

Figure 1: Examples of univariate Gaussian PDFs  $\mathcal{N}(x; \mu, \sigma^2)$ .

## The Gaussian distribution

Probably the most-important distribution in all of statistics is the *Gaussian distribution*, also called the *normal distribution*. The Gaussian distribution arises in many contexts and is widely used for modeling continuous random variables.

The probability density function of the univariate (one-dimensional) Gaussian distribution is

$$p(x \mid \mu, \sigma^2) = \mathcal{N}(x; \mu, \sigma^2) = \frac{1}{Z} \exp\left(-\frac{(x - \mu)^2}{2\sigma^2}\right).$$

The normalization constant Z is

$$Z = \sqrt{2\pi\sigma^2}.$$

The parameters  $\mu$  and  $\sigma^2$  specify the mean and variance of the distribution, respectively:

$$\mu = \mathbb{E}[x]; \qquad \sigma^2 = \text{var}[x].$$

Figure 1 plots the probability density function for several sets of parameters  $(\mu, \sigma^2)$ . The distribution is symmetric around the mean and most of the density ( $\approx 99.7\%$ ) is contained within  $\pm 3\sigma$  of the mean.

We may extend the univariate Gaussian distribution to a distribution over d-dimensional vectors, producing a multivariate analog. The probablity density function of the multivariate Gaussian distribution is

$$p(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \mathcal{N}(\mathbf{x}; \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{1}{Z} \exp \left( -\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^{\top} \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu}) \right).$$

The normalization constant Z is

$$Z = \sqrt{\det(2\pi \Sigma)} = (2\pi)^{d/2} (\det \Sigma)^{1/2}.$$

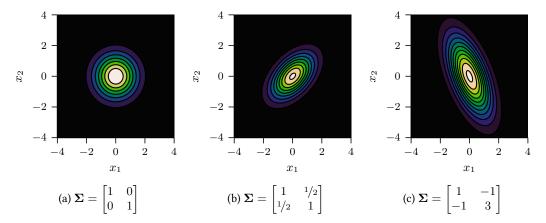


Figure 2: Contour plots for example bivariate Gaussian distributions. Here  $\mu = 0$  for all examples.

Examining these equations, we can see that the multivariate density coincides with the univariate density in the special case when  $\Sigma$  is the scalar  $\sigma^2$ .

Again, the vector  $\mu$  specifies the mean of the multivariate Gaussian distribution. The matrix  $\Sigma$  specifies the *covariance* between each pair of variables in  $\mathbf{x}$ :

$$\Sigma = \text{cov}(\mathbf{x}, \mathbf{x}) = \mathbb{E}[(\mathbf{x} - \boldsymbol{\mu})(\mathbf{x} - \boldsymbol{\mu})^{\top}].$$

Covariance matrices are necessarily symmetric and positive semidefinite, which means their eigenvalues are nonnegative. Note that the density function above requires that  $\Sigma$  be positive definite, or have strictly positive eigenvalues. A zero eigenvalue would result in a determinant of zero, making the normalization impossible.

The dependence of the multivariate Gaussian density on x is entirely through the value of the quadratic form

$$\Delta^2 = (\mathbf{x} - \boldsymbol{\mu})^{\top} \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu}).$$

The value  $\Delta$  (obtained via a square root) is called the *Mahalanobis distance*, and can be seen as a generalization of the Z score  $\frac{(x-\mu)}{\sigma}$ , often encountered in statistics.

To understand the behavior of the density geometrically, we can set the Mahalanobis distance to a constant. The set of points in  $\mathbb{R}^d$  satisfying  $\Delta=c$  for any given value c>0 is an ellipsoid with the eigenvectors of  $\Sigma$  defining the directions of the principal axes.

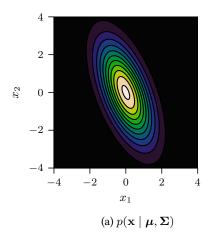
Figure 2 shows contour plots of the density of three bivariate (two-dimensional) Gaussian distributions. The elliptical shape of the contours is clear.

The Gaussian distribution has a number of convenient analytic properties, some of which we describe below.

# Marginalization

Often we will have a set of variables  $\mathbf{x}$  with a joint multivariate Gaussian distribution, but only be interested in reasoning about a subset of these variables. Suppose  $\mathbf{x}$  has a multivariate Gaussian distribution:

$$p(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \mathcal{N}(\mathbf{x}, \boldsymbol{\mu}, \boldsymbol{\Sigma}).$$



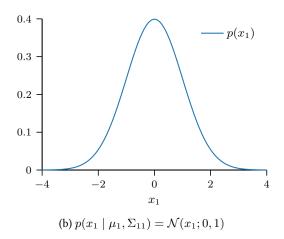


Figure 3: Marginalization example. (a) shows the joint density over  $\mathbf{x} = [x_1, x_2]^{\top}$ ; this is the same density as in Figure 2(c). (b) shows the marginal density of  $x_1$ .

Let us partition the vector into two components:

$$\mathbf{x} = \begin{bmatrix} \mathbf{x}_1 \\ \mathbf{x}_2 \end{bmatrix}$$
.

We partition the mean vector and covariance matrix in the same way:

$$oldsymbol{\mu} = egin{bmatrix} oldsymbol{\mu}_1 \ oldsymbol{\mu}_2 \end{bmatrix} \qquad oldsymbol{\Sigma} = egin{bmatrix} oldsymbol{\Sigma}_{11} & oldsymbol{\Sigma}_{12} \ oldsymbol{\Sigma}_{21} & oldsymbol{\Sigma}_{22} \end{bmatrix}.$$

Now the marginal distribution of the subvector  $\mathbf{x}_1$  has a simple form:

$$p(\mathbf{x}_1 \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \mathcal{N}(\mathbf{x}_1, \boldsymbol{\mu}_1, \boldsymbol{\Sigma}_{11}),$$

so we simply pick out the entries of  $\mu$  and  $\Sigma$  corresponding to  $x_1$ .

Figure 3 illustrates the marginal distribution of  $x_1$  for the joint distribution shown in Figure 2(c).

## Conditioning

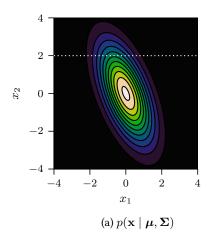
Another common scenario will be when we have a set of variables  $\mathbf{x}$  with a joint multivariate Gaussian prior distribution, and are then told the value of a subset of these variables. We may then condition our prior distribution on this observation, giving a posterior distribution over the remaining variables.

Suppose again that x has a multivariate Gaussian distribution:

$$p(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \mathcal{N}(\mathbf{x}, \boldsymbol{\mu}, \boldsymbol{\Sigma}),$$

and that we have partitioned as before:  $\mathbf{x} = [\mathbf{x}_1, \mathbf{x}_2]^{\top}$ . Suppose now that we learn the exact value of the subvector  $\mathbf{x}_2$ . Remarkably, the posterior distribution

$$p(\mathbf{x}_1 \mid \mathbf{x}_2, \boldsymbol{\mu}, \boldsymbol{\Sigma})$$



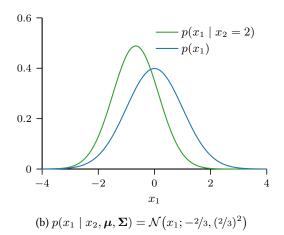


Figure 4: Conditioning example. (a) shows the joint density over  $\mathbf{x} = [x_1, x_2]^{\top}$ , along with the observation value  $x_2 = 2$ ; this is the same density as in Figure 2(c). (b) shows the conditional density of  $x_1$  given  $x_2 = 2$ .

is a Gaussian distribution! The formula is

$$p(\mathbf{x}_1 \mid \mathbf{x}_2, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \mathcal{N}(\mathbf{x}_1; \boldsymbol{\mu}_{1|2}, \boldsymbol{\Sigma}_{11|2}),$$

with

$$\begin{split} \boldsymbol{\mu}_{1|2} &= \boldsymbol{\mu}_1 + \boldsymbol{\Sigma}_{12}\boldsymbol{\Sigma}_{22}^{-1}(\mathbf{x}_2 - \boldsymbol{\mu}_2); \\ \boldsymbol{\Sigma}_{11|2} &= \boldsymbol{\Sigma}_{11} - \boldsymbol{\Sigma}_{12}\boldsymbol{\Sigma}_{22}^{-1}\boldsymbol{\Sigma}_{21}. \end{split}$$

So we adjust the mean by an amount dependent on: (1) the covariance between  $\mathbf{x}_1$  and  $\mathbf{x}_2$ ,  $\Sigma_{12}$ , (2) the prior uncertainty in  $\mathbf{x}_2$ ,  $\Sigma_{22}$ , and (3) the deviation of the observation from the prior mean,  $(\mathbf{x}_2 - \boldsymbol{\mu}_2)$ . Similarly, we reduce the uncertainty in  $\mathbf{x}_1$ ,  $\Sigma_{11}$ , by an amount dependent on (1) and (2). Notably, the reduction of the covariance matrix does *not* depend on the values we observe.

Notice that if  $\mathbf{x}_1$  and  $\mathbf{x}_2$  are independent, then  $\Sigma_{12} = \mathbf{0}$ , and the conditioning operation does not change the distribution of  $\mathbf{x}_1$ , as expected.

Figure 4 illustrates the conditional distribution of  $x_1$  for the joint distribution shown in Figure 2(c), after observing  $x_2 = 2$ .

## Convolutions

Gaussian probability density functions are closed under convolutions. Let  $\mathbf{x}$  and  $\mathbf{y}$  be d-dimensional vectors, with distributions

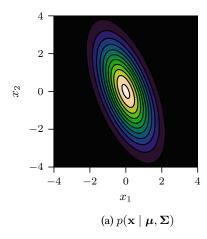
$$p(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \mathcal{N}(\mathbf{x}; \boldsymbol{\mu}, \boldsymbol{\Sigma}); \qquad p(\mathbf{y} \mid \boldsymbol{\nu}, \mathbf{P}) = \mathcal{N}(\mathbf{y}; \boldsymbol{\nu}, \mathbf{P}).$$

Then the convolution of their density functions is another Gaussian PDF:

$$f(\mathbf{y}) = \int \mathcal{N}(\mathbf{y} - \mathbf{x}; \boldsymbol{\nu}, \mathbf{P}) \, \mathcal{N}(\mathbf{x}; \boldsymbol{\mu}, \boldsymbol{\Sigma}) \, d\mathbf{x} = \mathcal{N}(\mathbf{y}; \boldsymbol{\mu} + \boldsymbol{\nu}, \boldsymbol{\Sigma} + \mathbf{P}),$$

where the mean and covariances add in the result. (This can be proven by considering the Fourier transform of a Gaussian, which is another Gaussian.)

This result will often come in handy.



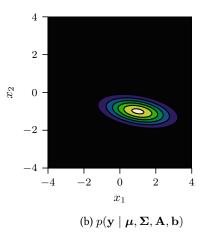


Figure 5: Affine transformation example. (a) shows the joint density over  $\mathbf{x} = [x_1, x_2]^{\top}$ ; this is the same density as in Figure 2(c). (b) shows the density of  $\mathbf{y} = \mathbf{A}\mathbf{x} + \mathbf{b}$ . The values of  $\mathbf{A}$  and  $\mathbf{b}$  are given in the text. The density of the transformed vector is another Gaussian.

#### Affine transformations

Consider a d-dimensional vector  $\mathbf{x}$  with a multivariate Gaussian distribution:

$$p(\mathbf{x} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \mathcal{N}(\mathbf{x}, \boldsymbol{\mu}, \boldsymbol{\Sigma}).$$

Suppose we wish to reason about an affine transformation of  $\mathbf{x}$  into  $\mathbb{R}^D$ ,  $\mathbf{y} = \mathbf{A}\mathbf{x} + \mathbf{b}$ , where  $\mathbf{A} \in \mathbb{R}^{D \times d}$  and  $\mathbf{b} \in \mathbb{R}^D$ . Then  $\mathbf{y}$  has a D-dimensional Gaussian distribution:

$$p(\mathbf{y} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}, \mathbf{A}, \mathbf{b}) = \mathcal{N}(\mathbf{y}, \mathbf{A}\boldsymbol{\mu} + \mathbf{b}, \mathbf{A}\boldsymbol{\Sigma}\mathbf{A}^{\top}).$$

Figure 5 illustrates an affine transformation of the vector  $\mathbf{x}$  with the joint distribution shown in Figure 2(c), for the values

$$\mathbf{A} = \begin{bmatrix} 1/5 & -3/5 \\ 1/2 & 3/10 \end{bmatrix}; \qquad \mathbf{b} = \begin{bmatrix} 1 \\ -1 \end{bmatrix}.$$

The density has been rotated and translated, but remains a Gaussian.

# Selecting parameters

The d-dimensional multivariate Gaussian distribution is specified by the parameters  $\mu$  and  $\Sigma$ . Without any further restrictions, specifying  $\mu$  requires d parameters and specifying  $\Sigma$  requires a further  $\binom{d}{2} = \frac{d(d-1)}{2}$ . The number of parameters therefore grows quadratically in the dimension, which can sometimes cause difficulty. For this reason, we sometimes restrict the covariance matrix  $\Sigma$  in some way to reduce the number of parameters.

Common choices are to set  $\Sigma = \operatorname{diag} \tau$ , where  $\tau$  is a vector of marginal variances, and  $\Sigma = \sigma^2 \mathbf{I}$ , a constant diagonal matrix. Both of these options assume independence between the variables in  $\mathbf{x}$ . The former case is more flexible, allowing a different scale parameter for each entry, whereas the latter assumes an equal marginal variance of  $\sigma^2$  for each variable. Geometrically, the densities are axis-aligned, as in Figure 2(a), and in the latter case, the isoprobability contours are spherical (also as in Figure 2(a)).