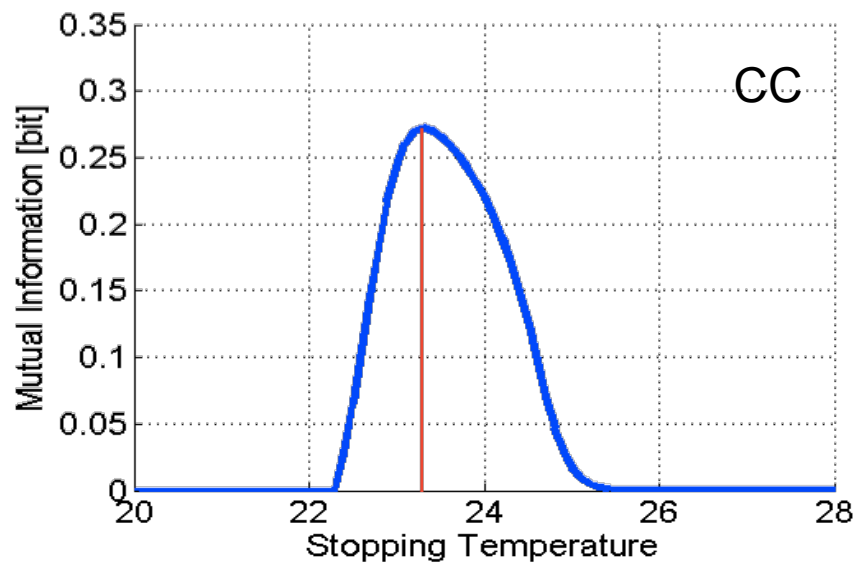
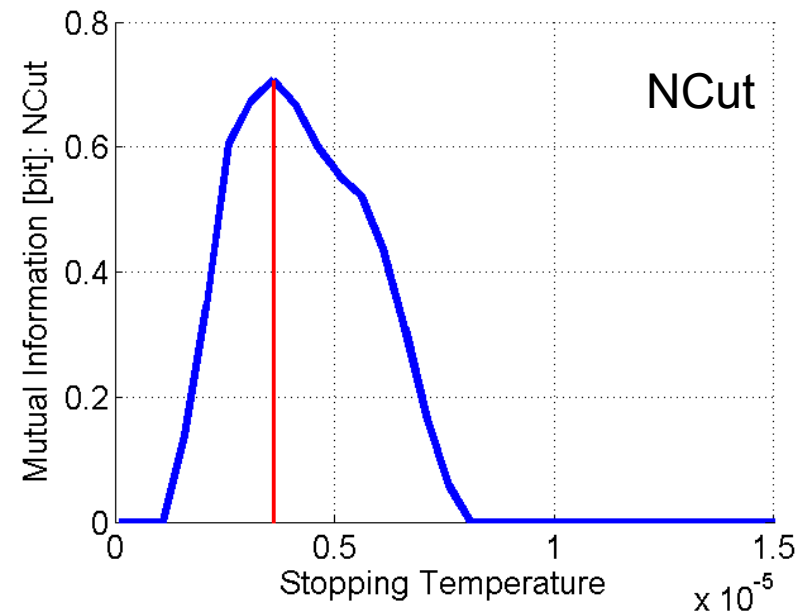
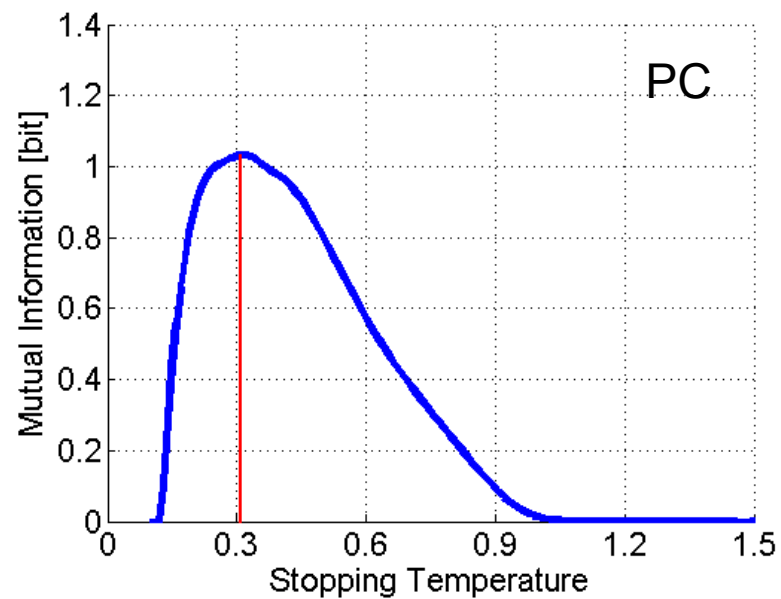
Defs.: $\mathcal{G}_k = \{i : c(i) = k\}$, $\mathcal{E}_{kl} = \{(i,j) : c(i) = k \wedge c(j) = l\}$

Information of Spectral Clusterings

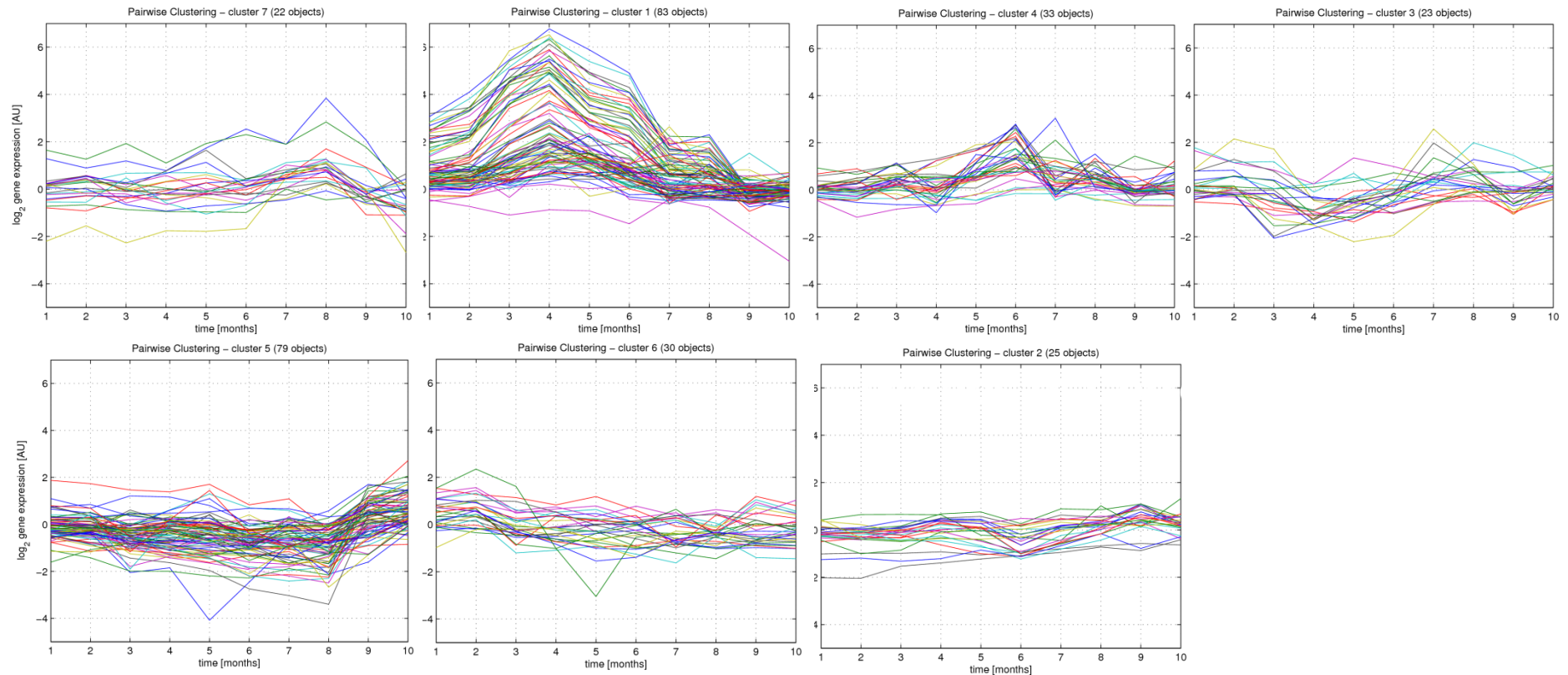
$$\hat{\mathcal{I}}_{\beta}(\tau_s, \hat{\tau}) \equiv \frac{1}{n} \log_2 \frac{|\mathcal{T}| Z_{\beta}^{(1\&2)}(\mathbf{X}^{(1)}, \mathbf{X}^{(2)})}{Z_{\beta}(\mathbf{X}^{(1)}) Z_{\beta}(\mathbf{X}^{(2)})}$$

- Pairwise Clustering extracts 30% more information than NCut and 3.5 times more than Correlation Clustering



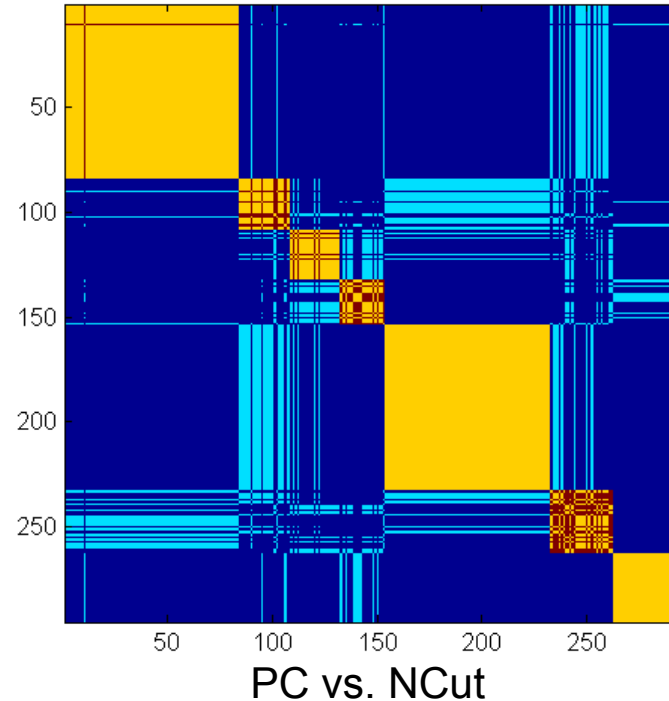
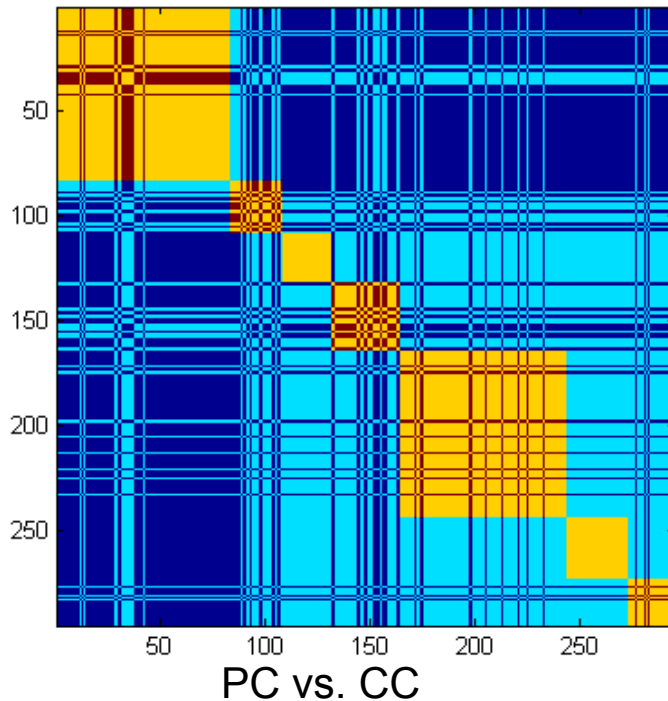
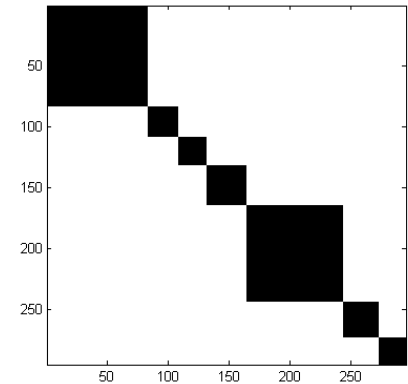


Visualization of PC clusters



Consistency of Clusterings

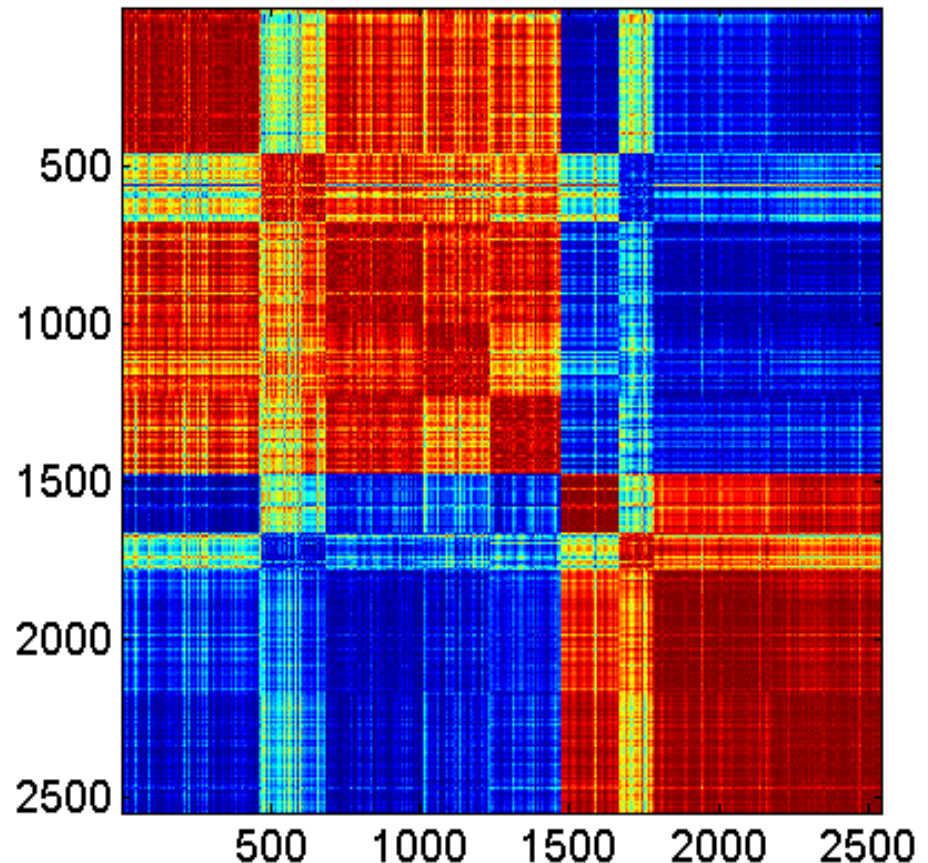
- objects (i,j) are clustered together by PC & CC,
- objects (i,j) are clustered differently by PC & CC,
- objects (i,j) in same cluster by PC and different by CC,
- objects (i,j) in different clusters by PC and same by CC.



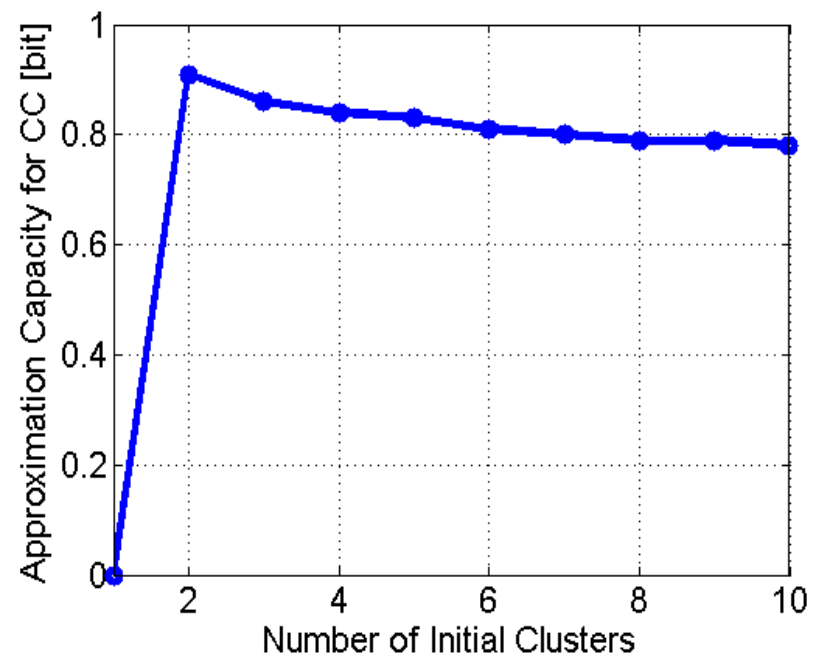
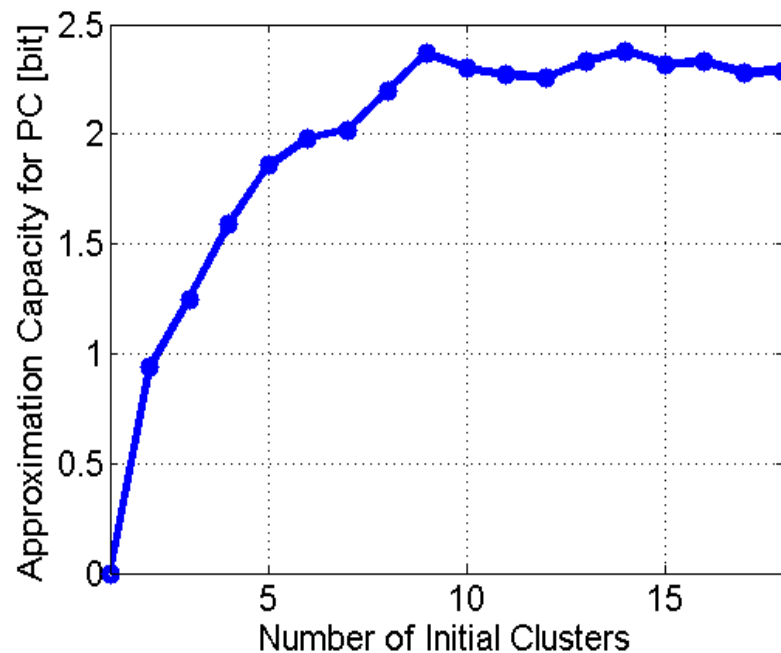
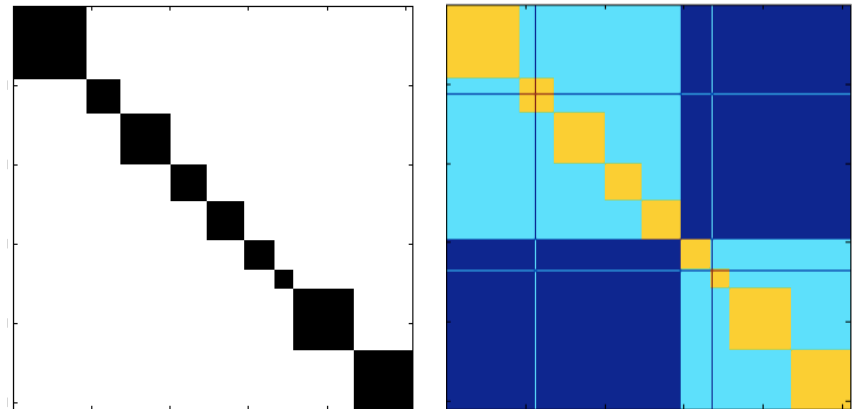
Clustering Yeast Genes for Metabolism

- gene expression values of approx. 2500 yeast genes are clustered to distinguish **dynamic behaviors** and **mechanistic interactions controlling cell growth**;
- Genes are measured at 18 different time points under three biological conditions (treatment with *proline*, *glutamine* and *rapamycin*).

Correlation matrix of gene expressions



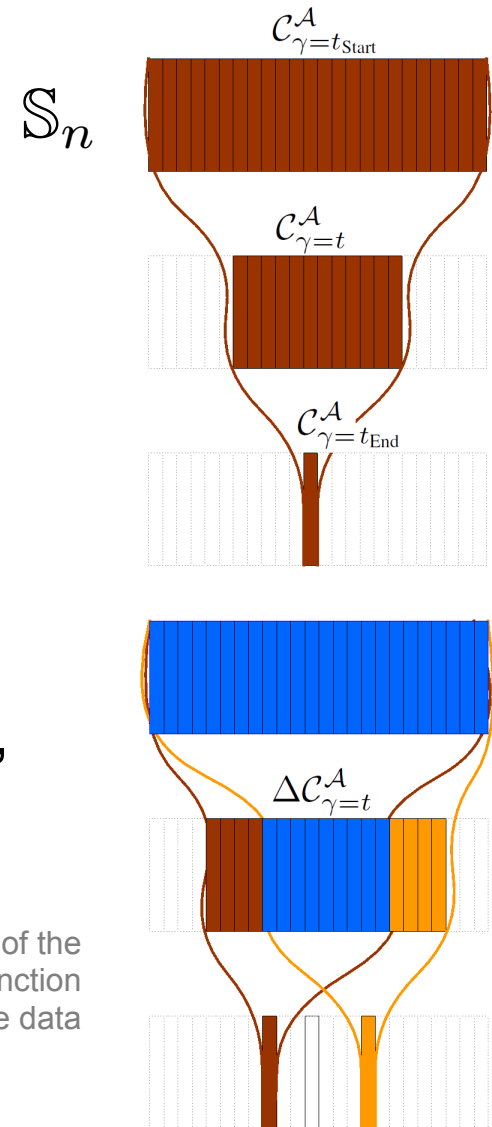
Approximation Capacities for CC and PC



Sorting in noisy conditions

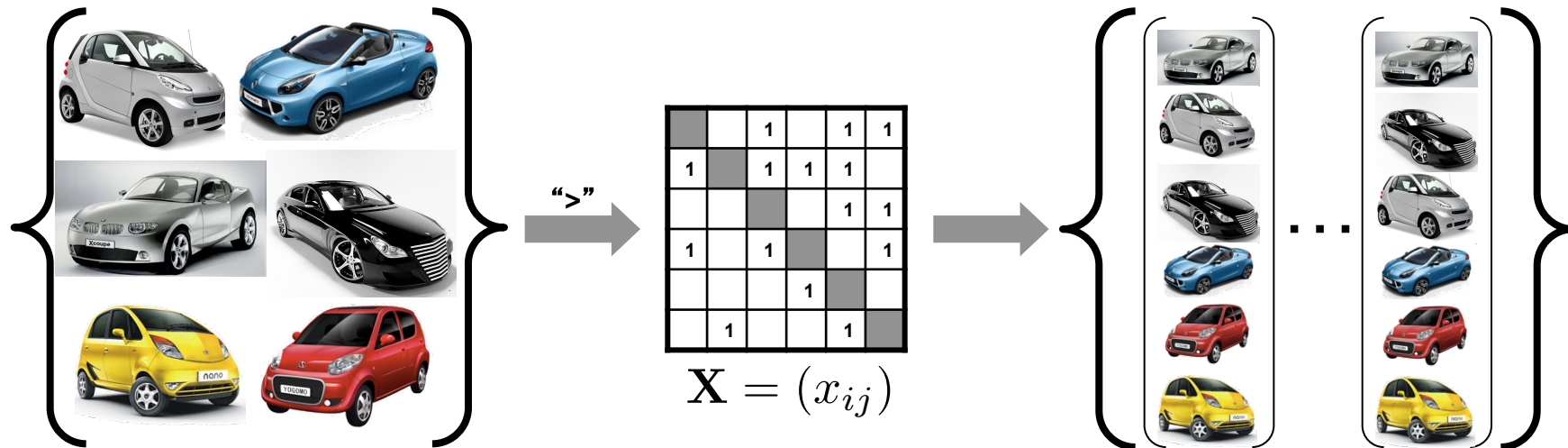
- *Sorting items into an order*
is **restricting the set of rankings**
based on pairwise comparisons
- New method to
optimally sort data in noisy conditions,
i.e. pairwise comparisons can be affected by noise!

Joint work with
L.M. Busse & M.H. Chehreghani



Cardinality of the
approximation set is a function
of the noise level in the data

Approximate Sorting

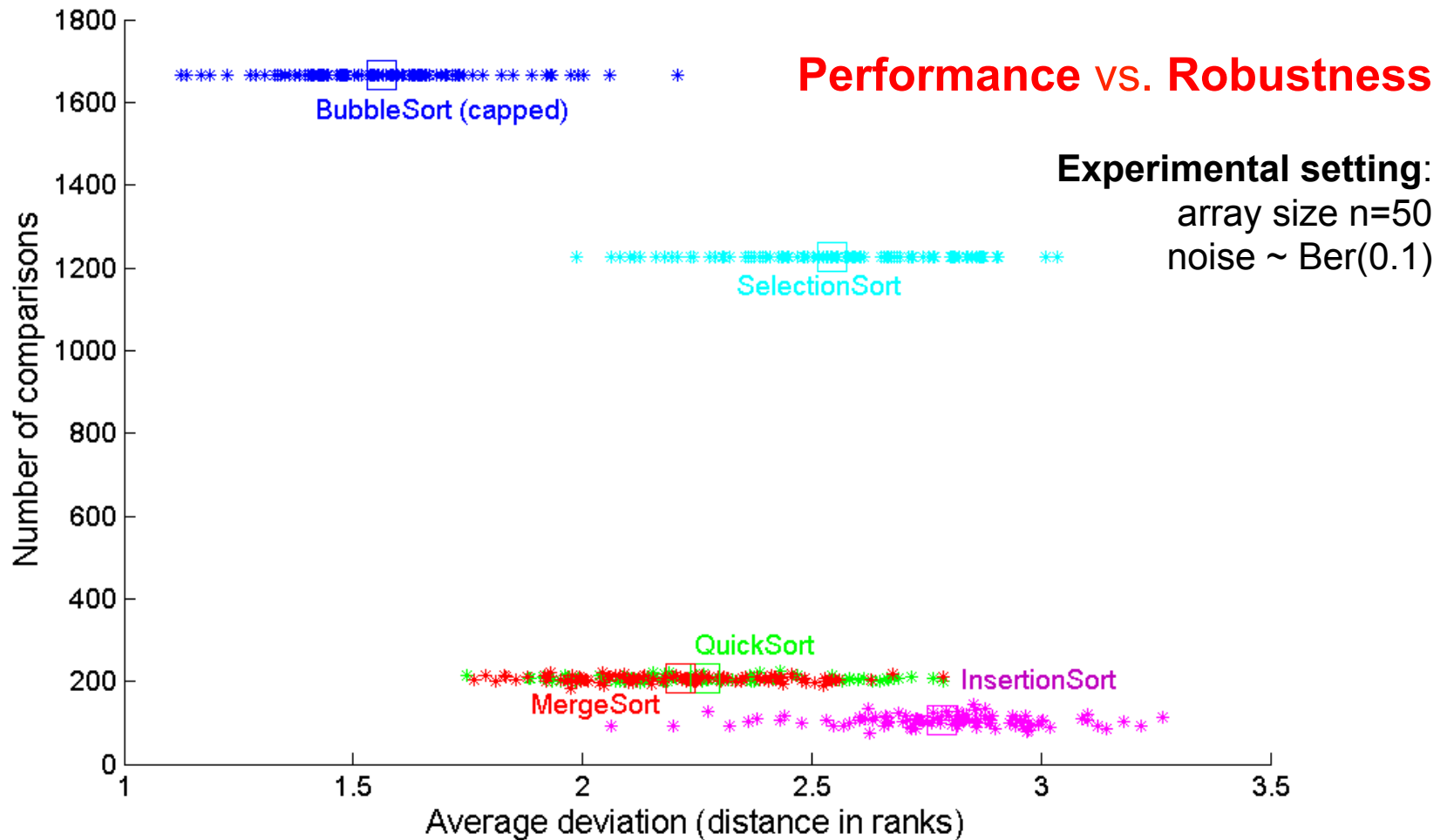


- Among all permutations π , find the one that **orders the items best!**

$$\mathcal{R}^{\text{Sorting}}(\pi | \mathbf{X}) = \sum_{i,j} x_{ij} \mathbb{I}_{\{\pi_i > \pi_j\}}$$

(This cost function counts the number of disagreements between the **provided pairwise comparisons** and the **pairwise comparisons induced by a proposal ranking** from the hypothesis class.)

Pareto-Optimality of Sorting Algorithms



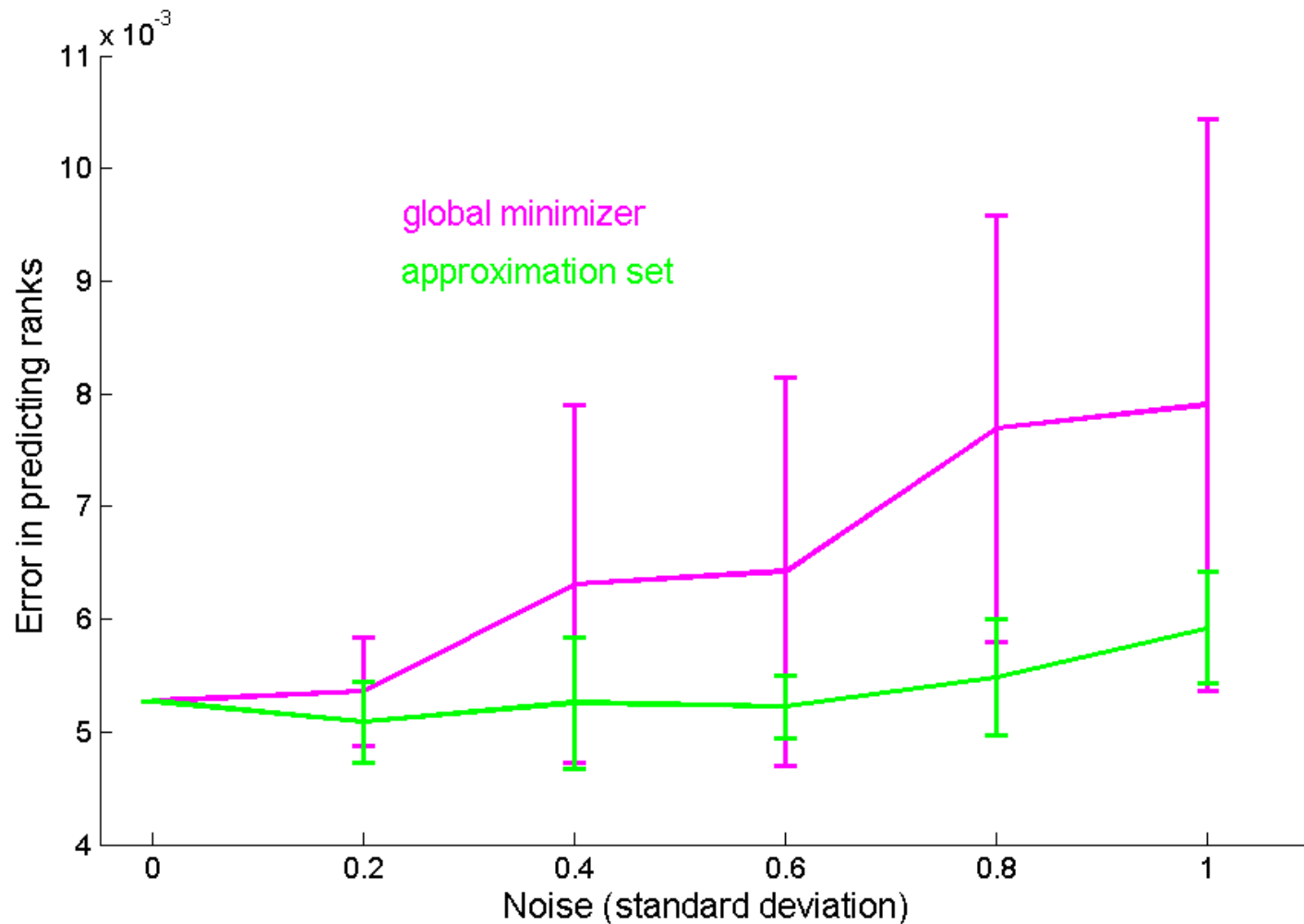
Approximate Sorting

- *Optimally balance the tradeoff between extracting as much reliable partial order information as possible from the data, while being **maximally robust against the noise type** actually present in the data.*

$$\begin{aligned}\mathcal{I}_\beta = & \frac{1}{n} \log |\mathcal{T}| + \frac{1}{n} \sum_{i=1}^n \log \sum_{k=1}^n \exp \left(-\beta (\mathcal{E}_{ik}^{(1)} + \mathcal{E}_{ik}^{(1)}) \right) \\ & - \frac{1}{n} \sum_{i=1}^n \log \left(\sum_{k=1}^n \exp \left(-\beta \mathcal{E}_{ik}^{(1)} \right) \sum_{k'=1}^n \exp \left(-\beta \mathcal{E}_{ik'}^{(2)} \right) \right)\end{aligned}$$

Prediction Performance

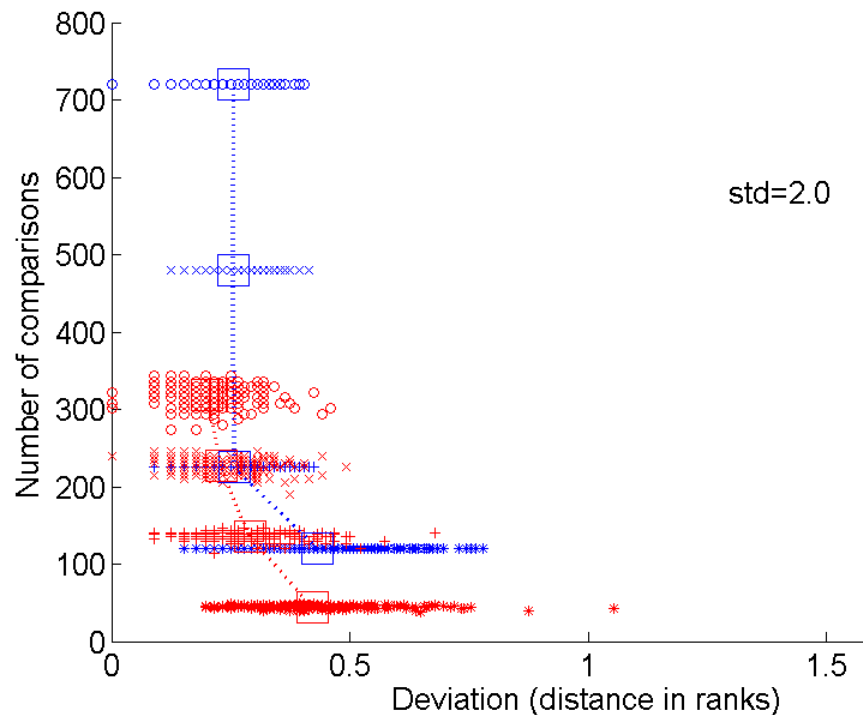
e.g. predicting soccer team ranks
based on records of wins and losses



Pareto-Optimality of Sorting Algorithms

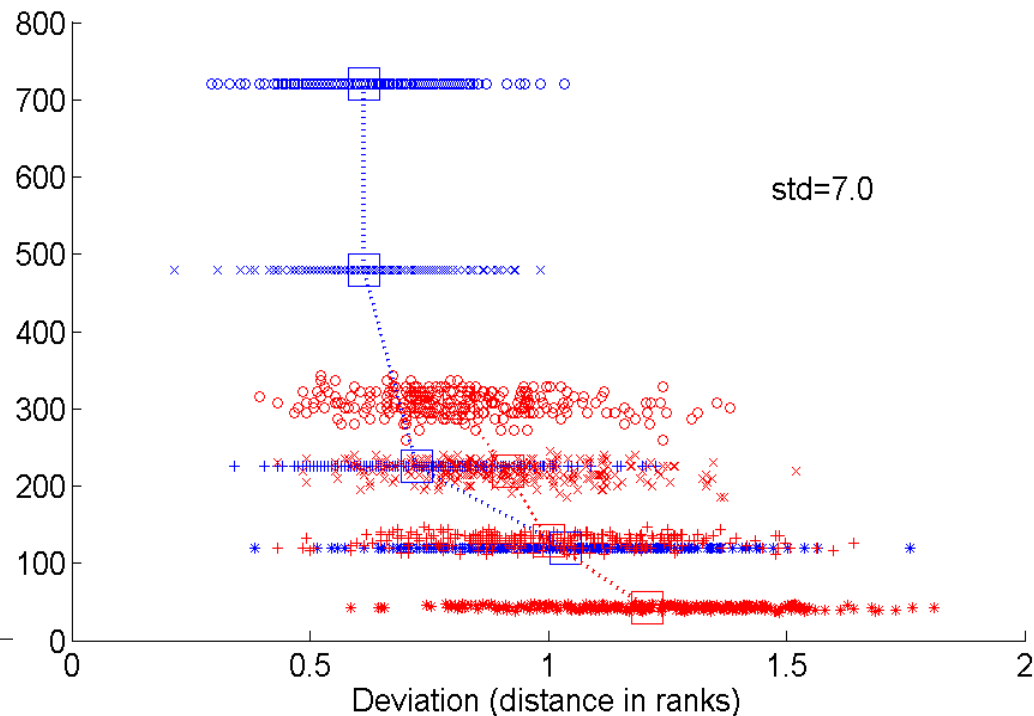
MergeSort

- 7x majority vote
- x 5x
- + 3x
- * 1x



BubbleSort

- 3n bubble steps
- x 2n
- + n
- * n/2



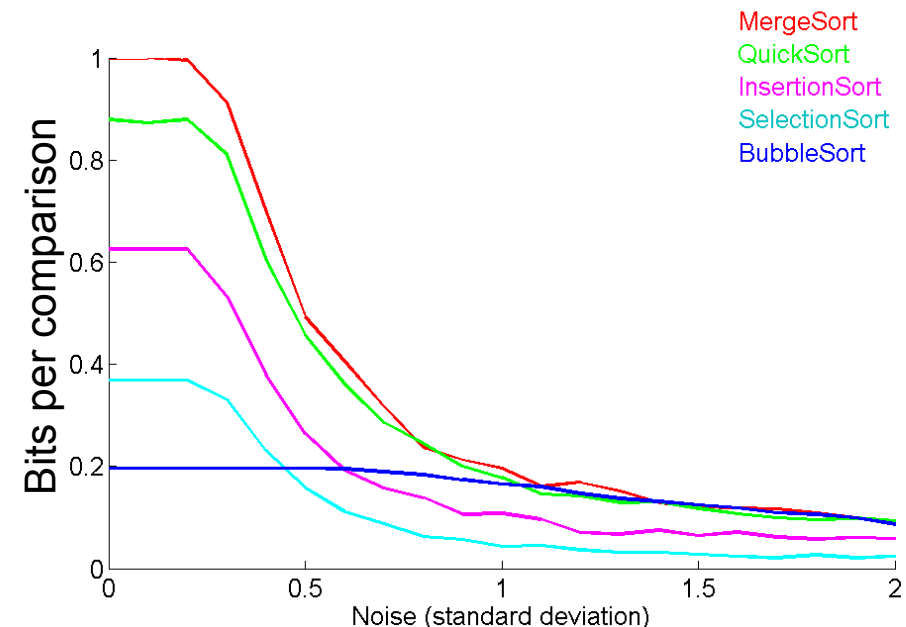
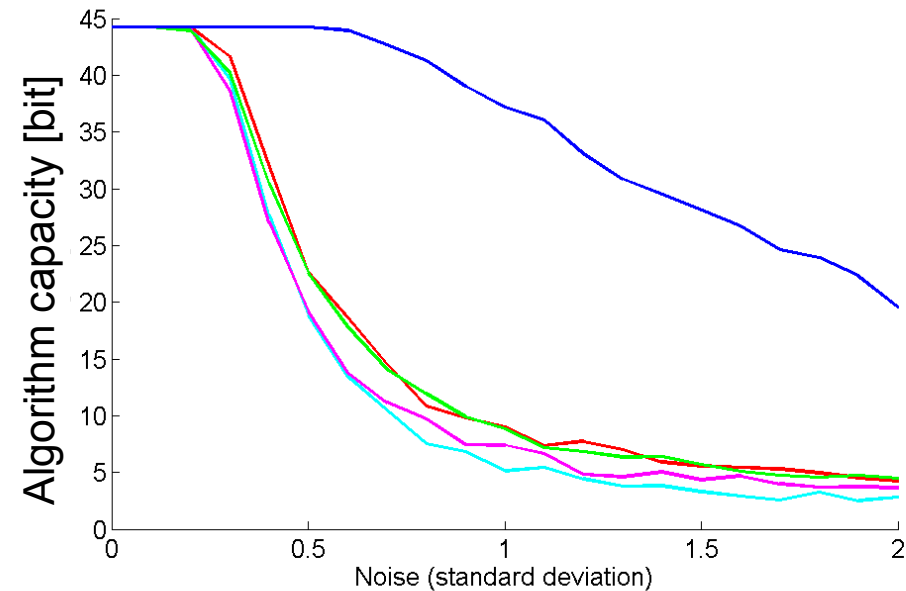
Information Capacity of Sorting Algorithms

Preliminary study: $n=16$

- Without runtime constraints

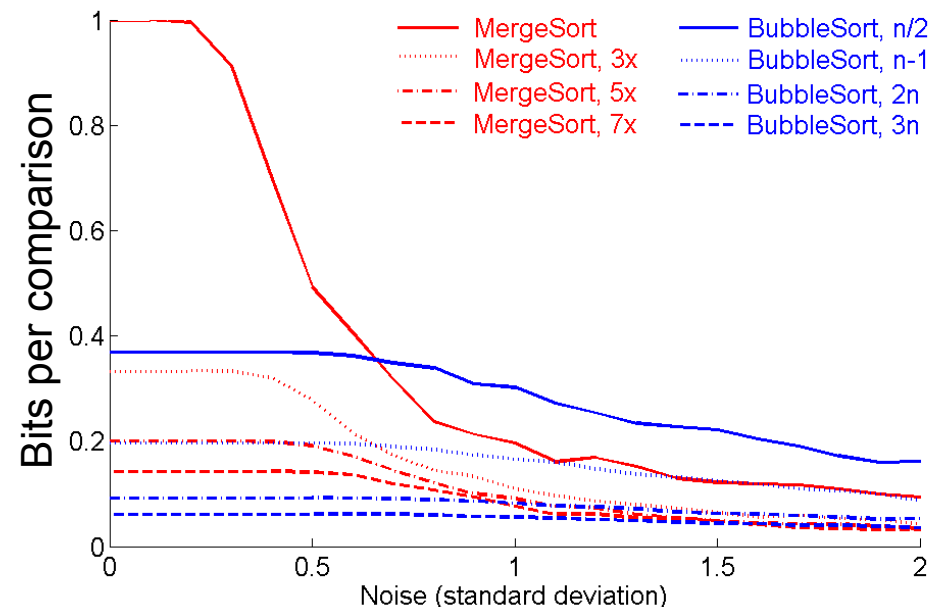
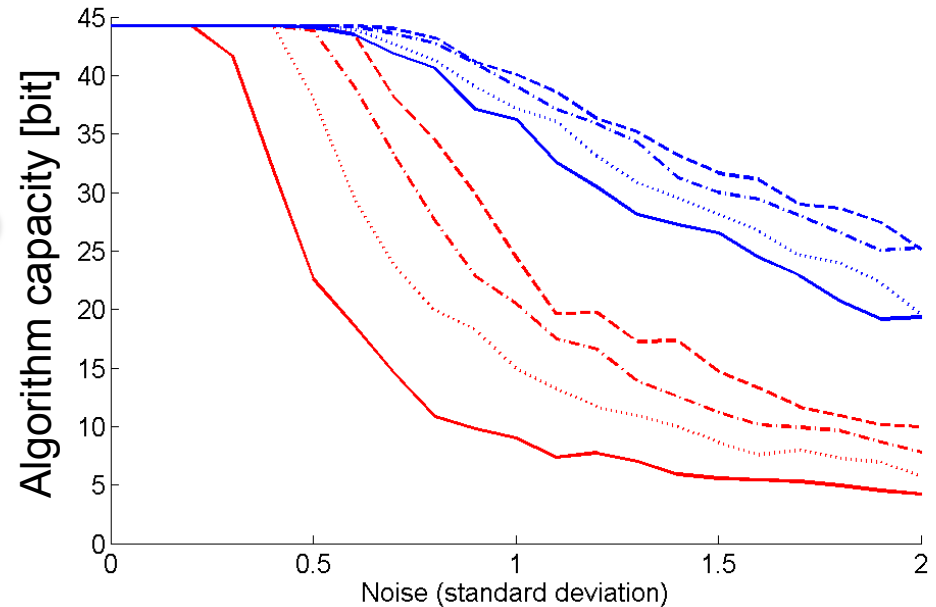
BubbleSort achieves maximal capacity.

- BubbleSort** is most robust at high noise levels, i.e., it extracts most information per comparison!



Information Content of BubbleSort ($O(n^2)$) to MergeSort ($O(n \log n)$)

- BubbleSort with resource budget outperforms MergeSort with majority voting.
- Algorithm design should be adapted to redundancy/efficiency tradeoff!



Conclusion

- **Quantization:** Noise quantizes mathematical structures (hypothesis classes) $\Rightarrow$ symbols
 - These symbols can be used for **coding!**
 - Optimal error free coding scheme determines **approximation capacity** of a model class.
- $\Rightarrow$ Bounds for robust optimization.
- $\Rightarrow$ **Quantization** of hypothesis class measures **structure specific information** in data.

Future Work

- **Generalization**: replace approximation sets based on cost functions by smoothed outputs of **algorithms** (“smoothed generalization”)
- **Model reduction** in dynamical systems: quantize sets of ODEs or PDEs (systems biology)
- Relate **statistical complexity**, i.e. the approximation capacity, to algorithmic or **computational complexity**.



**Credit to the
ETH Machine Learning Group**
Funding: SIMBAD (EU), FOR 916 (DFG & SNF)

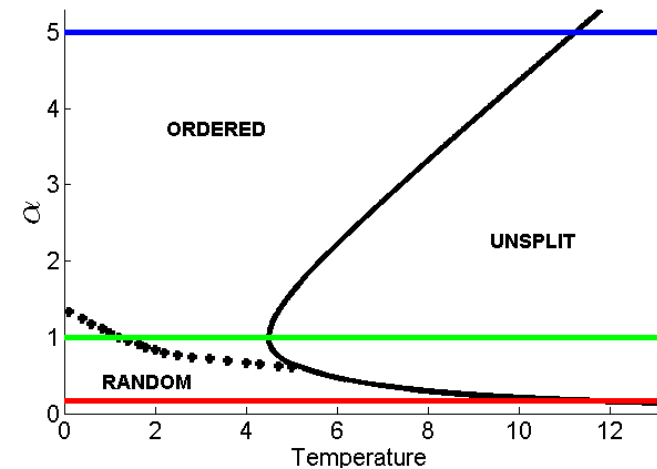
High Dimensional Density Estimation

Barkai, Sompolinsky, Phys Rev E 50:1766, 1994.

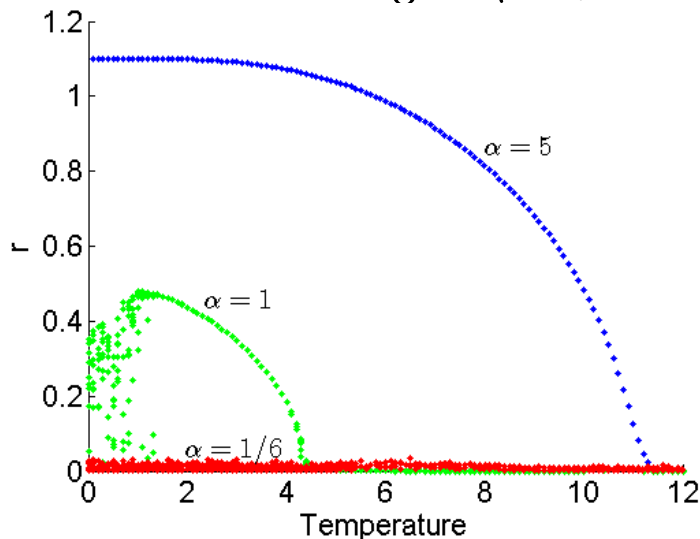
Phase Diagram:

mixture of 2 Gaussians

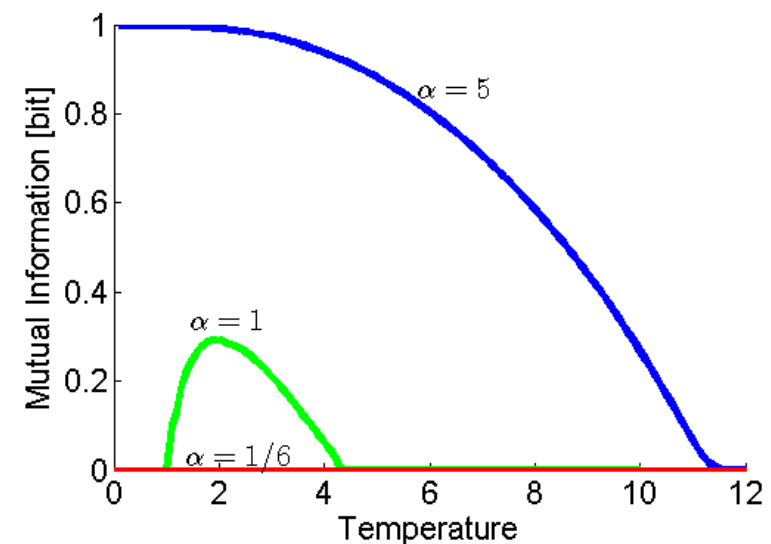
$n=500$; $d=100, 500, 3000$; $\alpha := n/d$



Overlap: $r = u_0^{-1} \langle \Delta\mu, \Delta\mu_0 \rangle$



Approximation Capacity



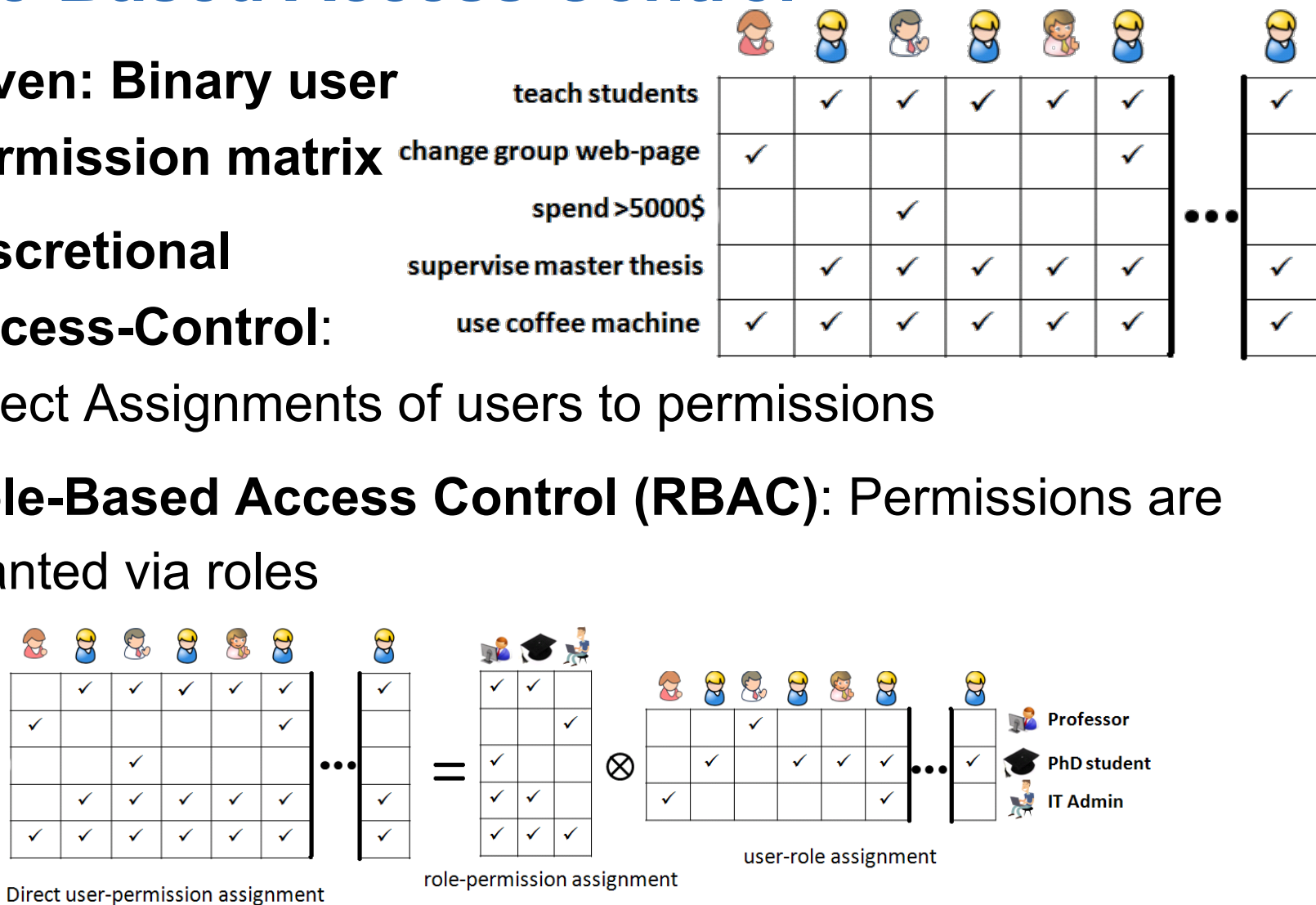
Role-Based Access Control

- Given: Binary user permission matrix

- Discretionary Access-Control:

Direct Assignments of users to permissions

- Role-Based Access Control (RBAC): Permissions are granted via roles

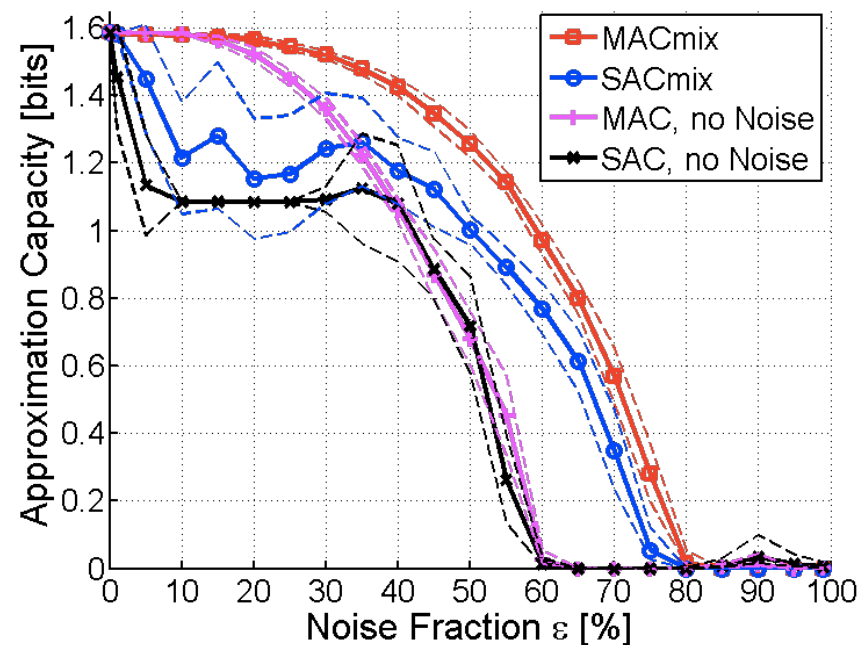
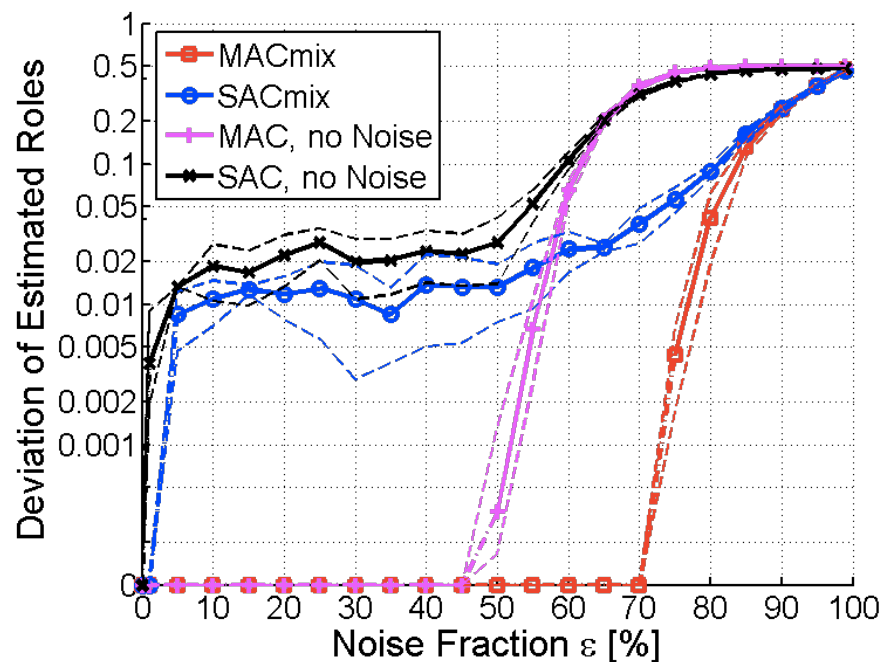


Role-Mining for RBAC

- **Role-Mining:** Given a user-permission assignment matrix $\mathbf{X}$, find a set of roles $\mathbf{U}$ and assignments $\mathbf{Z}$ such that
$$\mathbf{X} \approx \mathbf{U} \otimes \mathbf{Z}$$
- **Multi Assignment Clustering:** generative approach including noise model, inference with DA



Synthetic Data: Parameter Accuracy vs. Approximation Capacity



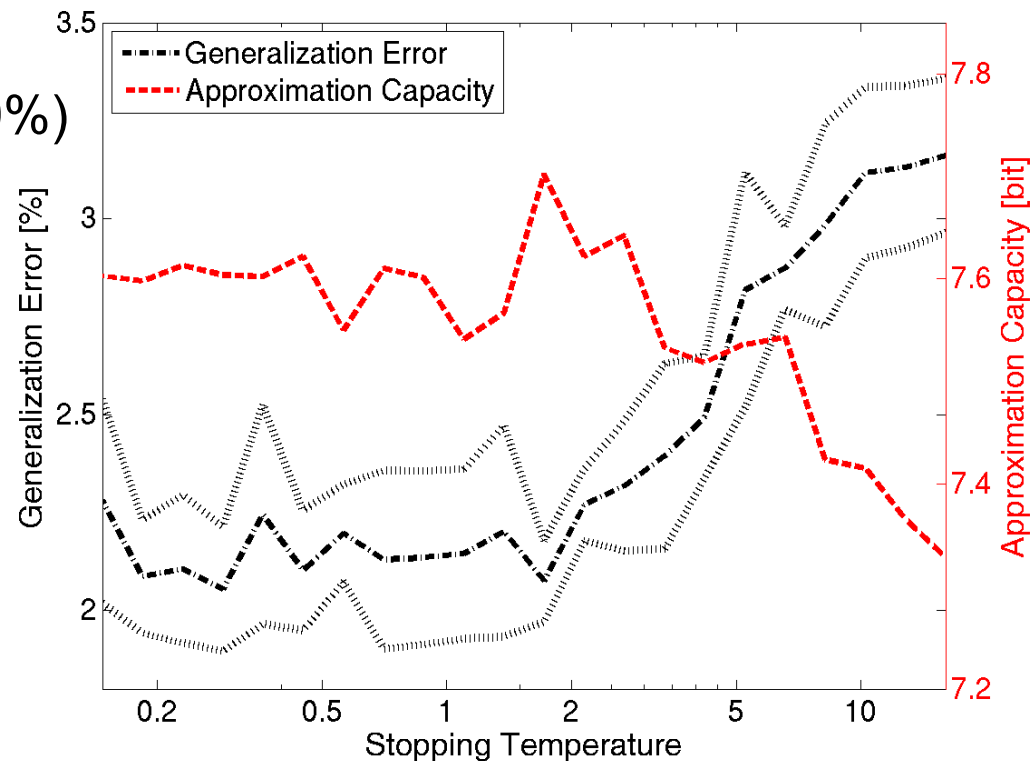
MAC: More accurate estimators for centroids, it yields higher approximation capacity than SAC.

Real-World Data: Prediction Error vs. Approximation Capacity

- **Generalization:** Can roles predict permissions of **new** users?

1. Use few permissions (20%) to determine role set
2. Predict hidden/missing permissions (80%).

- Centroids with maximal capacity yield minimal generalization error



Optimization approach to machine learning

- Given: **data** $\mathbf{X} \in \mathcal{X}$ in **data (input) space** $\mathcal{X}$
- **Goal: Learn structure from data**, i.e., interpret data relative to a hypothesis class
- **Hypothesis class** $\mathcal{C}$ with hypotheses (solutions)
$$c : \mathcal{X} \rightarrow \mathbb{K} \quad (\text{e.g., } \mathbb{B}^n \text{ or } \{1, \dots, k\}^n)$$

$$\mathbf{X} \mapsto c(\mathbf{X})$$

- **Cost function** to define a partial order on $\mathcal{C}$

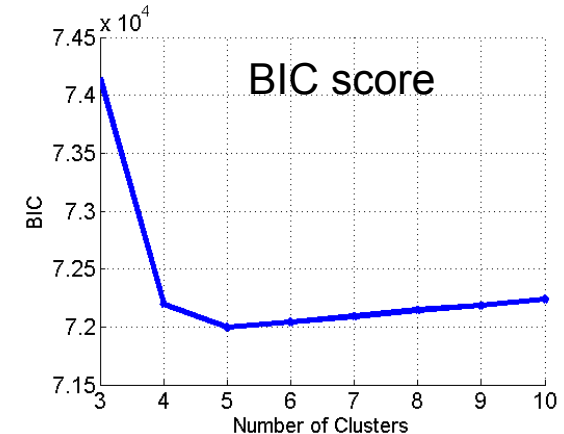
$$R : \mathcal{C} \times \mathcal{X} \rightarrow \mathbb{R}_{\geq 0}$$

$$(c, \mathbf{X}) \mapsto R(c, \mathbf{X})$$

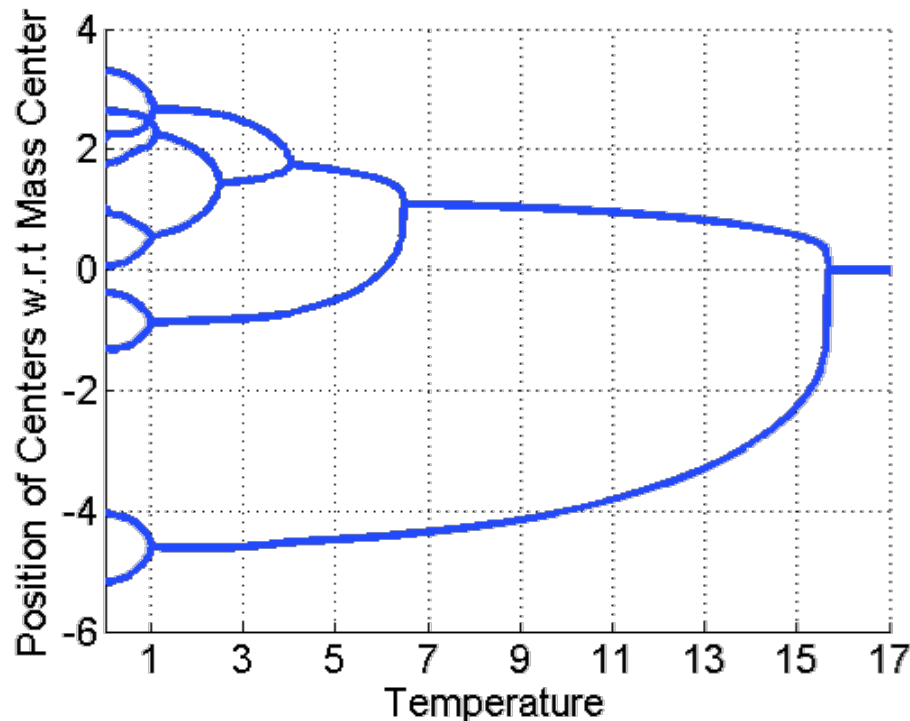
2d Mixture Model Estimation

Experimental Setting:

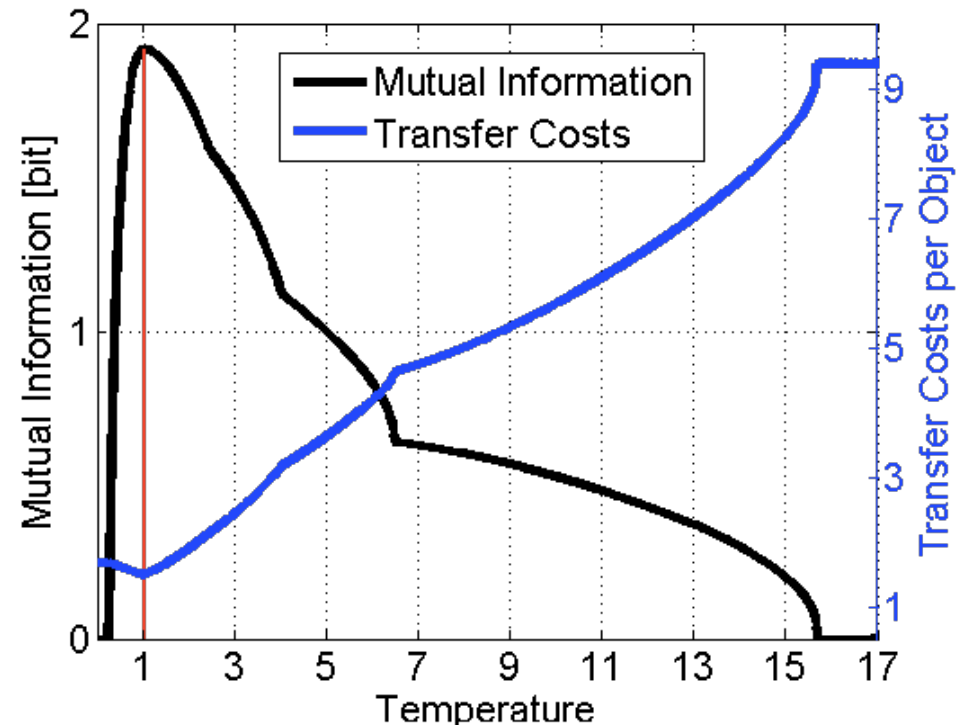
5 Gaussians, $n=10000$, $d=2$, $k^{\max}=10$



Cluster splitting

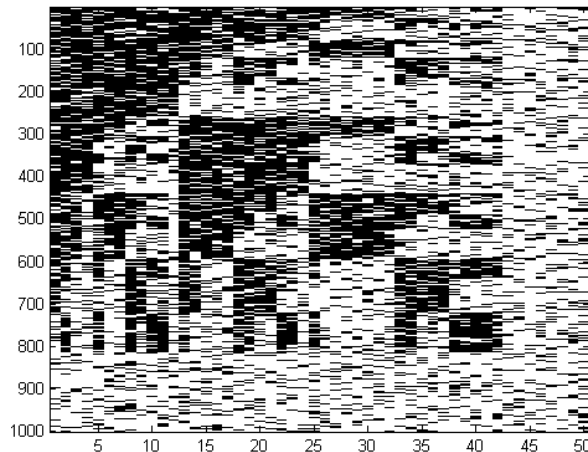


Approximation Capacity



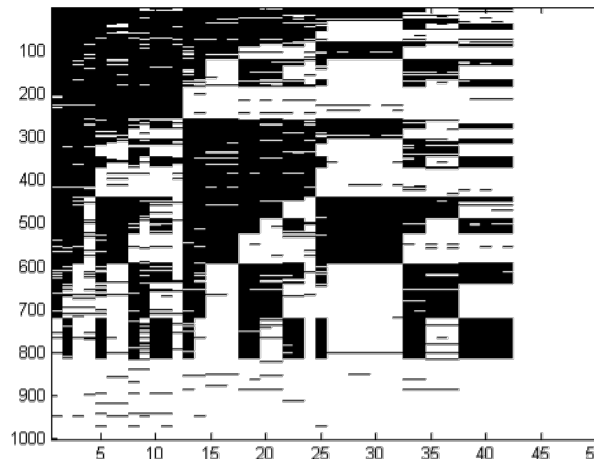
Denoising Binary Matrices by rank- k approximation

Boolean matrix with 40% random entries



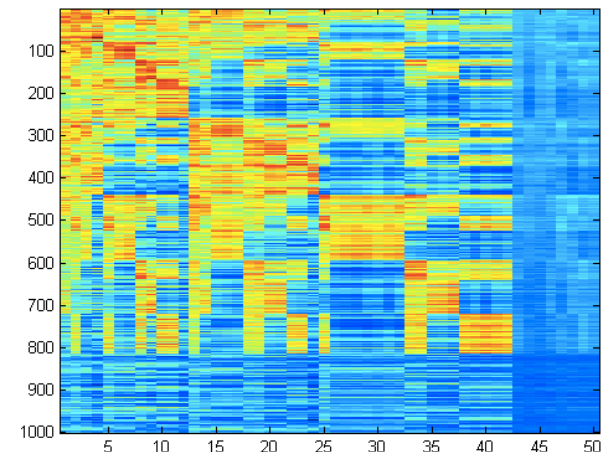
$$X = USV$$

Rounding as
approximation
 $g(X_k) = \text{round}(X_k)$



continuous rank- k approximation

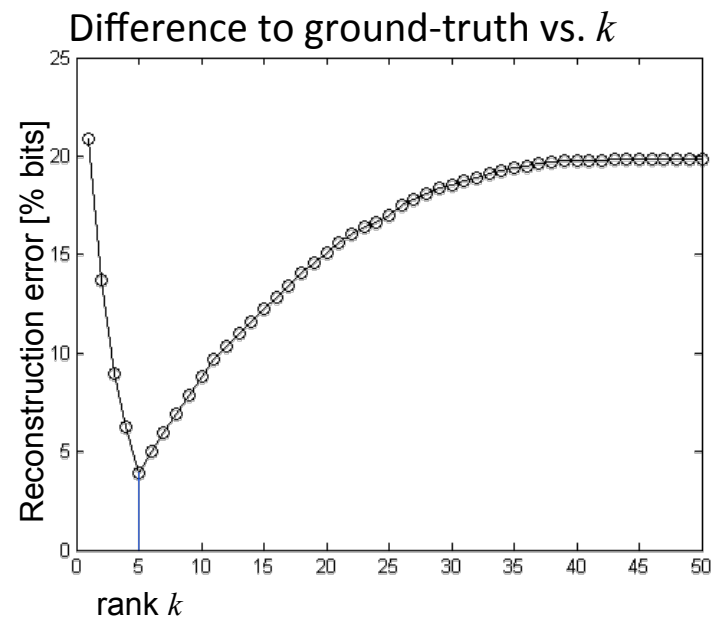
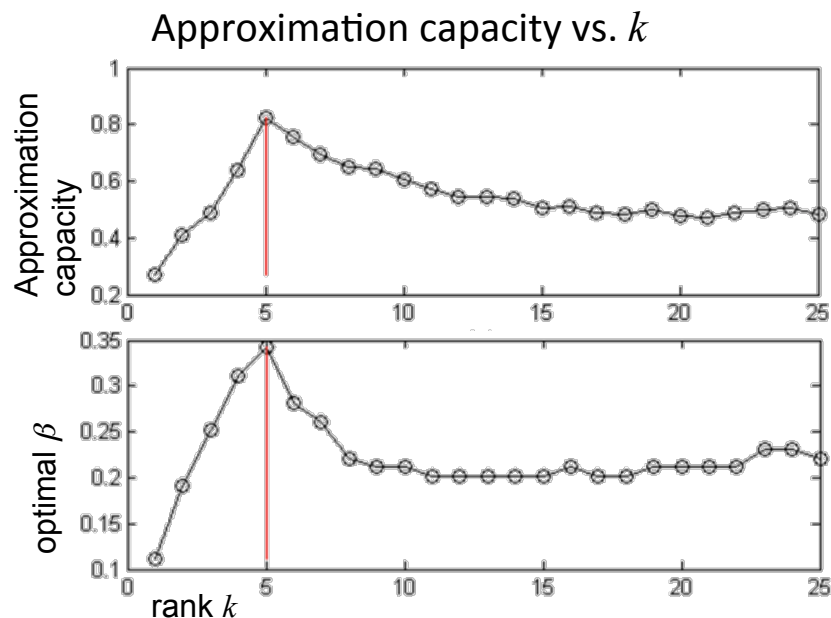
$$X_5 = U_5 S_5 V_5$$



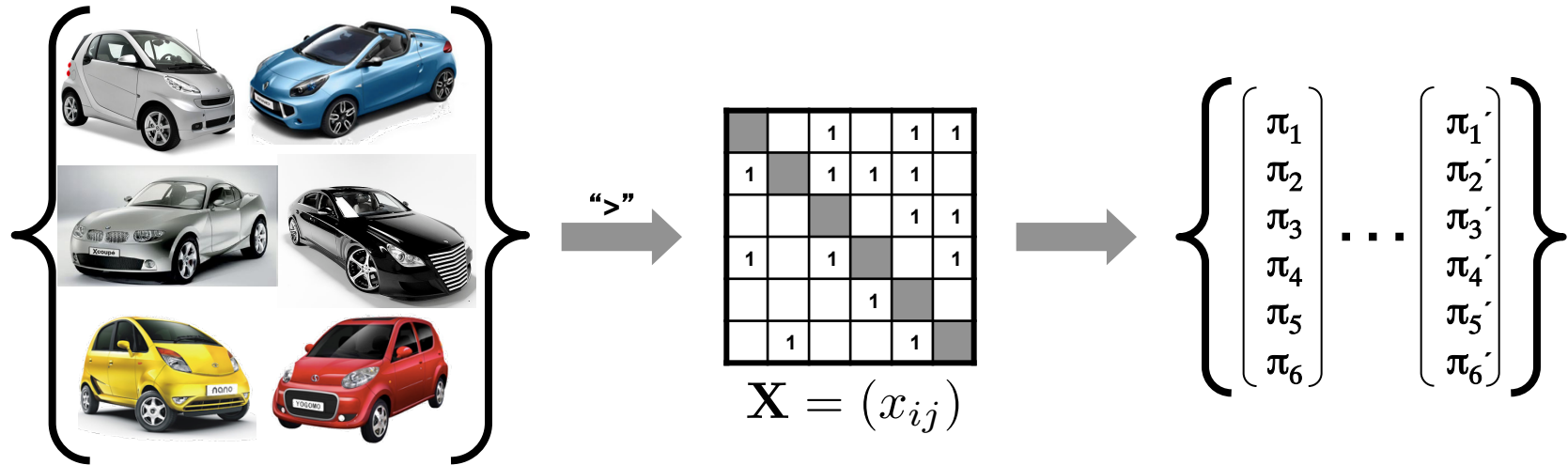
Maximum of approximation capacity selects optimal rank k

- Integrate over variations of the signal matrix $\mathbf{U}$.

$$\hat{\mathcal{I}}_{\beta}(\tau_s, \hat{\tau}) = \frac{1}{n} \log \frac{|\{\tau_s\}| Z_{\beta}^{1+2}(\mathbf{X}^{(1)}, \mathbf{X}^{(2)})}{Z_{\beta}(\mathbf{X}^{(1)}) Z_{\beta}(\mathbf{X}^{(2)})}$$



Approximate Sorting

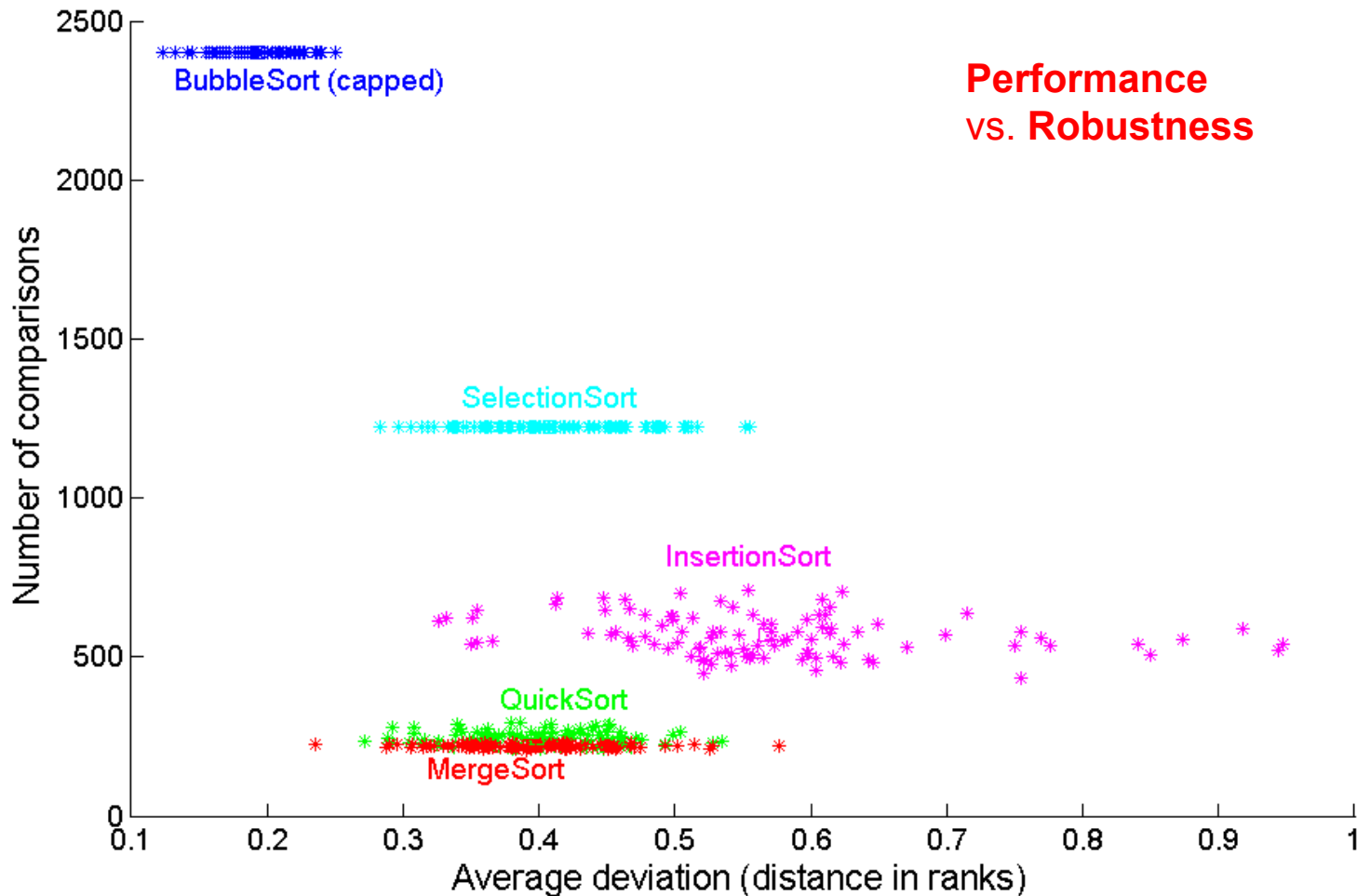


- Among all permutations π , find the one that **orders the items best!**

$$\mathcal{R}^{\text{Sorting}}(\pi | \mathbf{X}) = \sum_{i,j} x_{ij} \mathbb{I}_{\{\pi_i > \pi_j\}}$$

(This cost function counts the number of disagreements between the **provided pairwise comparisons** and the **pairwise comparisons induced by a proposal ranking** from the hypothesis class.)

Pareto-Optimality of Sorting Algorithms



Experimental setting: array size 50, noise 10%