

## Non-Linear Signal Processing: Exercise 2

This exercise is based on C.M. Bishop: *Pattern Recognition and Machine Learning*, sections 1.4, 2.3.0-2.3.4 and appendix C. The objective of the exercise is to become familiar with the 2D normal distribution, the notion of covariance, and using projections on eigenvectors as features.

Print and comment on the figures produced by the software `main2a.m` to `main2e.m` as outlined below at the four **Checkpoints**.

### Multivariate normal Distribution

Let  $\mathbf{x}$  be a  $d$ -dimensional variable, i.e.  $\mathbf{x} = (x_1, x_2, \dots, x_d)^T$ . The probability of the variable  $\mathbf{x}$  lying in a region,  $\mathcal{A}$ , which is a subspace of  $\mathbf{R}^d$  is given by

$$p(\mathbf{x} \in \mathcal{A}) = \int_{\mathcal{A}} p(\mathbf{x}) d\mathbf{x}, \quad (1)$$

where  $p(\mathbf{x})$  is the probability density function of the variable  $\mathbf{x}$ .

In one dimension, the normal probability density function is given by

$$\mathcal{N}(x|\mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(x - \mu)^2}{2\sigma^2}\right), \quad (2)$$

where  $\mu$  and  $\sigma^2$  are the mean and variance respectively. In  $d$  dimensions, the general multivariate normal probability density function is given by

$$\mathcal{N}(\mathbf{x}|\boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{1}{\sqrt{(2\pi)^d |\boldsymbol{\Sigma}|}} \exp\left(-\frac{1}{2}(\mathbf{x} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1}(\mathbf{x} - \boldsymbol{\mu})\right), \quad (3)$$

where  $\boldsymbol{\mu}$  is a  $d$ -dimensional vector, and  $\boldsymbol{\Sigma}$  is a  $d \times d$  covariance matrix.

### 2D normal Distribution

Let  $\mathbf{x}$  be a 2-dimensional variable, so that  $d = 2$  in the above equations. Let  $\mathcal{D}$  be a set of  $N$  samples from  $\mathbf{x}$ , so that  $\mathcal{D} = \{\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N\}$ , where  $\mathbf{x}_i = (x_{i,1}, x_{i,2})^T$ ,  $i = 1, \dots, N$ .

It is then possible to construct a 2D histogram of the data-set,  $\mathcal{D}$ , by defining a Cartesian grid of small areas  $\mathcal{A}_{j,k}$ , where  $j = 1, \dots, M_1$  and  $k = 1, \dots, M_2$ . The histogram is then given by

$$n_{j,k} = \sum_{\mathbf{x}_i \in \mathcal{A}_{j,k}} 1, \quad j = 1, \dots, M_1, \quad k = 1, \dots, M_2, \quad (4)$$

with  $n_{j,k}$  denoting the number of observations of  $\mathbf{x}_i$  falling in the region  $\mathcal{A}_{j,k}$ . The normalized histogram is given by

$$\tilde{n}_{j,k} = \frac{n_{j,k}}{\sum_{j',k'} n_{j',k'}}. \quad (5)$$

If the union of all the areas  $\mathcal{A}_{j,k}$  includes all the samples in  $\mathcal{D}$ , equation (5) simplifies to

$$\tilde{n}_{j,k} = \frac{n_{j,k}}{N}. \quad (6)$$

The normalized histogram can be compared with the histogram approximation to the probability density function

$$P_{j,k} = \int_{\mathcal{A}_{j,k}} p(\mathbf{x}) d\mathbf{x}, \quad j = 1, \dots, M_1, \quad k = 1, \dots, M_2. \quad (7)$$

Alternatively the histogram can be converted to a normalized probability density, simply by dividing the normalized histogram bins with their corresponding areas  $\mathcal{A}_{j,k}$

$$p_{j,k} = \frac{\tilde{n}_{j,k}}{\mathcal{A}_{j,k}}. \quad (8)$$

Hereby, we obtain a model for the density that is constant over the area  $\mathcal{A}_{j,k}$  of each bin.

### Checkpoint 2.1:

Use the program `main2a.m` to illustrate a 2-dimensional normal probability density function given by a mean,  $\boldsymbol{\mu}$ , and covariance matrix,  $\boldsymbol{\Sigma}$ .

Discuss the quality of the histogram as you vary the number of samples,  $N$ , from small to large values. Compare your findings with the results from exercise 1 for the 1D normal distribution and relate this to the curse of dimensionality.

## Interpretation of Covariance

A one-dimensional normal distribution is given by its mean,  $\mu$ , and its variance,  $\sigma^2$ . The variance describes the variation of the variable around its mean.

In two dimensions, each sample consists of two components. Each dimension has a mean and a variance just as in the one-dimensional case. Consider a sample,  $\mathbf{x} = (x_1, x_2)^T$  from a 2D normal distribution with mean  $\boldsymbol{\mu} = (\mu_1, \mu_2)$ . If the variance of  $x_1$ , namely  $\sigma_1^2$  is large, an individual sample of  $x_1$  may well be quite different from  $\mu_1$ , and similarly for  $x_2$ . However, there may be a trend that whenever  $x_1$  is larger than  $\mu_1$ ,  $x_2$  is also larger than  $\mu_2$ , and that whenever  $x_1$  is smaller than  $\mu_1$ ,  $x_2$  is also smaller than  $\mu_2$ . In such a case,  $x_1$  and  $x_2$  are not independent, and they are said to be correlated.

Another term is therefore needed to fully describe the variance of the variable,  $\mathbf{x}$ , namely the covariance between its components,  $\text{cov}[x_1, x_2] \equiv \mathbb{E}[(x_1 - \mu_1)(x_2 - \mu_2)]$ . The covariance matrix of  $\mathbf{x}$  is then given by

$$\boldsymbol{\Sigma} = \begin{pmatrix} \sigma_1^2 & \text{cov}[x_1, x_2] \\ \text{cov}[x_2, x_1] & \sigma_2^2 \end{pmatrix} \equiv \begin{pmatrix} \sigma_{11} & \sigma_{12} \\ \sigma_{21} & \sigma_{22} \end{pmatrix}. \quad (9)$$

The terms  $\sigma_{12}$  and  $\sigma_{21}$  are equal since they describe the covariance between the same components. The covariance matrix is therefore always symmetric. The magnitude of the covariance term for a given correlation between the two components also depends on the diagonal variance terms. A useful quantity describing the correlation between the components is the correlation coefficient,  $\rho$ . It is the normalized covariance and is given by

$$\rho = \frac{\text{cov}[x_1, x_2]}{\sqrt{\sigma_1^2 \sigma_2^2}} = \frac{\sigma_{12}}{\sqrt{\sigma_{11} \sigma_{22}}}. \quad (10)$$

where  $\rho \in [-1, 1]$ . However, the limiting case  $\rho = \pm 1$  corresponds to a perfect linear relationship between  $x_1$  and  $x_2$ . In this case the variable,  $\mathbf{x}$ , is not really 2-dimensional since one component completely defines the other.

### Checkpoint 2.2:

Use the program `main2b.m` to visualize the probability density functions of 2D normal distributions with different covariance matrices. For example, try to fix the variances,  $\sigma_1^2$  and  $\sigma_2^2$ , while only changing the covariance. Think of an example where there is no correlation between the components and implement this distribution. Comment on the dependence of the orientation and shape of the ellipsoids in the contour plots of quadratic form induced by the covariance matrix.

## Coordinate Transformation

For some non-linear signal detection algorithms it is desired that the input should have zero mean, unit variance and zero covariance. The advantage of this is that it is possible to use the same algorithm (and not changing the control parameters of it) for variables of very different origins and covariation.

Geometrically, such a normalization corresponds to a coordinate transformation to the system defined by the eigenvectors of the covariance matrix. Typically, the mean and covariance matrix are not known, and must therefore be estimated from the data-set,  $\mathcal{D} = \{\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N\}$ :

$$\hat{\mathbf{x}} = \frac{1}{N} \sum_{i=1}^N \mathbf{x}_i \quad (11)$$

$$\hat{\mathbf{\Sigma}} = \frac{1}{N-1} \sum_{i=1}^N (\mathbf{x}_i - \hat{\mathbf{x}})(\mathbf{x}_i - \hat{\mathbf{x}})^T. \quad (12)$$

The eigenvalue equation for the covariance matrix is

$$\hat{\mathbf{\Sigma}} \mathbf{u}_j = \lambda_j \mathbf{u}_j, \quad j = 1, \dots, d, \quad (13)$$

where  $\lambda_j$  is the  $j$ 'th eigenvalue and  $\mathbf{u}_j$  is the corresponding eigenvector of  $\hat{\mathbf{\Sigma}}$ . The transformed input variables are then given by

$$\tilde{\mathbf{x}}_i = \mathbf{\Lambda}^{-1/2} \mathbf{U}^T (\mathbf{x}_i - \hat{\mathbf{x}}), \quad (14)$$

where

$$\mathbf{U} = (\mathbf{u}_1, \dots, \mathbf{u}_d) \quad (15)$$

$$\mathbf{\Lambda} = \text{diag}(\lambda_1, \dots, \lambda_d). \quad (16)$$

It can be shown that the transformed data-set,  $\tilde{\mathcal{D}} = \{\tilde{\mathbf{x}}_1, \tilde{\mathbf{x}}_2, \dots, \tilde{\mathbf{x}}_N\}$  has zero mean and a covariance matrix given by the unit matrix.

### Checkpoint 2.3:

Use the program `main2c.m` to calculate the eigenvalues and eigenvectors of the covariance matrix for different distributions. Comment on the geometrical significance of the eigenvalues and eigenvectors. Compare the transformed data-sets from different distributions. What happens if the term  $\mathbf{\Lambda}^{-1/2}$  is removed from equation (14)?

## Projection on Eigenvectors

In some cases, the measured data is of a lower “true” dimension than the apparent dimension of the data vector. For example, imagine a data-set of a 3-dimensional variable. If all the data are on a straight line, the true dimension of the data is only 1D. If the data-set is transformed to a coordinate system, where the variation of the data is along one of the axes, the two other components can be ignored.

Let  $\lambda_1, \dots, \lambda_d$  be the ordered set of eigenvalues of the covariance matrix, such that  $\lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_d$ . If there exists a number  $m$ , such that  $\lambda_i \gg \lambda_j$ ,  $i = 1, \dots, m$ , and  $j = m+1, \dots, d$ , then the data-set can be transformed to a coordinate system, where most of the signal variance is in an  $m$ -dimensional linear subspace spanned by the  $m$ 'th first eigenvectors in the ordered list. This transformation is again given by the eigenvectors of the covariance matrix ( $\mathbf{U}$ ),

$$\tilde{\mathbf{x}}_i = \mathbf{U}^T(\mathbf{x}_i - \hat{\mathbf{x}}). \quad (17)$$

If we extract only the first  $m$  components of the transformed datavector  $\tilde{\mathbf{x}}$  we obtain a signal that carries most of the variation of the original signal. Such reduction of the effective dimensionality of the problem is also known as extraction of features.

### Checkpoint 2.4:

Use the programs `main2d.m` and `main2e.m` to transform 2D datasets into the eigenvector-space and comment on the “true” dimensionality of the classification problems.

### Challenges (not part of the curriculum)

- 1) Prove that the transformed data in Eq. (14) is zero mean and has a unit covariance matrix.
- 2) How would you make the decision on the number of principal components to retain? Hint: Check K.W. Jorgensen, L.K. Hansen: *Model selection for Gaussian kernel PCA denoising*. IEEE Transactions on Neural Networks and Learning Systems **23**(1):163-168 (2012), the references in this paper.

DTU, September 2009,

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