# Probabilities and Statistical Estimation Chapter 3

University of Amsterdam

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#### Probability distributions

- Introduction
- Mean, variance and Covariance
- Gaussian and  $\chi^2$

#### 2 Statistical Estimation

- Maximum Likelihood
- Maximum a Posteriori
- Bayesian inference

#### 3 Kalman Filter

- 4 Fisher Information Matrix
- 5 Akaike Information Criterion

# Introduction

#### Frequentist approach

- "The data comes from a distribution, let us find as best as we can which distribution that was"
- Find "estimators" for parameters, and try to figure out how good these estimators are.

Bayesian approach

- "The data could have come from any number of distributions, let us find what those distributions could have been, and how likely they are"
- Using Bayes' rule does not make an approach Bayesian

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Mean, variance and Covariance

# A few definitions (scalar variables)

Expectation

$$E[f(x)] \triangleq \int_{-\infty}^{\infty} f(x)p(x)dx$$

• Mean (Expectation of x):

$$E[x] = \int_{-\infty}^{\infty} x p(x) dx$$
 (2)

Variance

$$V[x] \triangleq E[(x - E[x])^2] = \int_{-\infty}^{\infty} (x - E[x])^2 p(x) dx$$
 (3)

Covariance

$$V[x,y] \triangleq E[(x - E[x])(y - E[y])$$
(4)

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Mean, variance and Covariance

## Multivariate version

Expectation

$$E[f(\mathbf{x})] \triangleq \int_{\mathbb{R}^n} f(\mathbf{x}) p(\mathbf{x}) \mathrm{d}\mathbf{x}$$
 (5)

• Mean (Expectation of x):

$$E[\mathbf{x}] = \int_{\mathbb{R}^n} \mathbf{x} p(\mathbf{x}) d\mathbf{x}$$
 (6)

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Covariance matrix

$$V[\mathbf{x}] \triangleq E[(\mathbf{x} - E[\mathbf{x}])(\mathbf{x} - E[\mathbf{x}])^{\top}]$$
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#### Law of large numbers

Law of large numbers:

As N grows large,

$$\frac{1}{N}\sum_{i=1}^{N}\mathbf{x}_{i} \to E[\mathbf{x}]$$
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Change of variables

If **x** is an *n*-vector and  $\mathbf{y} = \mathbf{A}\mathbf{x}$  is an *m*-vector, for an arbitrary *mn*-matrix  $\mathbf{A}$ 

$$E[\mathbf{y}] = \mathbf{A}E[\mathbf{x}] \tag{9}$$
$$V[\mathbf{y}] = \mathbf{A}V[\mathbf{x}]\mathbf{A}^{\top} \tag{10}$$

If we perform a functional transformation of variables,  $\mathbf{y} = \mathbf{y}(\mathbf{x})$ , then by defining  $\mathbf{x} = \bar{\mathbf{x}} + \Delta \mathbf{x}$  and  $\mathbf{y} = \bar{\mathbf{y}} + \Delta \mathbf{y}$ , we obtain to a first approximation

$$\mathbf{\bar{y}} = \mathbf{y}(\mathbf{\bar{x}}) \quad \Delta \mathbf{y} = \frac{\partial \mathbf{y}}{\partial \mathbf{x}} \Big|_{\mathbf{\bar{x}}} \Delta \mathbf{x} \quad V[\mathbf{y}] = \frac{\partial \mathbf{y}}{\partial \mathbf{x}} \Big|_{\mathbf{\bar{x}}} V[\mathbf{x}] \frac{\partial \mathbf{y}}{\partial \mathbf{x}} \Big|_{\mathbf{\bar{x}}}^{\top}$$
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# Principal Component Analysis

PCA is probably the most well-known method for dimensionality reduction.

- Also known as the Karhunen-Loève transform
- Orthogonal projection of the data into a lower-dimensional subspace, so that the variance of the projected data is maximised
- Equivalently: linear projection that minimises the mean-squared distance between data points and their projection

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### Maximising the projected variance

Consider projecting on  $\mathbf{u}_1$ , with unit length for convenience.

- Each vector  $\mathbf{x}_n$  is then projected into  $\mathbf{u}_1^\top \mathbf{x}_n$
- $\bullet\,$  The mean of the projected data equals the projected mean  $u_1\bar{x}$
- To maximise the variance of the projected data, we maximise

$$\frac{1}{N}\sum_{n=1}^{N} (\mathbf{u}_{1}^{\top}\mathbf{x}_{n} - \mathbf{u}_{1}^{\top}\bar{\mathbf{x}}_{n})^{2} = \mathbf{u}_{1}^{\top}V[\mathbf{x}]\mathbf{u}_{1}$$
(12)

• Using a Lagrange multiplier to constrain  $\mathbf{u}_1^\top \mathbf{u}_1 = 1$ , we get

$$\mathbf{u}_1^\top V[\mathbf{x}] \mathbf{u}_1 + \lambda_1 (1 - \mathbf{u}_1^\top \mathbf{u}_1)$$
(13)

resulting in  $V[\mathbf{x}]\mathbf{u}_1 = \lambda \mathbf{u}_1$ . That is,  $\mathbf{u}_1$  is an eigenvector of  $V[\mathbf{x}]$  and the maximum is obtained for the largest eigenvalue.

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### Interesting properties

- The eigenvalues indicate the variance of the data along the orientation of the corresponding eigenvector
- Since the eigenvectors form an orthonormal basis, the data is uncorrelated along the projection orientations (3.24)



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Mean, variance and Covariance

# Local distributions

- Definition of manifold
- Definition of dimension / codimension
- Non-singular: codimension = 1
- Singular: codimension  $\neq 1$
- Tangent space / Normal space
- Local distribution: Assume that the distribution is sufficiently "localised", so that all observations can be approximated as lying in the tangent space of the manifold

### 3D rotation

Consider a rotation **R** as a random variable, representing a rotation  $\mathbf{\bar{R}}$  perturbed by some small noise  $\Delta \mathbf{R}$ :  $\mathbf{R} = \mathbf{\bar{R}} + \Delta \mathbf{R}$ .  $\mathbf{\bar{R}}$ , **R** and  $\Delta \mathbf{R}$  are rotations, so that we can write:

$$\mathbf{R} = (\mathbf{I} + \Delta \Omega \mathbf{I} \mathbf{I} + O(\Delta \Omega^2))\bar{\mathbf{R}}$$
(14)  
$$\bar{\mathbf{R}} + \Delta \mathbf{R} = \bar{\mathbf{R}} + \Delta \Omega \mathbf{I} \bar{\mathbf{R}} + O(\Delta \Omega^2)$$
(15)

To a first approximation:

$$\Delta \mathbf{R} = \Delta \Omega \, \mathbf{I} \, \bar{\mathbf{R}} \tag{16}$$

The covariance matrix of the rotation can then be defined as:

$$V[\mathbf{R}] = E[\Delta \Omega^2 \mathbf{I} \mathbf{I}^\top]$$
(17)

The eigenvector with largest associated eigenvalue then A Sapproximates the vector around which the rotation is most likely  Akaike Information Criterion

Gaussian and  $\chi^2$ 

# The Gaussian Distribution



$$p(\mathbf{x}) = \frac{1}{(2\pi)^{n/2} |\mathbf{\Sigma}|^{1/2}} \exp{-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^{\top} \mathbf{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})}$$
(18)

where we can check that

$$E[\mathbf{x}] = \boldsymbol{\mu} \qquad V[\mathbf{x}] = \boldsymbol{\Sigma} \tag{19}$$

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### Interesting properties

• The quantity

$$(\mathbf{x} - \boldsymbol{\mu})^{\top} \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})$$
 (20)

is the squared *Mahalanobis distance* of  $\mathbf{x}$  from the mean.

- Uncorrelated normally distributed variables are always independent
- If Σ is not full rank, we define the Gaussian distribution in the space spanned by the data
- The *central limit theorem* states that the sum of a sufficiently large number of independent random variables with finite variance converges to a normal distribution.

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If  $x_1, \ldots, x_r$  are r independent samples from  $\mathcal{N}(0, 1)$ , then

$$R = x_1^2 + \dots + x_r^2 \tag{21}$$

has a  $\chi^2$  distribution. Some nice properties:

- *E*[*R*] = *r*
- V[R] = 2r
- The mode of the distribution is at R = r 2
- $\bullet\,$  The sum of independent  $\chi^2$  variables is  $\chi^2$  distributed

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• For a multivariate random variable  $\mathbf{x} \sim \mathcal{N}(\mathbf{0}, \mathbf{\Sigma})$  with  $\mathbf{\Sigma}$  of rank *r*, the quadratic sum

$$R = \mathbf{x}^{\top} \mathbf{\Sigma}^{-1} \mathbf{x}$$
 (22)

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is  $\chi^2$  distributed with *r* degrees of freedom





A frequentist approach to reject hypotheses:

- Construct random variables  $R = x_1^2 + \cdots + x_r^2$  such that each  $x_i$  has zero mean if the hypothesis holds
- If the variables do not have zero mean, E[R] becomes larger
- The hypothesis is rejected with significance level a (confidence) *level* (1-a) if R falls in the region  $(\chi^2_{r,a},\infty)$

Note that this allows you to reject hypotheses, not to accept them!



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Maximum Likelihood

# Maximum Likelihood

#### Find the parameter by maximising the likelihood

$$I(\boldsymbol{\theta}) \triangleq p(\{\mathbf{y}\}|\boldsymbol{\theta})$$
(23)  
=  $\prod_{i} p(\mathbf{y}_{i}|\boldsymbol{\theta})$ (24)

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Maximum a Posteriori

# Maximum a Posteriori

The major problem with ML estimation is *overfitting*; learning the structure of the data extremely well, but performing poorly on new examples.

If we have prior knowledge, we can encode this in the model in the form of a prior probability distribution over the parameters, and update these with the observed data:

$$p(\theta|\mathbf{y}) = \frac{p(\mathbf{y}|\theta)p(\theta)}{p(\mathbf{y})}$$
(25)

where the marginal probability density  $p(\mathbf{y})$  is a constant. The MAP estimate is obtained by maximising  $p(\theta|\mathbf{y})$ . This reduces overfitting, but this is not Bayesian inference.

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Bayesian inference

### **Bayesian Inference**

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In Bayesian inference, we learn a distribution over parameters. We consider that all parameters we are not interested in are "nuisance parameters". To obtain the distribution over the quantity of interest,  $\theta_i$ , we *marginalise out* the other parameters:

$$p(\theta_i | \mathbf{y}) \propto \int_{\theta_{\neg i}} p(\mathbf{y} | \theta) \mathrm{d}\theta_{\neg i}$$
 (26)



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Bayesian inference

# "Statistical Estimation"

If we have a model  $\mathbf{y} = \mathbf{A}\mathbf{x} + \boldsymbol{\epsilon}$ , where  $\mathbf{A}$  is known and  $\boldsymbol{\epsilon}$  has a Gaussian distribution with known parameters, how do we find  $\mathbf{x}$ ?

Find the parameter by maximising the likelihood:

$$\ell(\boldsymbol{\theta}) \triangleq p(\mathbf{y}|\boldsymbol{\theta}), \text{ where } \boldsymbol{\theta} = \mathbf{x}$$
 (27)

$$= \frac{1}{(2\pi)^{n/2} |\mathbf{\Sigma}|^{1/2}} \exp{-\frac{1}{2} (\mathbf{y} - \mathbf{A}\mathbf{x})^{\top} \mathbf{\Sigma}^{-1} (\mathbf{y} - \mathbf{A}\mathbf{x})}$$
(28)

Ignoring constants and since  $exp(\cdot)$  is a monotonically increasing function, this is equivalent to minimising the Mahalanobis distance, resulting in:

$$\hat{\mathbf{x}} = (\mathbf{A}^{\top} \mathbf{\Sigma}^{-1} \mathbf{A})^{-1} \mathbf{A} \mathbf{\Sigma}^{-1} \mathbf{y}$$
(29)

and the error has a  $\chi^2$  distribution Intelligent Autonomous Systems

# Kalman Filter

Linear dynamical system defined as:

$$\mathbf{x}_{t+1} = \mathbf{A}_t \mathbf{x}_t + \mathbf{B}_t \mathbf{v}_t \tag{30}$$

$$\mathbf{y}_t = \mathbf{C}_t \mathbf{x}_t + \mathbf{w}_t \tag{31}$$

where  $\mathbf{A}_t$ ,  $\mathbf{B}_t$  and  $\mathbf{C}_t$  are fixed matrices;  $\mathbf{v}_t$  and  $\mathbf{w}_t$  are normally distributed with known means and covariances.

The filter updates its estimators  $\hat{\mathbf{x}}_t$  and  $V[\hat{\mathbf{x}}_t]$  at each time step by first estimating them given past observations, and updating it with the current observation.

The Kalman "smoother" additionally performs a backward pass to include information from the future in the estimate of  $x_t$ 

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# Fisher Information Matrix

Score:

$$\mathbf{I} = \nabla_{\boldsymbol{\theta}} \log p(\mathbf{x}; \boldsymbol{\theta}) \tag{32}$$

Fisher Information Matrix

$$\mathbf{J} = E[\mathbf{I}\mathbf{I}^{\top}] \tag{33}$$

can be written, if the log-likelihood is twice differentiable, as

$$\mathbf{J} = E[-\nabla_{\boldsymbol{\theta}}^2 \log p(\mathbf{x}; \boldsymbol{\theta})]$$
(34)

Intuitively: the more peaked the log-likelihood, the more informative the distribution

Cramér-Rao Lower Bound The Fisher Information Matrix provides a lower bound on the variance of an estimator of a parameter. If A othe bound is attained, the estimator is said to be *efficient*  Statistical Estimation

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# Akaike Information Criterion

The Akaike Information Criterion

$$AIC = 2m' - 2\sum_{i} \log p(\mathbf{x}_{i}; \hat{\boldsymbol{\theta}})$$
(35)

where m' is the rank of the fisher information matrix **J** 

- This penalises models that are too flexible, and optimises the expected likelihood of future data.
- Other information criteria are also commonly used, such as the BIC, which penalise complex models slightly differently. The BIC penalises a high number of parameters less as more data becomes available.

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