

Introduction to Statistical Data Science

October 25, 2018

1 Lecture 1

Basic Rules of Probability

Frequently we will say $p(x) \propto f(x)$ for some non-negative function $f(x)$

Then we can conclude that:

$$p(x) = \frac{f(x)}{\sum_y f(y)}$$

For joint distribution,

$$\sum_x p(x) = \sum_x \sum_y p(x, y) = 1$$

Independence

If $p(x|y) = p(x)$ for all states of x and y , then the variables x and y are said to be independent as $x \perp\!\!\!\perp y$.

If x and y are independent, then x and y are uncorrelated. However, in general, x and y are uncorrelated, then cannot conclude that x and y are independent.

Conditional Independence

$$\mathcal{X} \perp\!\!\!\perp \mathcal{Y} | \mathcal{Z}$$

$$p(\mathcal{X}, \mathcal{Y} | \mathcal{Z}) = p(\mathcal{X} | \mathcal{Z}) p(\mathcal{Y} | \mathcal{Z})$$

and

$$p(\mathcal{X} | \mathcal{Y}, \mathcal{Z}) = p(\mathcal{X} | \mathcal{Z})$$

Conditional independence does not imply marginal independence:

$$p(x, y) = \sum_z p(x|z)p(y|z)p(z) \neq \sum_z p(x|z)p(z) \sum_z p(y|z)p(z)$$

2 Lecture 2

Graphs

Definition:

A graph consists of nodes (vertexes) and undirected or directed links (edges) between nodes.

Path:

A path from X_i to X_j is a sequence of connected nodes starting at X_i and ending at X_j . (no direction)

Directed Acyclic Graph:

Graph in which by following the direction of the arrows a node will **never** be visited **more than once**.

Parents and Children:

X_i is a parent of X_j if there is a link from X_i to X_j . X_i is a child of X_j if there is a link from X_j to X_i .

Ancestors and Descendants:

The ancestors of a node X_i are the nodes with a directed path ending at X_i . The descendants of X_i are the nodes with a directed path beginning at X_i .

Undirected Graph:

Clique:

A clique is a fully connected subset of nodes.

Maximal Clique:

Clique which is not a subset of a larger clique.

Connected graph:

There is a path between every pair of vertices.

Connected components:

In a non-connected graph, the connected components are the connected-subgraphs.

Connectedness: Singly-connected

There is only one path from any node a to another node b

Multiply-connected

A graph is multiply-connected if it is not singly connected.

Belief Networks (Bayesian Networks)

A belief network is a **directed acyclic graph** in which each node is associated with the conditional probability of the node given its parents.

Processing the network

Firstly write the whole joint distribution such as:

$$p(A, R, E, B) = p(A|R, E, B)p(R|E, B)p(E|B)p(B)$$

Then, according to the assumption, remove some independent variable from the joint distribution. **It does matter that the order of joint distribution influence the processing.**

Uncertain Evidence

In soft/uncertain evidence the variable is in more than one state, with the strength of our belief about each state being given by probabilities. For example, if y has the states $dom(y) = \{\text{red, blue, green}\}$ the vector $(0.6, 0.1, 0.3)$ could represent the probabilities of the respective states

In the calculation, we can do this: Given $P(A = tr) = 0.7$

$$p(B = tr|\tilde{A}) = \sum_A p(B = tr|A)p(A|\tilde{A})$$

Independence

If C has more than one incoming link, then $A \perp\!\!\!\perp B$ and A is not conditional independent with B under C condition. In this case C is called collider. If C has at most one incoming link, then

$A \perp\!\!\!\perp B|C$ and A is not independent with B. In this case C is called non-collider.

d-connected/separated

X and Y are d-connected by Z if there is any path from X to Y that is not blocked by Z

If all of the paths are blocked then we say X and Y are d-separated by Z.

Markov Equivalence

skeleton

Formed from a graph by removing the arrows.

immorality

An immorality in a DAG is a configuration of three nodes, A,B,C such that C is a child of both A and B, with A and B not directly connected.

Markov Equivalence

Two graphs represent the same set of independence assumptions if and only if they have the same skeleton and the same set of immoralities.

BN representation

BN cannot represent whatever independence statements are present in p .

Fundamentally, the actual **numerical distribution** p **contains much more information than a graph can represent.**

Markov Network

A Markov Network is an undirected graph in which there is a potential (non-negative function) ψ defined on each maximal clique.

The joint distribution is proportional to the product of all clique potentials:

$$P(Y) = \frac{1}{Z} \prod_C \Psi_C(Y_C)$$
$$z = \sum_Y \prod_C \Psi_C(Y_C)$$

where the $\Psi_C(Y_C)$ is a potential function (strict positive). The potential function would usually indicated as

$$\Psi_C(Y_C) = \exp\{-E(Y_C)\}$$

For the image recovery, the probability of pixel $Pr(x)$ is only related to the sum of neighboring points and the y .

Iterated conditional Modes

To find the minimum of a function $f(x_1, \dots, x_D)$, firstly make a guess x_1^*, \dots, x_D^* . Then look at the single variable x_i keeping all others fixed. Then set $x_i^* = \operatorname{argmin}_{x_i} F(x_i)$. Then repeat this, sweeping through all variables, usually in a random order. Then repeat the whole process until convergence.

Boltzmann machine

Ising model

Given $x_i \in +1, -1$, then the joint distribution would be:

$$p(x_1, \dots, x_9) = \frac{1}{Z} \prod_{i \sim j} \phi_{ij}(x_i, x_j)$$

$$\phi_{ij}(x_i, x_j) = e^{-\frac{1}{2T}(x_i - x_j)^2}$$

Rule for independence in Markov Networks

Remove all links neighboring the variables in the conditioning set Z .

If there is no path from any member of x to any member of y , then x and y are conditionally independent given z .

Factor Graphs

A square node represents a factor (non negative function) of its neighboring variables. FGs are most commonly used for inference.

Markov Models

Time-Series

A time-series is an ordered sequence:

$$x_{a:b} = \{x_a, x_{a+1}, \dots, x_b\}$$

For the time series data, we need a model $p(v_{1:T})$. For the causal consistency, it is meaningful to consider the decomposition.

$$p(v_{1:T}) = \prod_{t=1}^T p(v_t | v_{1:t-1})$$

with the convention $p(v_t | v_{1:t-1}) = p(v_1)$ for $t = 1$.

Only the recent past is relevant:

$$p(v_t | v_1, \dots, v_{t-1}) = p(v_t | v_{t-L}, \dots, v_{t-1})$$

where $L \geq 1$ is the order of the Markov chain.

$$p(v_{1:T}) = p(v_1)p(v_2|v_1)p(v_3|v_2) \dots p(v_T|v_{T-1})$$

For a stationary Markov chain the transitions $p(v_t = s' | v_{t-1} = s) = f'(s', s)$ are time-independent. $p(v_t | v_{t-1})$ is the transition matrix.

Equilibrium distribution

$$\mathbf{p}_t = \mathbf{M}^{t-1} \mathbf{p}_1$$

where $t \rightarrow \infty$ is independent of initial distribution \mathbf{p}_1 , then \mathbf{p}_∞ is called the equilibrium distribution of the chain:

$$\mathbf{p}_\infty = \mathbf{M} \mathbf{p}_\infty$$

Due to the Markov convergence (transition matrix not change, finite states, transit to any states, not a simple loop), the matrix eigenvalue has 1 value at most. Therefore, it would converge.

Mixture of Markov models

$$p(v_{1:T}) = \sum_{h=1}^H p(h)p(v_{1:T}|h) = \sum_{h=1}^H p(h) \prod_{t=1}^T p(v_t|v_{t-1}, h)$$

Applying EM later.

Hidden Markov Models

$$p(h_{1:t}, v_{1:T}) = p(v_1|h_1)p(h_1) \prod_{t=2}^T p(v_t|h_t)p(h_t|h_{t-1})$$

where $p(h_t|h_{t-1})$ and $p(v_t|h_t)$ are constant through time.

Inference

Inference corresponds to using the distribution to answer question about environment.

Computational Efficiency

Inference can be computationally very expensive and we wish to characterize situation in which inferences can be computed efficiently.

For **singly-connected** graphical models, and certain inference questions, there (usually) exist **efficient algorithms** based on the concept of **message passing**.

In general, the case of **multiply-connected** models is computationally **inefficient**.