

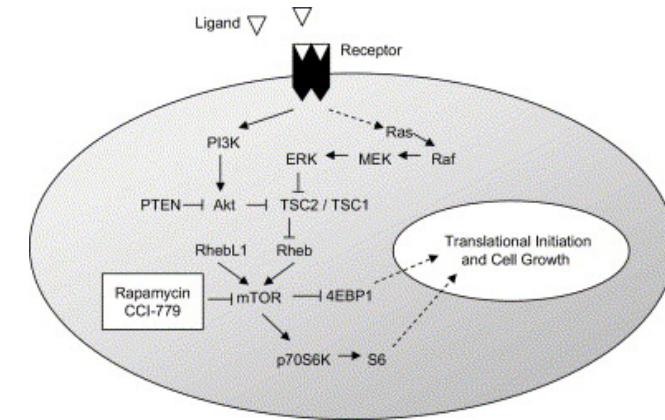
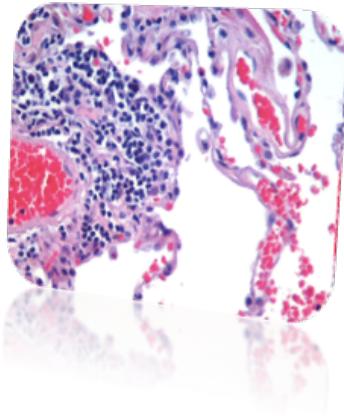
# Information theory of algorithms

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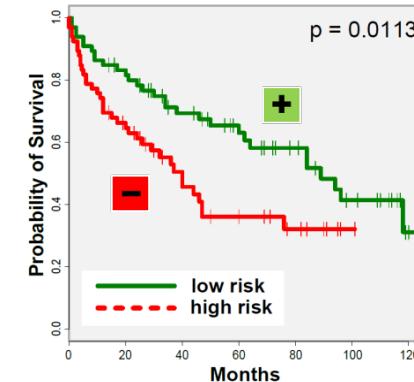
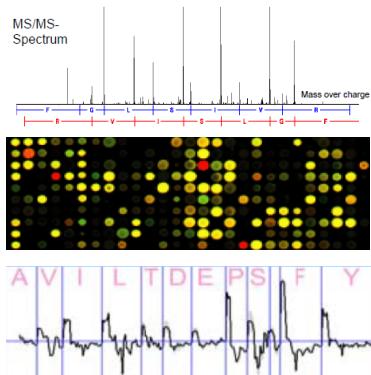


# IT value for personalized medicine



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

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



*my Value*

# Roadmap

- **Robust computation versus learning**  
Measuring the information content of algorithms
- **Algorithm/Model validation** by information theory
- **Learning optimal algorithms:** open challenge!
  - Validating approximate spanning trees
  - Graph cut for gene expression analysis
  - Robust sorting

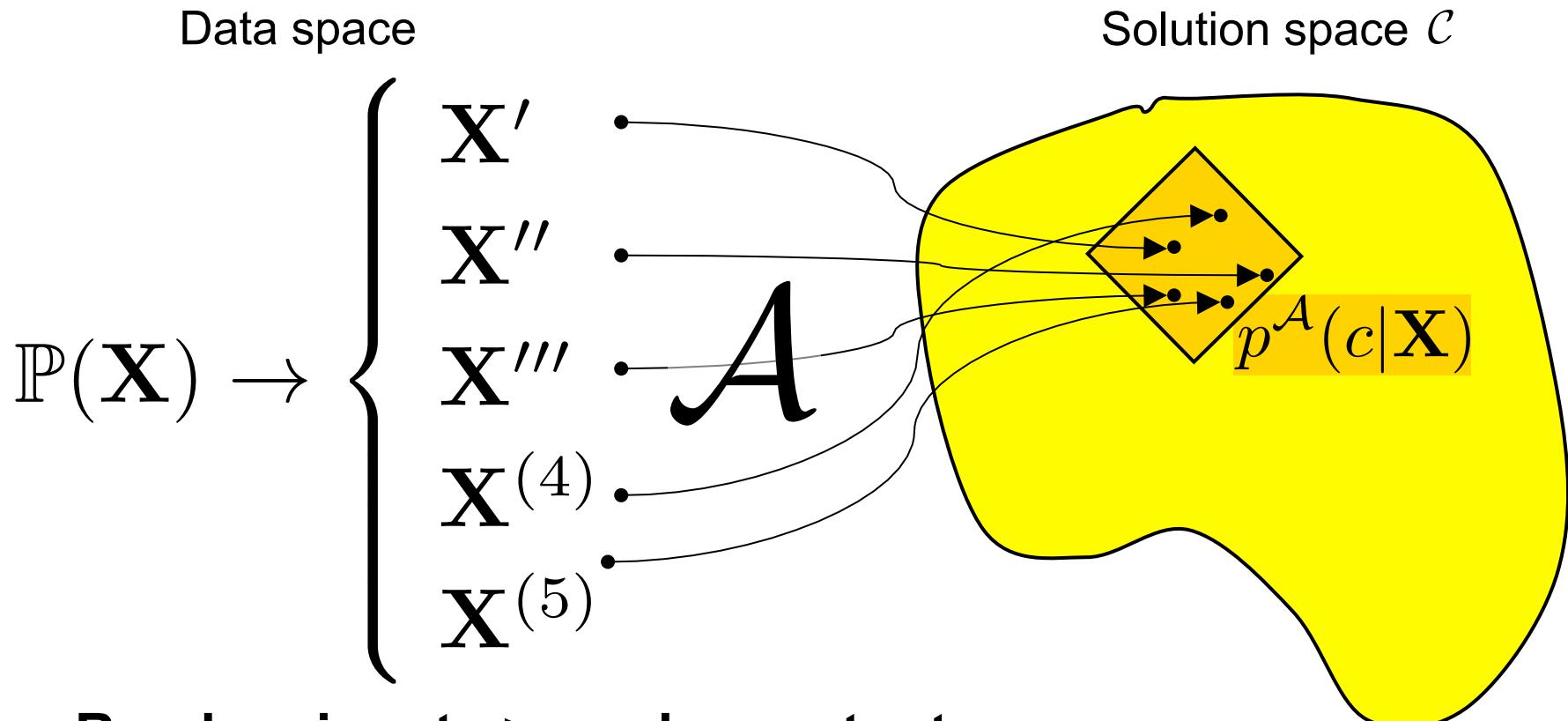
# What is an algorithms?

- Informally, an **algorithm** is any well-defined computational procedure that takes some value(s) as **input** and produces some value(s) as **output** (CLRS' 01)
- **Classical view:** algorithm  $\mathcal{A}$  maps (stochastic) data (input) to a solution / hypothesis (output).

$$\text{input } \mathbf{X} \rightarrow \mathcal{A} \rightarrow \text{output } c \in \mathcal{C}$$

# Challenge of robust algorithm design

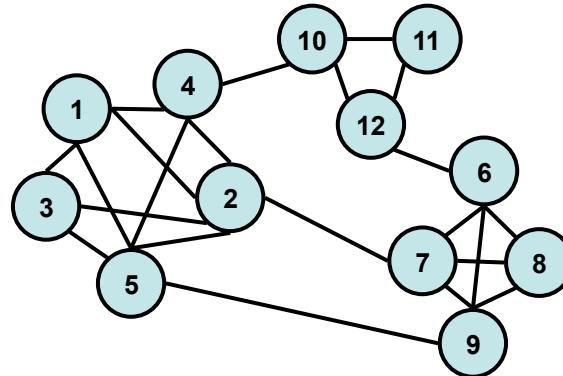
Algorithmic processing: input  $\mathbf{X} \rightarrow \mathcal{A} \rightarrow$  output  $c \in \mathcal{C}$



# Random input => random output

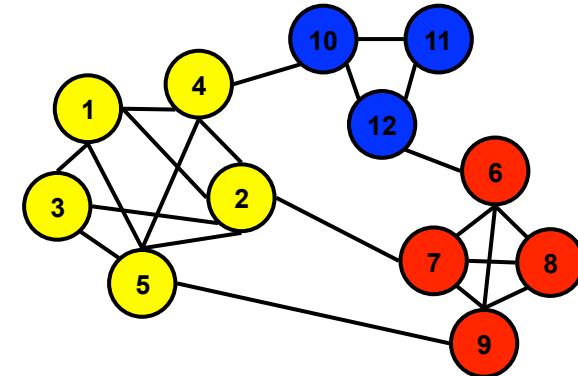
# Core problem in discriminative learning?

- Often, the data space is much larger than the solution space



Space of graphs  $\mathfrak{G}_n = \{(\mathcal{V}, \mathcal{E}, \mathcal{W})\}$   
with  $n$  vertices

$$|\mathfrak{G}_n| = \mathbb{R}^{\binom{n}{2}}$$

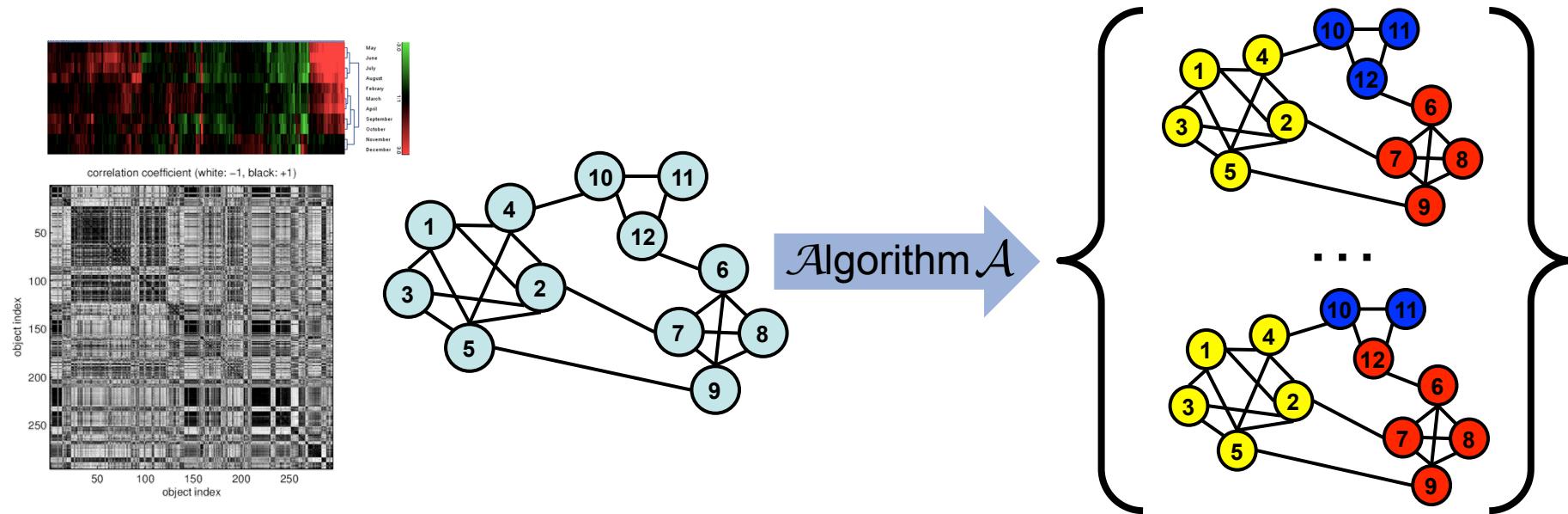


Space of  $k$  colorings  $\mathcal{C}$

$$|\mathcal{C}| \leq k^n$$

# What is learning?

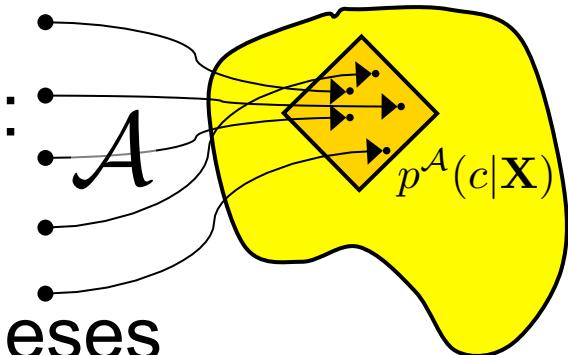
- Given: data  $\mathbf{X} \sim \mathbb{P}(\mathbf{X})$  and a hypothesis class  $\mathcal{C}$



- Modeling in data analysis requires
  - quantization:** given  $\mathcal{A}$ , identify a set of good hypotheses;
  - learning:** find an  $\mathcal{A}$  that specifies an informative set!

# Robust algorithms

- For **stochastic input**, an algorithm  $\mathcal{A}$  returns a **stochastic output**; we interpret  $\mathcal{A}$  as a trajectory in the powerset of the solution space  $\mathcal{C}$ .
- **Discriminative probabilistic view:** algorithm  $\mathcal{A}$  maps stochastic data (input) to a set of solutions / hypotheses  $A_t(\mathbf{X})$  (output) still considered at time  $t$   
 $\text{input } (\mathbf{X}, t^*) \rightarrow \mathcal{A} \rightarrow \text{output posterior } p^{\mathcal{A}}(c|\mathbf{X})$



# Algorithms as sets of feasible solutions

- Let  $X$  denote data and  $A_t(\mathbf{X})$  the set of feasible solutions at iteration  $t$

algorithm  $\mathcal{A}(\mathbf{X}) = \langle A_0(\mathbf{X}), \dots, A_T(\mathbf{X}) \rangle,$

**init**  $A_0(\mathbf{X}) = \mathcal{C},$

$A_t(\mathbf{X}) \subseteq \mathcal{C}, \quad 0 \leq t \leq T,$

**return**  $A_T(\mathbf{X}) = \{c^\perp(\mathbf{X})\}.$

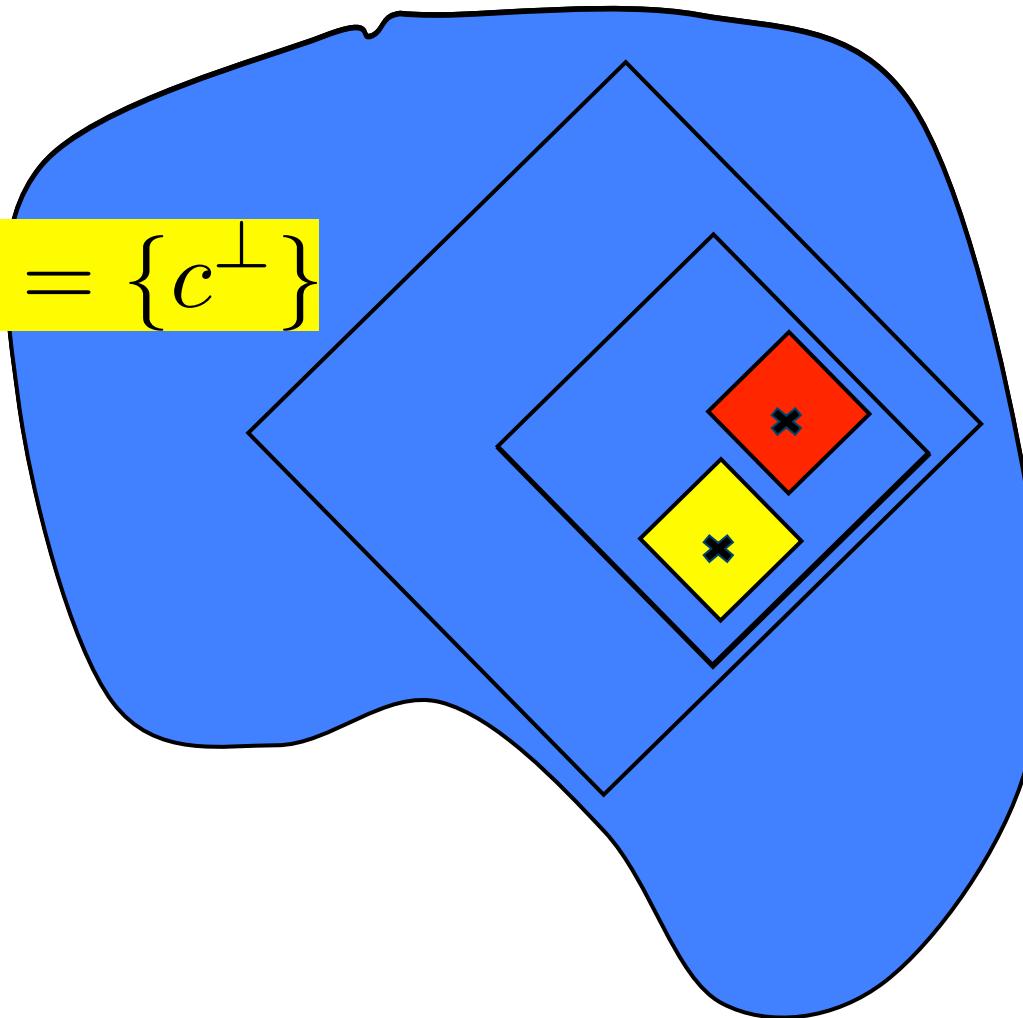
- Monotonically contractive algorithms

$\mathcal{A}(\mathbf{X}) = \langle A_0(\mathbf{X}) \supseteq \dots \supseteq A_{t^*}(\mathbf{X}) \supseteq \dots \supseteq A_T(\mathbf{X}) \rangle$

# Hypotheses explored by an algorithm $\mathcal{A}$

data  $\mathbf{X}'$

$$A_T(\mathbf{X}') = \{c^\perp\}$$

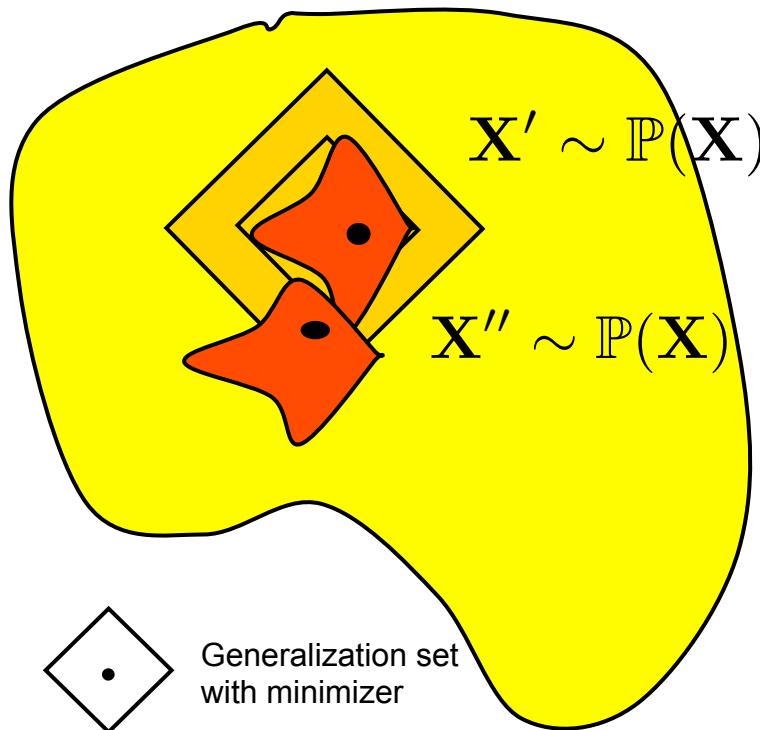


data  $\mathbf{X}''$

$$A_T(\mathbf{X}'')$$

# Coarsening of hypothesis classes and the two instances test

- Quantize hypothesis class by generalization sets



- Size: “partition function” at iteration  $t$  is  $|A_t(\mathbf{X}')|$

- Weight overlap models joint approximations

$$|A_t(\mathbf{X}') \cap A_t(\mathbf{X}'')|$$

- Posterior of hypothesis  $c$

$$p^{\mathcal{A}}(c|\mathbf{X}') = \begin{cases} \frac{1}{|A_t(\mathbf{X}')|} & \text{if } c \in A_t(\mathbf{X}') \\ 0 & \text{otherwise} \end{cases}$$

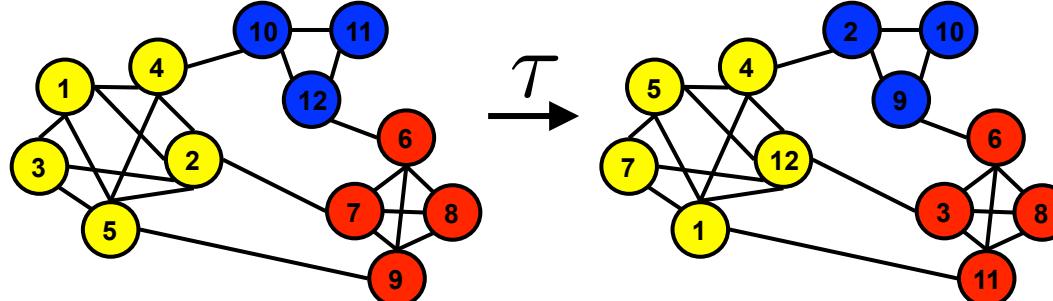
# Information theory: structures as symbols

- 1) sample a hypothesis  $\tilde{c} \sim p^{\mathcal{A}}(c|\mathbf{X})$  Cover **hypothesis class**
- 2) for  $j = 1 \dots M$  densely, but identifiably!

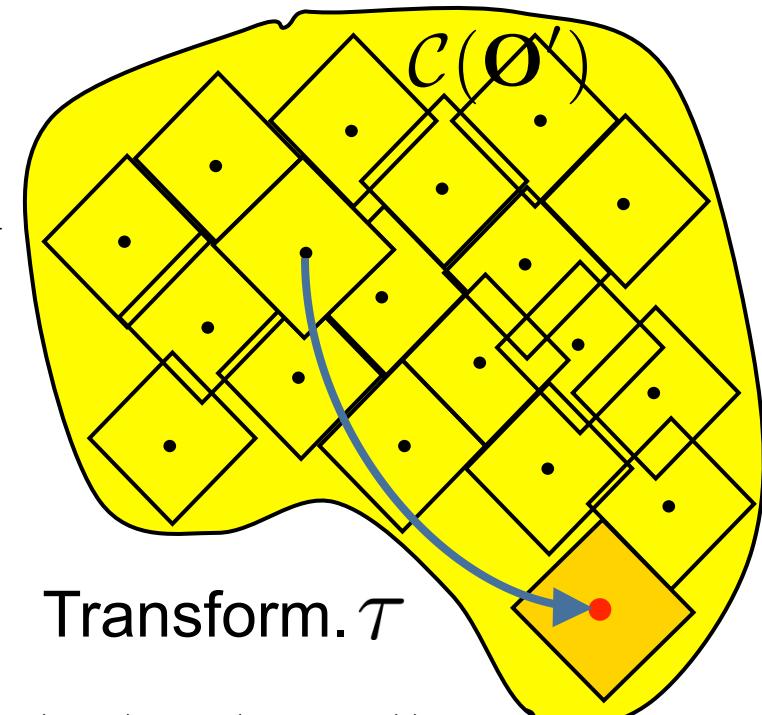
Select a random transf.  $\tau_j \in \mathbb{T}(\tilde{c})$

define code vector  $\tilde{c}_j := \tau_j \circ \tilde{c}$

- 3) return codebook  $\mathcal{T} := \{\tilde{c}_1, \dots, \tilde{c}_M\}$

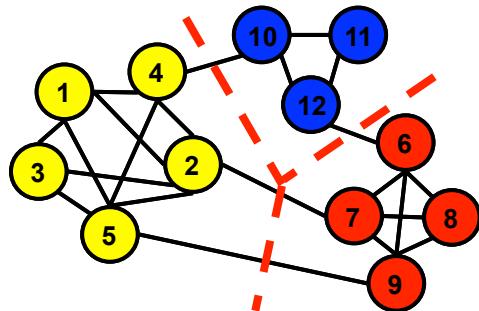


$$\mathbb{T}(\tilde{c}) = \{\tau : \forall \beta, c, w_\beta(c, \mathbf{X}) = w_\beta(\tau \circ c, \tau \circ \mathbf{X}) \wedge D(w_\beta(., \mathbf{X}), w_\beta(., \tau \circ \mathbf{X})) > \rho\}$$



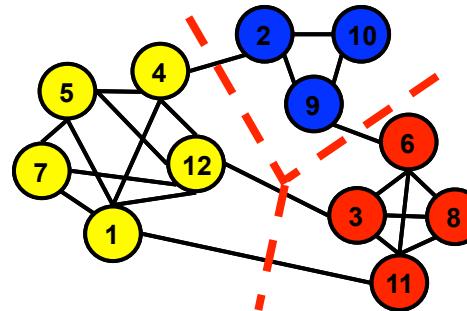
# Code problem generation for graph cut

graph cut code problems

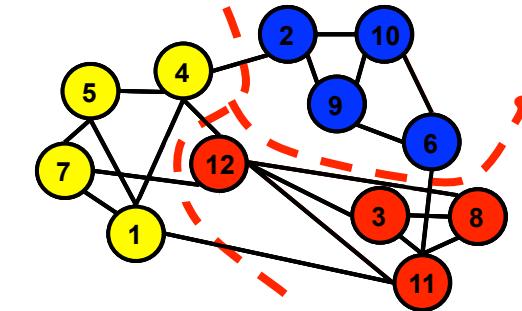


...

$M$



graph cut  
*test* problem



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# Communication process and decoding

- Sender sends transformation  $\tau_s$
- Receiver accepts instance  $\tilde{\mathbf{X}} := \tau_s \circ \mathbf{X}''$  with  $\mathbf{X}', \mathbf{X}'' \sim P(\mathbf{X})$  and decodes the transformation by **maximizing expected posterior**

$$\hat{\tau} \in \arg \max_{\tau \in \mathcal{T}} \mathbb{E}_{\tilde{c} | \tau \circ \mathbf{X}'} p(\tilde{c} | \tau_s \circ \mathbf{X}'')$$

- **Error event:**  $\hat{\tau} \neq \tau_s$

# Generalization capacity from typicality

- **Theorem:** Asymptotic error free *identification*

$\lim_{n \rightarrow \infty} P(\hat{\tau} \neq \tau_s | \tau_s) = 0$  of *code structures* is possible for

$$\begin{aligned} P(\hat{\tau} \neq \tau_s | \tau_s) &\leq M \mathbb{E}_{\mathbf{X}', \mathbf{X}''} \left( |\mathbb{T}| k^{\mathcal{A}}(\mathbf{X}', \mathbf{X}'') \right)^{-1} \\ &\stackrel{\text{typicality}}{\leq} \exp(-(I - \log M)) \quad \text{with} \end{aligned}$$

$$I = \mathbb{E}_{\mathbf{X}', \mathbf{X}''} \log \left( |\mathbb{T}| k^{\mathcal{A}}(\mathbf{X}', \mathbf{X}'') \right)$$

$$k^{\mathcal{A}}(\mathbf{X}', \mathbf{X}'') = \sum_{c \in \mathcal{C}} p^{\mathcal{A}}(c | \mathbf{X}') p^{\mathcal{A}}(c | \mathbf{X}'') \in [0, 1]$$

# Behavior of the generalization capacity

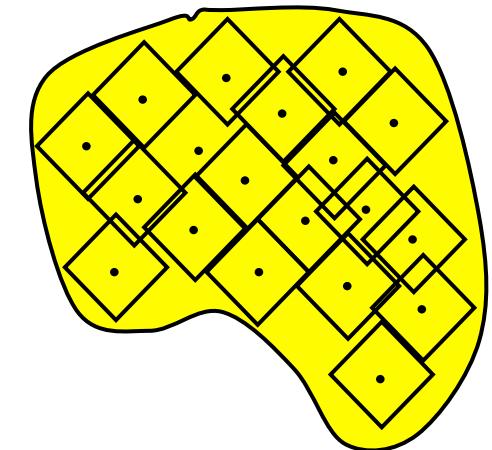
- Let us assume that  $\mathbf{X}', \mathbf{X}''$  are very similar.

$$\mathbf{X}' \approx \mathbf{X}'' \approx \mathbf{X}$$

$$k^{\mathcal{A}}(\mathbf{X}, \mathbf{X}) = \sum_{c \in \mathcal{C}} p^{\mathcal{A}}(c|\mathbf{X})^2 = \frac{|A_{t^*}(\mathbf{X}) \cap A_{t^*}(\mathbf{X})|}{|A_{t^*}(\mathbf{X})|^2}$$

$$\mathcal{I} = \mathbb{E}_{\mathbf{X}} \log \left( \frac{|\mathbb{T}|}{|A_{t^*}(\mathbf{X})|} \right)$$

- 2<sup>nd</sup> order terms yield maximum of GC  $\mathcal{I}$  for parameter selection



# Learning an algorithm: open challenge!

- **Statistical behavior of an algorithm** is described by its posterior  $p^{\mathcal{A}}(c|\mathbf{X}')$
- **Adapt posterior**  $p^{\mathcal{A}}(c|\mathbf{X}')$  s.t. generalization capacity is maximized

$$p^* \in \arg \max_{\substack{p^{\mathcal{A}}: \mathcal{X} \times \mathcal{C} \rightarrow [0, 1] \\ \sum_c p^{\mathcal{A}}(c|\mathbf{X}) = 1}} \mathbb{E}_{\mathbf{X}', \mathbf{X}''} \log(|\mathbb{T}| k^{\mathcal{A}}(\mathbf{X}', \mathbf{X}''))$$

- **Problem:** We cannot evaluate  $\mathbb{E}_{\mathbf{X}', \mathbf{X}''} \log \dots$  since  $P(\mathbf{X}', \mathbf{X}'')$  is unknown!

# Roadmap

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  - Validating approximate spanning trees
  - Graph cut for gene expression analysis
  - Robust sorting

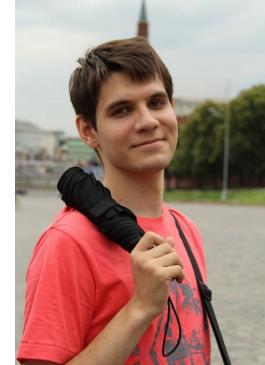
# Example: Robust Spanning Trees

- **Given** are graphs with stochastic edge weights.
- **Question:** How robust are MST algorithms in this stochastic setting?
- Measure **robustness of algorithms** like Prim's, Kruskal's algorithm or the Reverse-Delete algorithm.
- Determine a **stable set of approximate spanning trees** by early stopping of an MST algorithm.

# Learning to span a graph

Consider **Minimum Spanning Tree** algorithms

- **Prim's “Growing tree” strategy:** add minimal edge to tree.
- **Kruskal's “Joining trees” strategy:** add minimal edge connecting two trees in a forest.
- **Reverse-Delete:** “Reducing graph” strategy: delete maximal edge without destroying connectivity.



Alexey Gronskiy

# MST Algorithm as a sequence of approximate spanning tree sets

- Let  $\mathbf{X}$  be a graph and  $A_t(\mathbf{X})$  the set of spanning trees at iteration  $t$

$$\mathcal{A}(\mathbf{X}) = \langle A_0(\mathbf{X}), \dots, A_T(\mathbf{X}) \rangle,$$

$$A_t(\mathbf{X}) \subseteq \mathcal{C}, \quad 0 \leq t \leq T,$$

$$A_0(\mathbf{X}) = \mathcal{C}, \quad A_T(\mathbf{X}) = \{c^\perp\}.$$

- Monotonically contractive algorithms

$$\mathcal{A}(\mathbf{X}) = \langle A_0(\mathbf{X}) \supseteq A_1(\mathbf{X}) \supseteq \dots \supseteq A_T(\mathbf{X}) \rangle$$

# Cardinality of AST sets

- **Matrix tree theorem:** the number of spanning trees is equal to (any) cofactor of the matrix

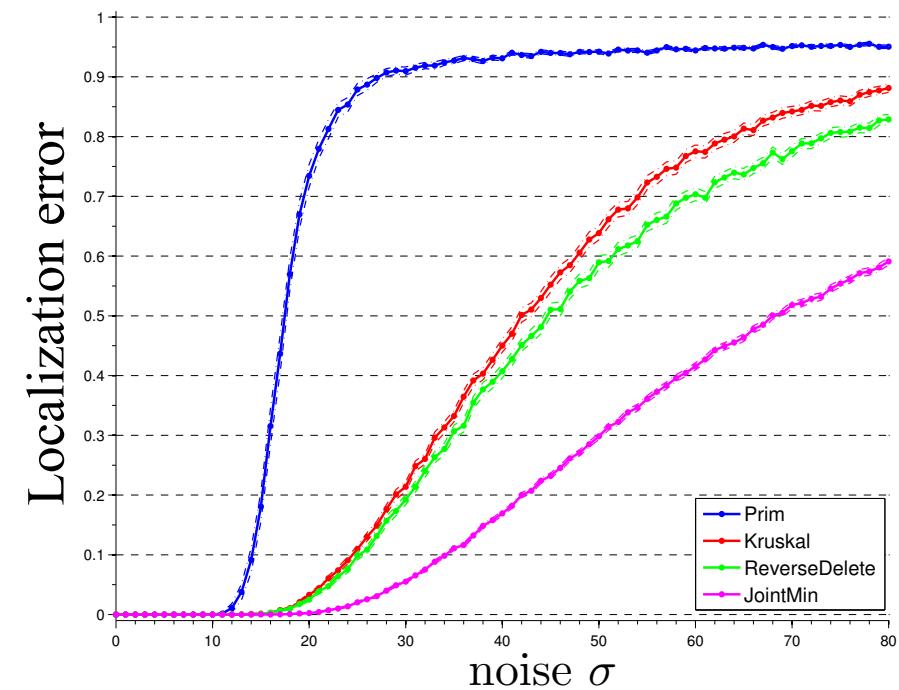
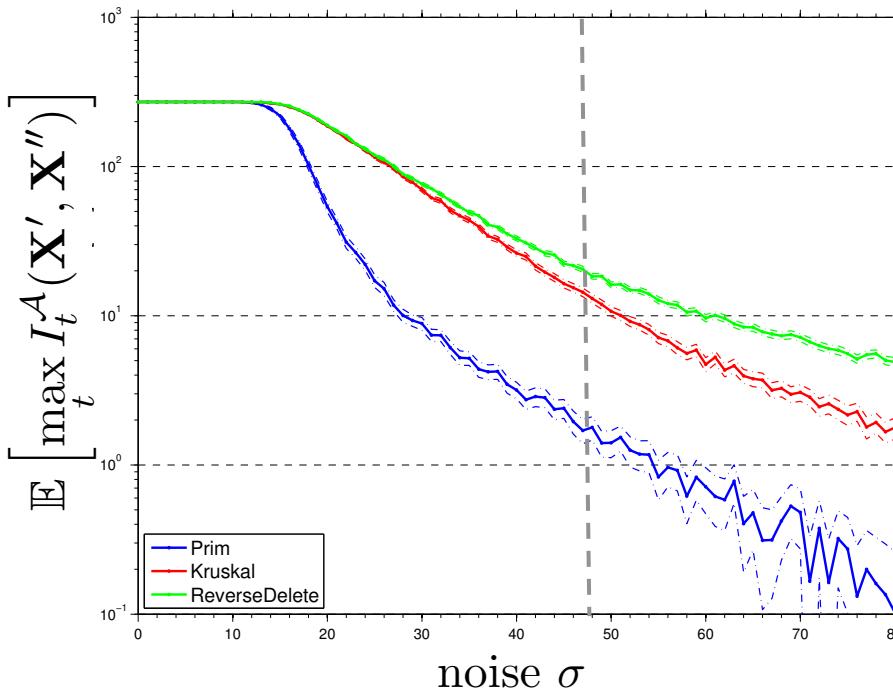
$$L = M_{\text{deg}}^X - M_{\text{adj}}^X$$

- Calculate the cofactor of  $L$  after  $t$  steps for the effective  $X(t)$  where selected edges are contracted (Prim, Kruskal) or removed (Reverse-Delete).

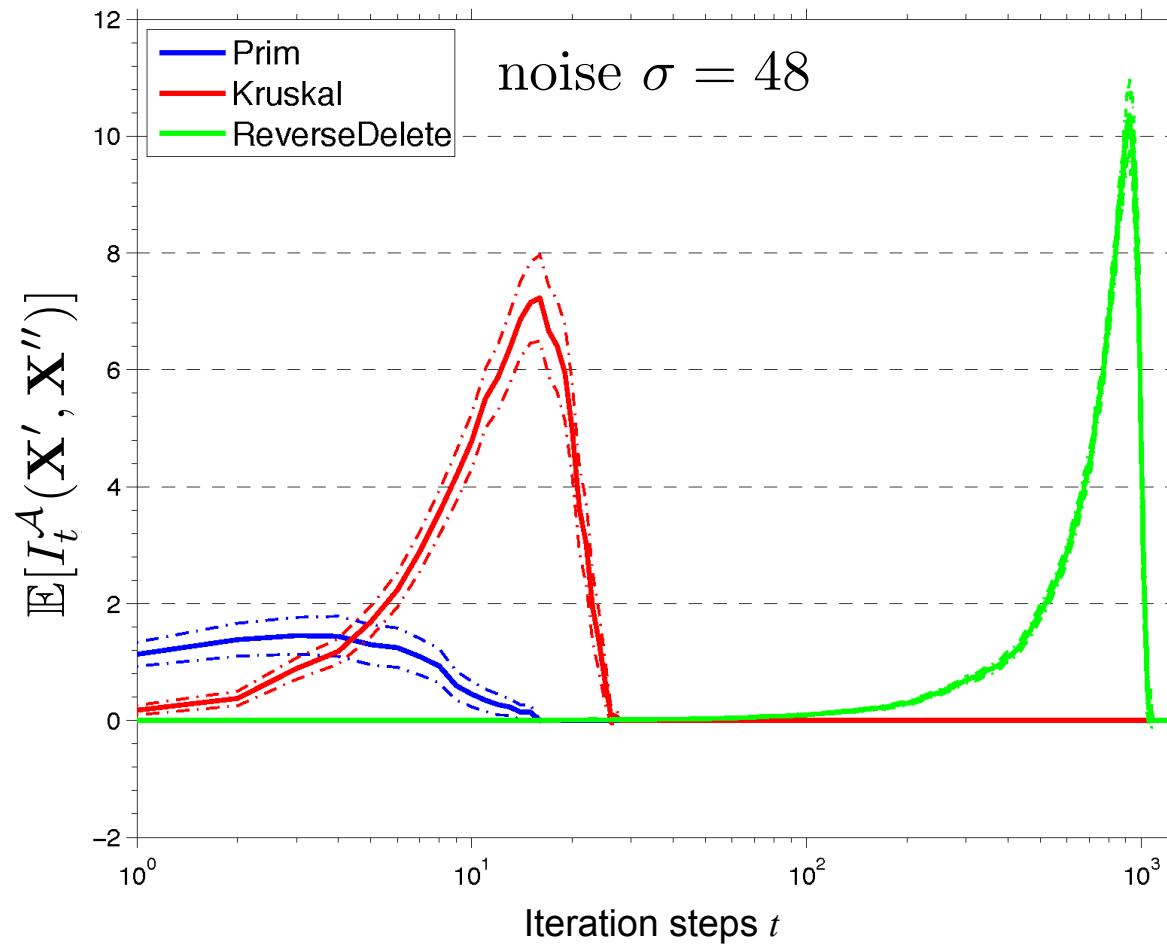
# Algorithmic informativeness

- Hierarchical graph generation:

- ground truth graph: 50 vertices, i.i.d. normal weights  $\mathcal{N}(100, 100)$
- Additive Gaussian noise  $\mathcal{N}(0, \sigma^2) \Rightarrow \mathbf{X}', \mathbf{X}''$



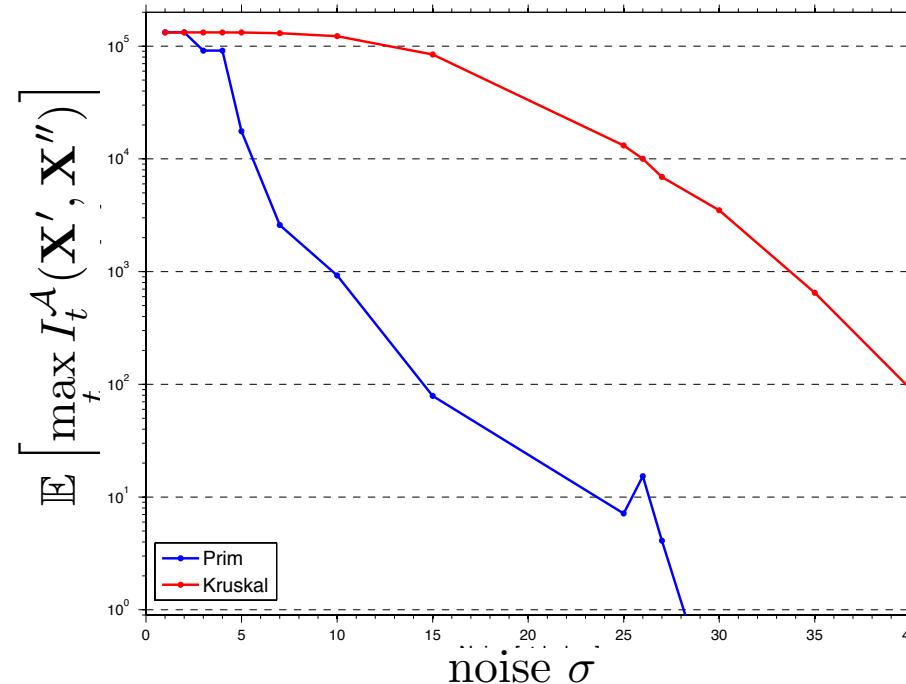
# Dynamics of algorithmic informativeness



# Informativeness of ASTs with $10^4$ vertices

- Hierarchical graph generation:

- ground truth graph:  $10^4$  vertices, i.i.d. normal weights  $\mathcal{N}(100, 100)$
- Additive Gaussian noise  $\mathcal{N}(0, \sigma^2)$



# Conclusion

- **Quantization:** Noise quantizes mathematical structures (hypothesis classes) => symbols
- **Coding** with these symbols defines a **generalization capacity for algorithms**  
⇒ **Quantization** of hypothesis class measures **structure specific information** in data.
- How to relate **statistical complexity** to algorithmic or **computational complexity** ?