$Probabilistic \ Graphical \ Models$

Jerónimo Hernández-González

Outline

- Structural learning: why?
- Structural learning based on scoring functions
- Structural learning based on conditional independence tests
- Structural learning for supervised classification

Model learning

Learning algorithm

Approximate P^* by learning a PGM, M, from data $D = \{x^1, \dots, x^N\}$ which is assumed to be i.i.d. sampled from P^*

$$A:D\to M\equiv (G,\Theta)$$

We want to extract information from D about P^* to encode in:

- ▶ The structure of the PGM, G (Structural learning):
 - ▶ NP-complete combinatorial optimization problem
 - ► Efficient heuristics: Local search, genetic algorithms, ...
- \triangleright The parameters of the PGM, Θ (Parametric learning)
 - ** It might be complemented with domain expert information

Structural learning Importance of the right structure

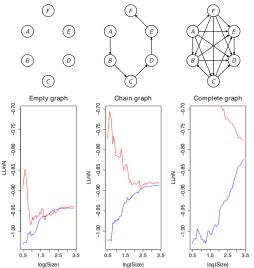
Missing an edge

- Incorrect independences?
- ► Reduces no. parameters
- ► Might not learn the real distribution *P**
- ► Tends to better generalization

Adding an extra edge

- Spurious dependencies?
- Increases no. parameters
- ► Might correctly learn *P**
- ► Tends to worse generalization

Fitting and generalization



Training vs. validation normalized Log-Likelihood: $nLL(\mathcal{M}; D) = \frac{LL(\mathcal{M}; D)}{v \cdot N}$

Fitting and generalization

Fitting - training LogLikelihood: $nLL(A(D_{tr}) = \mathcal{M}; D_{tr})$

- ► Tends to increase with number of parameters
- ► Tends to decrease with number of training cases

Same data is used for model training and scoring

Flexibility (model expressiveness) increases with no. parameters

Generalization - validation LogLikelihood: $nLL(\mathcal{M}; D_{va})$

- ▶ (Upper) constrained by the number of parameters
- ► Tends to increase with number of training cases

Fitting is an upper bound

Note that generalization is unknown!! (might be estimated, e.g., by CV)

Fitting and generalization

Overfitting

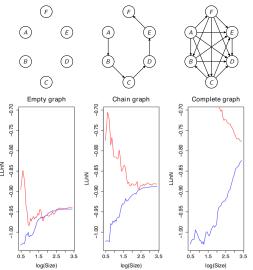
$$nLL(\mathcal{M}, D_{tr}) - nLL(\mathcal{M}, D_{va})$$

- Fitting and generalization, unbalanced
- Need to learn too many parameters with not enough training samples

Trade off!

Number of parameters vs. number of training cases

Fitting and generalization



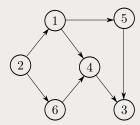
Training vs. validation normalized Log-Likelihood: $nLL(\mathcal{M}; D) = \frac{LL(\mathcal{M}; D)}{v \cdot N}$

Exercise

Counting parameters

Let $X_1, ..., X_6$ be binary random variables. Calculate the increase in the no. parameters after the addition of the edges:

- **▶** (2,5)
- **▶** (2, 4)



What?

Find the best structure (which fixes the no. parameters) with a dataset D sampled from a distribution of interest, P^* .

» Note the constrain relationship

When domain expertise cannot provide it

Why?

- Because we need a model to answer future queries
- Because we are interested into structure discovery: the structure is a goal itself

How? A NP-complete problem in the general case

- Based on a scoring function
- ► Based on conditional independence tests
 - » Exact polynomial algorithms in specific conditions. Otherwise: heuristic search.

 $Probabilistic \ Graphical \ Models$

Jerónimo Hernández-González

General formulation

1. Define a scoring function that evaluates the ability of a structure to describe the observed data

E.g.: Likelihood, penalized likelihood, ...

Search for the structure that maximizes the value of the scoring function

E.g.: Hill-climbing search, Genetic algorithms, ...

General formulation

1. Define a scoring function that evaluates the ability of a structure to describe the observed data

E.g.: Likelihood, penalized likelihood, ...

Search for the structure that maximizes the value of the scoring function

E.g.: Hill-climbing search, Genetic algorithms, ...

Scoring function: Likelihood score

Objective: Maximum likelihood learning

Select the model \mathcal{M} that maximizes:

$$L(\mathcal{M}; D) = p(D; \mathcal{M}) = \prod_{\mathbf{x} \in D} p_{\mathcal{M}}(\mathbf{x}) = \prod_{\mathbf{x} \in D} \prod_{i=1}^{\mathbf{v}} p(x_i | \mathbf{pa}_i; \Theta_i)$$

or

$$\log L(\mathcal{M}; D) = N \cdot \sum_{i=1}^{v} -H(X_i) + MI(X_i; PA_i)$$

For the purpose of model selection (structure comparison):

$$rg \max_{\mathcal{M}} \log L(\mathcal{M}; D) \propto rg \max_{\mathcal{M}} \sum_{i=1}^{v} \mathrm{MI}(X_i; PA_i)$$

» The score of a structure is $score(\mathcal{G}; D) = máx_{\Theta} score((\mathcal{G}, \Theta); D)$

Likelihood score

► If the likelihood is used as scoring function, it reduces to the sum of mutual information for each factor

$$score_L(\mathcal{G}; D) = N \cdot \sum_{i=1}^{v} -H(X_i) + MI(X_i; PA_i)$$

 $\approx \sum_{i=1}^{v} MI(X_i; PA_i)$

Mutual information quantifies the **strength** of the dependence relationship $X_i \not\perp \!\!\! \perp PA_i$

Scoring function: Likelihood score

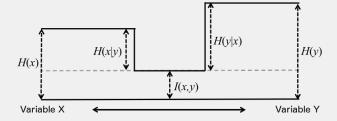
Likelihood score

Entropy

$$H(X) = -\sum_{x \in X} p(x) \log p(x)$$

Mutual Information

$$MI(X; Y) = \sum_{x,y} p(x,y) \log \frac{p(x,y)}{p(x)p(y)}$$
$$= H(Y) - H(Y|X) = H(X) - H(X|Y)$$



Scoring function: Likelihood score

Pros: Additively decomposable

$$score(\mathcal{G}; D) = \sum_{i=1}^{v} local_score(\mathcal{G}_{X_i}; D)$$

- Local changes in the graph only affect locally to the score
- Efficient search heuristics

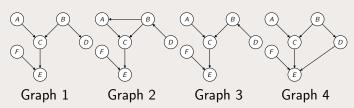
Cons: Monotone increasing

- ▶ Mutual information, MI(X; Y), is always ≥ 0
 - Only MI(X; Y) = 0, if X, Y are independent in the empirical distribution
- By adding edges, almost always the scoring value increases
- ► The complete graph maximizes this score → Overfitting!!

Exercise

Likelihood score

Which statement about the likelihood scores of these graphs is true?



- a) $Score_L(G_1; D) = Score_L(G_3; D)$, for every dataset D
- b) $Score_L(G_1; D) \geq Score_L(G_4; D)$, for every dataset D
- c) $Score_L(G_2; D) \geq Score_L(G_3; D)$, for every dataset D
- d) $Score_L(G_4; D) \geq Score_L(G_2; D)$, for every dataset D

Trade-off in overfitting (rev.)

Number of parameters vs. number of training cases

Avoiding overfitting

Restrict the hypothesis space by limiting the complexity of \mathcal{M} :

Explicitly: To control it.

E.g.:

- Set a limit on the no. parents
- Set a limit on the no. parameters as a function of no. cases
- Implicitly: To penalize it.

E.g.:

- Add a penalty per each additional edge
- A Bayesian score that averages the score of all the parameters

Scoring function: Penalized log-likelihood score

Penalized log-likelihood score

$$score(G; D) = score_L(G; D) - f(dim(G))$$

Add a penalization term which depends on the complexity of the graph \mathcal{G} . E.g.:

- No. parameters
- No. edges

Scoring function: BIC Score

BIC Score

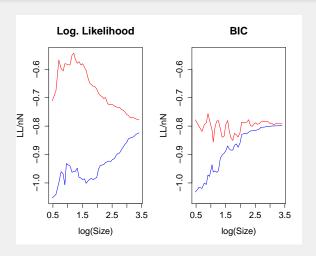
Logarithmic penalization of the number of parameters

$$score_{BIC}(\mathcal{G}; D) = score_{L}(\mathcal{G}; D) - \frac{\log N}{2} \# param(\mathcal{G})$$

- ► As N grows, more emphasis is given to fit to data
- Asymptotically,
 - Required edges are added due to linear growth of likelihood term vs. logarithmic growth of complexity penalization term
 - Edges with spurious contribution to likelihood are left out
 - Any *I*-equivalent structure of the true \mathcal{G}^* maximizes the score

Scoring function: BIC Score

Fitting and generalization

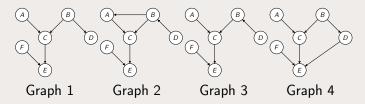


Training vs. validation normalized Log-Likelihood

Exercise

BIC score

Which statement about the BIC scores of these graphs is true?



- a) $Score_{BIC}(G_1; D) = Score_{BIC}(G_3; D)$, for every dataset D
- b) $Score_{BIC}(G_1; D) \geq Score_{BIC}(G_4; D)$, for every dataset D
- c) $Score_{BIC}(G_2; D) \neq Score_{BIC}(G_3; D)$, for every dataset D
- d) $Score_{BIC}(G_1; D) \geq Score_{BIC}(G_2; D)$, for every dataset D

Scoring function: Bayesian Score

Bayesian Score

Bayesian learning assumes a prior to obtain the posterior after observing D:

$$P(\mathcal{G}|D) \propto P(D|\mathcal{G})P(\mathcal{G})$$

We can define a score as:

$$score_B(G; D) = P(D|G)P(G)$$

where

$$P(D|\mathcal{G}) = \int P(D|\mathcal{G}, \Theta_{\mathcal{G}}) P(\Theta_{\mathcal{G}}|\mathcal{G}) d\Theta_{\mathcal{G}}$$

** In the general case, this integral is difficult to assess. Easier if conjugate priors are used: e.g., Dirichlet-Multinomial

Scoring function: Bayesian Score

The prior over structures, P(G)

- Uniform (uninformative)
- ▶ Penalize no. edges $P(\mathcal{G}) \propto c^{\#edges(\mathcal{G})}$ with 0 < c < 1
- ► Penalize no. parameters

General formulation

1. Define a scoring function that evaluates the ability of a structure to describe the observed data

E.g.: Likelihood, penalized likelihood, ...

2. Search for the structure that maximizes the value of the scoring function

E.g.: Hill-climbing search, Genetic algorithms, ...

Search method: Finding the tree with optimal score

Score equivalence

It means that the $score(X \to Y) = score(Y \to X)$, and Equivalent *I*-structures \Longrightarrow Same score $score_L$, $score_{BIC}$, and $score_B$ satisfy score equivalence

Optimal forest structure

Using a decomposable score,

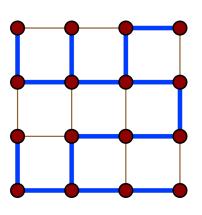
- 1. Build the complete graph and weigh each edge (X, Y) with $\max(score(X \to Y), 0)$
- 2. Find the Maximum Spanning Tree on that graph
- 3. Remove edges with weight 0

Search method: Minimum Spanning Tree

Minimum Spanning Tree

Given a connected graph $\ensuremath{\mathcal{U}}$ with weighted undirected edges,

a spanning tree of \mathcal{U} is a **tree subgraph** with all the vertices of \mathcal{U} and the minimum possible number of edges



Search method: Minimum Spanning Tree

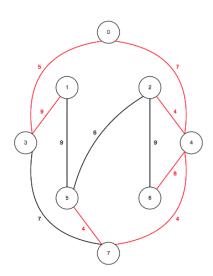
Minimum Spanning Tree

Given a connected graph ${\cal U}$ with weighted undirected edges,

a spanning tree of $\mathcal U$ is a **tree subgraph** with all the vertices of $\mathcal U$ and the minimum possible number of edges

a minimum spanning tree is a **spanning** tree of $\mathcal U$ having minimum weight

If $\mathcal U$ is unconnected, a minimum spanning forest is the union of a MST for each connected component



Search method: Minimum Spanning Tree

Kruskal's algorithm

Given an undirected edge-weighted graph $\mathcal U$ with n nodes,

- 1. Let L be the sorted list of edges of \mathcal{U} by increasing weight
- 2. Set $\mathcal{T} = \emptyset$, where \mathcal{T} is the subgraph of the MST
- 3. Pull the top edge from L and add it to T
- 4. Remove from L any edge that forms a loop in $\mathcal T$
- 5. If \mathcal{T} has n-1 edges, return \mathcal{T} (a tree) Elself $L=\emptyset$, return \mathcal{T} (a forest) Otherwise, go to step 3

https://www.cs.usfca.edu/~galles/visualization/Kruskal.html

Exercise

Recovering directionality

After we find an undirected spanning tree, which is the most efficient way to transform it into a directed spanning tree?

- a) Evaluate all possible directions for the edges (at most, 2^n possible sets of edge directions) by iterating over them (within $O(2^n)$ time).
- b) Exploit score decomposability to evaluate all possible directions for the edges (within O(n) time).
- c) Pick any arbitrary direction for each edge (within O(n) time). Due to score equivalence, all possible directed versions of the optimal undirected spanning forest have the same score.
- d) Pick any arbitrary root, and direct all edges away from it (within O(n) time).

Search method: Finding the tree with optimal score

Score equivalence

It means that the $score(X \to Y) = score(Y \to X)$, and Equivalent *I*-structures \Longrightarrow Same score $score_L$, $score_{BIC}$, and $score_B$ satisfy score equivalence

Optimal forest structure

Using a decomposable score,

- 1. Build the complete graph and weigh each edge (X, Y) with $\max(score(X \to Y), 0)$
- 2. Find the Maximum Spanning Tree on that graph
- 3. Remove edges with weight 0

Finding the maximal scoring network with at most k > 1 parents is NP-hard!

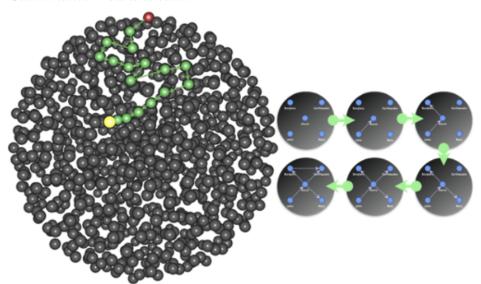
Search method: **Heuristic** search

Local search

- ▶ Based on the concept of neighborhood: each graph $\mathcal G$ has a few neighbor graphs $\mathcal G'$
 - Defined by operators such as: edge addition, removal or reversal
- ➤ Search technique

 Greedy hill-climbing, Best first search, Genetic, Simulated annealing, ...
- ► Efficient: Polynomial complexity
- Suboptimal solutions: Local optima

Search method: **Heuristic** search



Search method: **Heuristic** search

Greedy hill climbing

- 1. Start with an initial network. E.g.,
 - the empty or a random network
 - best tree
 - prior knowledge
- 2. Repeat at each iteration:
 - 2.1 Calculate the score^(**) of any possible structure obtained by a single local change. E.g.,
 - add an edge
 - remove an edge
 - reverse an edge
 - 2.2 Apply the change that most improves the score

Until no change improves the score

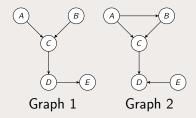
Local maxima reached

Thanks to $score^{(**)}$ decomposability, we can assess the scoring value by reevaluating only the part affected by the local change

Exercise

Calculating Likelihood Differences

We need to choose between these two graphs with likelihood score:



What is $Score_L(G_1; D) - Score_L(G_2; D)$ given dataset D of size N?

- a) $N \cdot [\mathrm{MI}_{\hat{p}}(C; A, B) + \mathrm{MI}_{\hat{p}}(D; C) + \mathrm{MI}_{\hat{p}}(E; D) \mathrm{MI}_{\hat{p}}(B; A) \mathrm{MI}_{\hat{p}}(D; C, E)]$
- b) $N \cdot [\mathrm{MI}_{\hat{p}}(D;C) + \mathrm{MI}_{\hat{p}}(E;D) \mathrm{MI}_{\hat{p}}(A;B) \mathrm{MI}_{\hat{p}}(D;C,E) \mathrm{H}_{\hat{p}}(A,B,C,D,E)]$
- c) $N \cdot [\operatorname{MI}_{\hat{p}}(D; C) + \operatorname{MI}_{\hat{p}}(E; D) \operatorname{MI}_{\hat{p}}(B; A) \operatorname{MI}_{\hat{p}}(D; C, E)],$
- d) $N \cdot [MI_{\hat{p}}(A; B) H_{\hat{p}}(A, B)]$
- e) $N \cdot \mathrm{MI}_{\hat{\rho}}(A; B)$

 $Probabilistic \ Graphical \ Models$

Jerónimo Hernández-González

What?

Find the best structure (which fixes the no. parameters) with a dataset D sampled from a distribution of interest, P^* .

» Note the constrain relationship

When domain expertise cannot provide it

Why?

- Because we need a model to answer future queries
- Because we are interested into structure discovery: the structure is a goal itself

How? A NP-complete problem in the general case

- Based on a scoring function
- ▶ Based on conditional independence tests
 - » Exact polynomial algorithms in specific conditions. Otherwise: heuristic search.

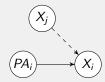
Based on conditional independence tests

PGMs as models that fulfill a set of independencies

Objective: Model the most important dependencies of p

Iterative procedure:

- ightharpoonup given the current structure $\mathcal G$
- ▶ test each possible inclusion of a new parent: should we move from $p(x_i|pa_i)$ to $p(x_i|pa_i \cup \{x_j\})$?



▶ i.e., check the statistical independence: $X_i \perp \!\!\! \perp X_j \mid PA_i$

Independence test: χ squared

χ squared: independence test, $X_A \perp \!\!\! \perp X_B$

A chi-square (χ^2) statistic measures the difference of the observed and expected frequencies of the outcomes of two sets of variables. Null hypothesis (H_0) assumes independence:

$$\forall \mathbf{x}, p(\mathbf{x}_{A,B}) = p(\mathbf{x}_A) \cdot p(\mathbf{x}_B)$$

Statistic:

$$X^{2} = \sum_{i=1}^{r_{A}} \sum_{j=1}^{r_{B}} \frac{(O_{ij} - E_{ij})^{2}}{E_{ij}}$$
 $X^{2} \sim \chi^{2}(v)$

i.e., X^2 is distributed according to a χ^2 distribution with ν degrees of freedom, where $\nu = (r_A - 1) \cdot (r_B - 1)$

Unlikelihood of the independence according to observed data

Independence test: χ squared

χ squared: independence test, $X_A \perp \!\!\! \perp X_B$

A chi-square (χ^2) statistic measures the difference of the observed and expected frequencies of the outcomes of two sets of variables. Null hypothesis (H_0) assumes independence:

$$\forall \mathbf{x}, p(\mathbf{x}_{A,B}) = p(\mathbf{x}_A) \cdot p(\mathbf{x}_B)$$

Alternative statistic (likelihood-ratio):

$$G^{2} = 2 \sum_{i=1}^{r_{A}} \sum_{j=1}^{r_{B}} O_{ij} \log \frac{O_{ij}}{E_{ij}} = 2N \cdot MI(\mathbf{X}_{A}; \mathbf{X}_{B})$$

$$G^2 \sim \chi^2(v)$$

i.e., G^2 is distributed according to a χ^2 distribution with v degrees of freedom, where $v = (r_A - 1) \cdot (r_B - 1)$

Unlikelihood of the independence according to observed data

Assuming independence: which are the expected counts?

	favor	indifferent	opposed	total
democrat				285
republican				215
total	202	150	148	500

Assuming independence: which are the expected counts?

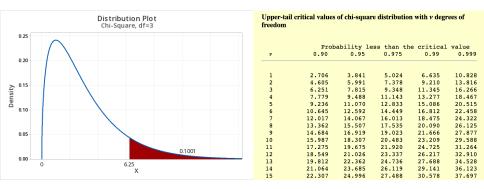
	favor	indifferent	opposed	total
democrat	$\frac{285}{500} \cdot \frac{202}{500} \cdot 500 = 115,14$	85,50	84,36	285
republican	86,86	64,50	63,64	215
total	202	150	148	500

Observed counts vs. Expected counts (given indep. assumption)

	favor	indifferent	opposed	total
democrat	138 - 115,14	83 - 85,50	64 - 84,36	285
republican	64 - 86,86	67 - 64,50	84 - 63,64	215
total	202	150	148	500

$$X^{2} = \frac{(138 - 115,14)^{2}}{115,14} + \frac{(83 - 85,50)^{2}}{85,50} + \frac{(64 - 84,36)^{2}}{84,36} + \frac{(64 - 86,86)^{2}}{86,86} + \frac{(67 - 64,50)^{2}}{64,50} + \frac{(84 - 63,64)^{2}}{63,64} = 22,152$$

$$X^{2} = \frac{(138 - 115,14)^{2}}{115,14} + \frac{(83 - 85,50)^{2}}{85,50} + \frac{(64 - 84,36)^{2}}{84,36} + \frac{(64 - 86,86)^{2}}{86,86} + \frac{(67 - 64,50)^{2}}{64,50} + \frac{(84 - 63,64)^{2}}{63,64} = 22,152$$



Unlikelihood of the independence according to observed data

Based on conditional independence tests

Conditional independence test, $X_i \perp \!\!\! \perp X_j | PA_i$

A chi-square (χ^2) which assumes (null hypothesis, H_0) conditional independence:

$$\forall \mathbf{x}, p(x_i, x_j | \mathbf{x}_{PA_i}) = p(x_i | \mathbf{x}_{PA_i}) \cdot p(x_j | \mathbf{x}_{PA_i})$$

Statistic:

$$X^{2} = \sum_{l=1}^{r_{i}} \sum_{m=1}^{r_{j}} \sum_{k=1}^{r_{PA_{i}}} \frac{(O_{lmk} - E_{lmk})^{2}}{E_{lmk}} \text{ with } E_{lmk} = N_{\cdot mk} N_{l \cdot k} / N_{\cdot \cdot k}$$

$$X^2 \sim \chi^2(v)$$

i.e., X^2 is distributed according to a χ^2 distribution with v degrees of freedom, where $v=(r_A-1)\cdot(r_B-1)\cdot r_C$

Unlikelihood of the independence according to observed data

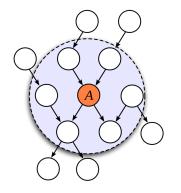
Independence test: χ squared

χ squared: independence test, $\textbf{\textit{X}}_{A} \perp \!\!\! \perp \textbf{\textit{X}}_{B}$

Results depend on the magnitude of the difference between actual and observed frequencies (X^2 and/or G^2), the degrees of freedom (ν), and the sample size (N).

- The degrees of freedom increase exponentially with the number of parents: $(r_i 1)(r_j 1)(r_{PA_i})$ The test becomes more demanding
- ► The difference X² (and G²) increases linearly with the sample size N
 The test becomes less demanding

Specific approaches: Markov blanket



Markov blanket

Set of nodes including parents, children, and other parents of its children.

The Markov blanket renders a node independent of the rest of the network.

The joint distribution of the variables in its Markov blanket is sufficient for calculating the distribution of a node.

 $Probabilistic \ Graphical \ Models$

Jerónimo Hernández-González

A generative approach

Advantages of $p_M(x, c)$ (vs. of $p_M(c|x)$)

- Flexibility
- ▶ Discover relationships between vars.: $p(x_A|c)p(x_B|c)p(c)$
- Incorporate a priori knowledge: Bayesian statistics
- ▶ Obtain the conditional distribution $p_M(c|\mathbf{x})$
- ▶ Deal with missing values, outliers, ...
- ▶ Rejection region, $p_M(c|\mathbf{x}) > t$

Disadvantages of $p_M(x, c)$ (vs. of $p_M(c|x)$)

- Harder problem
- ▶ Models irrelevant information, $p_M(x)$, for classification
- Requires more parameters

Structures biased towards classification

Going discriminative: Modeling p(C|x) but not p(X)

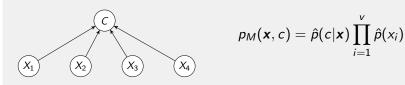
Capture specific knowledge about classification

Few parameters with highly discriminative information:

- ▶ Model the dependencies of the form $X_i \not\perp\!\!\!\perp C | \mathbf{X}_S$
- ▶ Model the most relevant dependencies like $X_A \not\perp \!\!\! \perp X_B \mid C$
- Avoid redundancy $\{C \perp \!\!\! \perp X_A | X_{V \setminus A}\}$

Structures biased towards classification: Informed Bayes

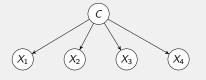
Informed Bayes



- Models all and only the important dependencies
- Exponential number of parameters w.r.t. v
- ► Given enough data resembles the Bayes classifier
- ▶ In normal scenarios, tends to overfit!!

Structures biased towards classification: Naive Bayes

Naive Bayes

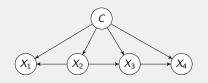


$$p_M(\mathbf{x},c) = \hat{p}(c) \prod_{i=1}^{r} \hat{p}(x_i|c)$$

- Assumes that features are independent given the class: $\{X_i \perp \!\!\! \perp X_i | C\}$
- ▶ Models the most important dependences: $\{X_i \not\perp\!\!\!\perp C | \mathbf{X}_S\}$
- No. parameters is linear with No. variables
- Worse generalization with large training sets
- Low risk of overfitting

Structures biased towards classification: Tree-augmented naive Bayes

Tree-augmented naive Bayes



$$p_{M}(\boldsymbol{x},c) = \hat{p}(c) \prod_{i=1}^{v} \hat{p}(x_{i}|x_{j},c)$$

- ► Generalization of naive Bayes
- Breaks the strong conditional independence assumption of NB
- Apart from C, each feature can have a parent (k = 1) among the rest of features

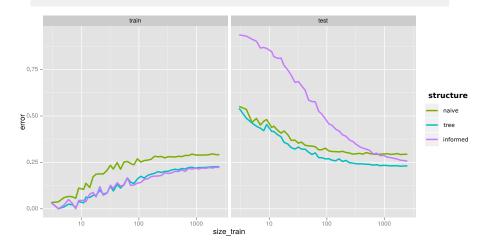
 MST among features with weights $CMI(X_i, X_j | C)$

Further generalization: k-dependence Bayesian classifier restricts the maximum number of parents to k

Structures biased towards classification

Fitting and generalization

Training (left) and test (right) classification error



Exercise

Hidden variable

Consider a generative naive Bayes model with 10 binary-valued features $\{X_1, \ldots, X_{10}\}$, but the class variable C is not observed. Still, C is strongly correlated with its children.

Suppose we learn a structure directly on $\{X_1, \ldots, X_{10}\}$ (without C). Which structure are we likely to learn if we use the likelihood score as the structure learning criterion?

- a) The empty network, i.e., a network consisting of only the variables but no edges between them.
- b) Some connected network over $\{X_1, \ldots, X_{10}\}$ which is not fully connected nor empty.
- c) A fully connected network, i.e., one with an edge between every pair of nodes.

Summary

Structural learning

- Usually done by scored based approaches and local search
- ▶ Many alternatives, some exact in controlled scenarios
- > Structures predefined specific for supervised classification

 $Probabilistic \ Graphical \ Models$

Jerónimo Hernández-González