Bayesian Inference and Hypothesis Testing

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Conditional Probability and the Theorem of Bayes

Simple Bayesian hypothesis testing

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So we can reverse the order of conditioning, i.e. relate to the probability of A given B to that of B given A.

The cards problem

- 1. Print out a number of cards, with either [A|A], [A|B] or [B|B] on their sides.
- 2. If you have an A, what is the probability of an A on the other side?
- 3. Have the students perform the experiment with:
 - 3.1 Draw a random card.
 - 3.2 Count the number of people with A.
 - 3.3 What is the probability that somebody with an A on one side will have an A on the other?
 - 3.4 Half of the people should have an A?

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The prior and posterior probabilities

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Prior elicitation

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- All those that think the accused is guilty, raise their hand.
- Divide by the number of people in class
- ▶ Let us call this $P(H_1)$.
- This is a purely subjective measure!

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DNA test properties

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- ▶ What is your belief now that the suspect is guilty?

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- What is your belief that the people with the positive test are guilty?

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The posterior can always be updated with more data!



Python example

```
# the input to the function is the prior, the likelihood function
# Input:
# - prior for hypothesis 0 (scalar)
# - data (single data point)
# - likelihood[data][hypothesis] array unction
# Returns:
# - posterior for the data point (if multiple points are given,
def get_posterior(prior, data, likelihood):
                marginal = prior * likelihood[data][0] + (1 - prior) * likel
                posterior = prior * likelihood[data][0] / marginal
                return posterior
import numpy as np
prior = 0.9
likelihood = np.zeros([2, 2])
# pr of negative test if not a match
likelihood[0][0] = 0.9
# pr of positive test if not a match
likelihood[1][0] = 0.1
# pr of negative test if a match
                                                                                                                                                                <□ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ >
```

Types of hypothesis testing problems

Simple Hypothesis Test

Example: DNA evidence, Covid tests

- ightharpoonup Two hypothesese H_0, H_1
- $ightharpoonup P(D|H_i)$ is defined for all i

Multiple Hypotheses Test

Example: Model selection

- $ightharpoonup H_i$: One of many mutually exclusive models
- \triangleright $P(D|H_i)$ is defined for all i

Null Hypothesis Test

Example: Are men's and women's heights the same?

- $ightharpoonup H_0$: The 'null' hypothesis
- $ightharpoonup P(D|H_0)$ is defined
- ► The alternative is undefined

Problem definition

▶ Defining the models $P(D|H_i)$ incorrectly.

The garden of many paths

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- ▶ Using an "unreasonable" prior $P(H_i)$

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The garden of many paths

- Having a huge hypothesis space
- ► Selecting the relevant hypothesis after seeing the data

Bayesian Inference

- ▶ Model family $\{P_{\theta} | \theta \in \Theta\}$
- **Each** model θ assigns probabilities $P_{\theta}(x)$ to possible $x \in X$.
- ightharpoonup We also have a (subjective) prior distribution eta over the parameters.
- Given x, we calculate the posterior distribution

$$\beta(\theta|x) = \frac{P_{\theta}(x)\beta(\theta)}{\sum_{\theta' \in \Theta} P_{\theta'}(x)\beta(\theta')}, \qquad \text{(finite } \Theta)$$

$$\beta(\theta|x) = \frac{P_{\theta}(x)\beta(\theta)}{\int_{\Theta} P_{\theta'}(x)\beta(\theta')d\theta'}, \qquad \text{(continuous } \Theta)$$

$$\beta(B|x) = \frac{\int_{B} P_{\theta'}(x)d\beta(\theta)}{\int_{C} P_{\theta'}(x)d\beta(\theta)}, \qquad B \subset \Theta \qquad \text{(arbitrary } \Theta)$$

Alternative notation for different probability spaces

- ▶ The prior $\beta(\theta) = \mathbb{P}(\theta)$ and posterior $\beta(\theta \mid x) = \mathbb{P}(\theta \mid x)$ belief.
- ▶ The likelihood $P_{\theta}(x) = \mathbb{P}(x \mid \theta)$
- ▶ The marginal $\mathbb{P}_{\beta}(x) = \sum_{\theta} P_{\theta}(x)\beta(\theta)$.



Probabilistic machine learning

Setting

- ▶ Model family $\{P_{\theta} | \theta \in \Theta\}$
- ightharpoonup Prior β on Θ
- ightharpoonup Observations $x = x_1, \dots, x_t$

Maximum likelihood approach

- ▶ Model selection: $\theta_{ML}^*(x) = \arg\max_{\theta} P_{\theta}(x)$.
- ▶ Model prediction: $P_{\theta_{MI}^*(x)}(x_{t+1})$

Maximum a posteriori approach

- ▶ Model selection: $\theta_{MAP}^*(x) = \arg\max_{\theta} P_{\theta}(x)\beta(\theta)$.
- ▶ Model prediction: $P_{\theta_{MAP}^*(x)}(x_{t+1})$

Bayesian approach

- ▶ Posterior calculation: $\beta(\theta|x) = P_{\theta}(x)\beta(\theta)/\mathbb{P}_{\beta}(x)$
- ▶ Model prediction: $\mathbb{P}_{\beta}(x_{t+1}|x) = \sum_{\theta} P_{\theta}(x_{t+1})\beta(\theta|x)$



Differences between approaches

Maximum likelihood approach

- Ignores model complexity
- ► Is an optimisation problem

Maximum a posteriori approach

- Regularises model selection using the prior
- Can be seen as solving the optimisation problem

$$\max_{\theta} \ln P_{\theta}(x) + \ln \beta(\theta),$$

where the prior term $\ln \beta(\theta)$ acts as a regulariser.

Bayesian approach

- ► Does not select a single model
- Averages over all models according to their fit and the prior
- ▶ Does not result in an optimisation problem.



The n-meteorologists problem

- ▶ Consider *n* meteorological stations $\{\mu\}$ predicting rainfall.
- $ightharpoonup x_t \in \{0,1\}$ with $x_t = 1$ if it rains on day t.
- \blacktriangleright We have a prior distribution $\beta(\mu)$ for each station.
- lacktriangle At time t, station μ makes as a prediction $P_{\mu}(x_{t+1}|x_1,\ldots,x_t)$
- We observe x_{t+1} and calculate the posterior $\beta(\mu|x_1,\ldots,x_t,x_{t+1})$.

The marginal distribution

To take into account all stations, we can marginalise:

$$\mathbb{P}_{\beta}(x_{t+1} \mid x_1, \dots x_t) = \sum_{\mu} P_{\mu}(x_{t+1} | x_t) \beta(\mu)$$

The posterior

► Show that

$$\beta(\mu \mid x_1, \dots, x_{t+1}) = \frac{P_{\mu}(x_t \mid x_1, \dots, x_t)\beta(\mu \mid x_1, \dots, x_t)}{\sum_{\mu'} P_{\mu'}(x_t \mid x_1, \dots, x_t)\beta(\mu' \mid x_1, \dots, x_t)}$$

► How would you implement an ML or a MAP solution to this problem?



Sufficient statistics

A statistic f

This is any function $f: X \to S$ where

- X is the data space
- \triangleright S is an arbitrary space

Example statistics for $X = \mathbb{R}^*$ (the set of all real-valued sequences)

- ▶ The sample mean of a sequence $1/T \sum_{t=1}^{T} x_t$
- ► The total number of samples T

Sufficient statistic

f is sufficient for a family $\{P_{\theta}: \theta \in \Theta\}$ when

$$f(x) = f(x') \Rightarrow P_{\theta}(x) = P_{\theta}(x') \forall \theta \in \Theta.$$

If there exists a finite-dimensional sufficient statistic, Bayesian and ML learning can be done in closed form within the family.



Conjugate priors

Consider a parametrised family of priors $\mathcal B$ on Θ and a distribution family $\{P_\theta\}$ The pair is conjugate if, for any prior $\beta \in \mathcal B$, and any observation x, there exists $\beta' \in \mathcal B$ such that $\beta'(\theta) = \beta(\theta|x)$

Standard Parametric conjugate families

Prior	Likelihood	Parameters $ heta$	Observations x
Beta	Bernoulli	[0, 1]	$\{0,1\}^{T}$
Multinomial	Dirichlet	\triangle^n	$\{1,\ldots,n\}^T$
Gamma	Normal	\mathbb{R},\mathbb{R}	\mathbb{R}^T
Wishart	Normal	\mathbb{R}^n , $\mathbb{R}^{n \times n}$	$\mathbb{R}^{n \times T}$

The Simplex $\mathbb{\Delta}^n = \{ \boldsymbol{\theta} \in [0,1]^n : \|\boldsymbol{\theta}\|_1 \}$ is the set of all *n*-dimensional probability vectors.

Extensions

- Discrete Bayesian Networks.
- Linear-Gaussian Models (i.e. Bayesian linear regression)
- Gaussian Processes.



Beta-Bernoulli



Definition of the Bernoulli distribution

If $x_t \mid \theta \sim \text{Bernoulli}(\theta)$. $\theta \in [0, 1]$, $x_t \in \{0, 1\}$ and:

$$P_{\theta}(x_t = 1) = \theta$$

Definition of the Beta density

If $\theta \sim \text{Beta}(\alpha_1, \alpha_0)$, $\alpha_0, \alpha_1 > 0$ and

$$p(\theta|\alpha_1,\alpha_0) \propto \theta^{\alpha_1-1} (1-\theta)^{\alpha_0-1}$$

Beta-Bernoulli conjugate pair

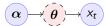
- $\triangleright x_t \mid \theta \sim \text{Bernoulli}(\theta).$

Then, for any $x = x_1, \dots, x_T$, the posterior distribution is

 $\blacktriangleright \theta \mid x \sim \text{Beta}(\alpha_1 + \sum_t x_t, \alpha_0 + T - \sum_t x_t).$



Dirichlet-Multinomial



Definition of the Multinomial distribution

If $x_t \mid \boldsymbol{\theta} \sim \operatorname{Mult}(\boldsymbol{\theta})$, with $\boldsymbol{\theta} \in \mathbb{\Delta}^n$ and $x_t \in \{1, \ldots, n\}$ and:

$$P_{\theta}(x_t = i) = \theta_i$$

Definition of the Dirichlet density

If $\boldsymbol{\theta} \sim \operatorname{Dir}(\boldsymbol{\alpha})$, with $\boldsymbol{\alpha} \in \mathbb{R}^n_+$ then

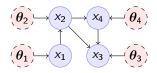
$$p(heta|lpha) \propto \prod_i heta_i^{lpha_i-1}$$

Dirichlet-Multinomial conjugate pair

- \bullet $\theta \sim \text{Dir}(\alpha)$.
- $\triangleright x_t \mid \theta \sim \text{Bernoulli}(\theta).$

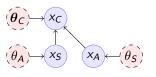
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Discrete Bayesian Networks



- ▶ A directed acyclic graph (DAG) defined on variables $x_1, ..., x_n$ with each x_n taking a finite number of values,
- ▶ Let S_i be the indices corresponding to parent variables of x_i .
- $\triangleright x_i \mid \theta_i, x_{S_i} = k \sim \text{Mult}(\theta_{i,k}).$

Example: Lung cancer, smoking and asbestos

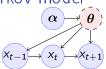


$$P_{\theta_A}(x_A = 1) = \theta_A \qquad (1)$$

$$P_{\theta_S}(x_S=1)=\theta_S \qquad (2)$$

$$P_{\theta_C}(x_C = 1 \mid X_A = j, X_S = k) = \theta_{C,j,k}$$
 (3)

Markov model



A Markov model obeys

$$\mathbb{P}_{\theta}(x_{k+1}|x_k,\ldots,x_1) = \mathbb{P}_{\theta}(x_{k+1}|x_k)$$

i.e. the graphical model is a chain. We are usually interested in homogeneous models, where

$$\mathbb{P}_{\boldsymbol{\theta}}(x_{k+1} = i \mid x_k = j) = \theta_{i,j} \qquad \forall k$$

Inference for finite Markov models

- ▶ If $x_t \in [n]$ then $x_{t+1} \mid \theta, x_t = i \sim \text{Mult}(\theta_i), \theta_i \in \mathbb{A}^n$
- ▶ Prior $\theta_i \mid \alpha \sim \text{Dir}(\alpha)$ for all $i \in [n]$.
- ▶ Posterior $\theta_i \mid x_1, \dots, x_t, \alpha \sim \text{Dir}(\alpha^{(t)})$ with

$$\alpha_{i,j}^t = \alpha_{i,j} + \sum_{k=1}^t \mathbb{I}\left\{x_k = i \land x_{k+1} = j\right\}, \qquad \alpha^0 = \alpha.$$