# Non-parametric Methods

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### Introduction

- ▶ Density estimation with parametric models assumes that the forms of the underlying density functions are known.
- However, common parametric forms do not always fit the densities actually encountered in practice.
- In addition, most of the classical parametric densities are unimodal, whereas many practical problems involve multimodal densities.
- Non-parametric methods can be used with arbitrary distributions and without the assumption that the forms of the underlying densities are known.

- ▶ Suppose that n samples  $\mathbf{x_1}, \dots, \mathbf{x_n}$  are drawn i.i.d. according to the distribution  $p(\mathbf{x})$ .
- ▶ The probability P that a vector  $\mathbf{x}$  will fall in a region  $\mathcal{R}$  is given by

$$P = \int_{\mathcal{R}} p(\mathbf{x'}) d\mathbf{x'}.$$

▶ The probability that k of the n will fall in  $\mathcal{R}$  is given by the binomial law

$$P_k = \binom{n}{k} P^k (1 - P)^{n-k}.$$

▶ The expected value of k is E[k] = nP and the MLE for P is

$$\hat{P} = \frac{k}{n}$$
.

▶ If we assume that  $p(\mathbf{x})$  is continuous and  $\mathcal{R}$  is small enough so that  $p(\mathbf{x})$  does not vary significantly in it, we can get the approximation

$$\int_{\mathcal{R}} p(\mathbf{x'}) d\mathbf{x'} \simeq p(\mathbf{x}) V$$

where x is a point in  $\mathcal{R}$  and V is the volume of  $\mathcal{R}$ .

► Then, the density estimate becomes

$$p(\mathbf{x}) \simeq \frac{k/n}{V}.$$



- Let n be the number of samples used,  $\mathcal{R}_n$  be the region used with n samples,  $V_n$  be the volume of  $\mathcal{R}_n$ ,  $k_n$  be the number of samples falling in  $\mathcal{R}_n$ , and  $p_n(\mathbf{x}) = \frac{k_n/n}{V_n}$  be the estimate for  $p(\mathbf{x})$ .
- ▶ If  $p_n(\mathbf{x})$  is to converge to  $p(\mathbf{x})$ , three conditions are required:

$$\lim_{n \to \infty} V_n = 0$$

$$\lim_{n \to \infty} k_n = \infty$$

$$\lim_{n \to \infty} \frac{k_n}{n} = 0.$$



## Histogram Method

► A very simple method is to partition the space into a number of equally-sized cells (bins) and compute a histogram.

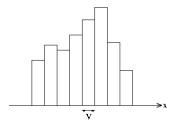


Figure 1: Histogram in one dimension.

▶ The estimate of the density at a point x becomes

$$p(\mathbf{x}) = \frac{k}{nV}$$

where n is the total number of samples, k is the number of samples in the cell that includes  $\mathbf{x}$ , and V is the volume of that cell.

# Histogram Method

- ► Although the histogram method is very easy to implement, it is usually not practical in high-dimensional spaces due to the number of cells.
- Many observations are required to prevent the estimate being zero over a large region.
- Modifications for overcoming these difficulties:
  - Data-adaptive histograms,
  - Independence assumption (naive Bayes),
  - Dependence trees.



- ▶ Other methods for obtaining the regions for estimation:
  - ► Shrink regions as some function of n, such as  $V_n = 1/\sqrt{n}$ . This is the *Parzen window* estimation.
  - ▶ Specify  $k_n$  as some function of n, such as  $k_n = \sqrt{n}$ . This is the k-nearest neighbor estimation.

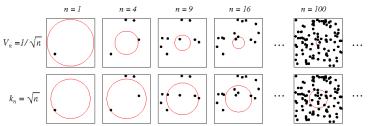


Figure 2: Methods for estimating the density at a point, here at the center of each square.

▶ Suppose that  $\varphi$  is a d-dimensional window function that satisfies the properties of a density function, i.e.,

$$\varphi(\mathbf{u}) \ge 0$$
 and  $\int \varphi(\mathbf{u}) d\mathbf{u} = 1$ .

► A density estimate can be obtained as

$$p_n(\mathbf{x}) = \frac{1}{n} \sum_{i=1}^n \frac{1}{V_n} \varphi\left(\frac{\mathbf{x} - \mathbf{x}_i}{h_n}\right)$$

where  $h_n$  is the window width and  $V_n = h_n^d$ .



The density estimate can also be written as

$$p_n(\mathbf{x}) = \frac{1}{n} \sum_{i=1}^n \delta_n(\mathbf{x} - \mathbf{x_i}) \quad \text{where} \quad \delta_n(\mathbf{x}) = \frac{1}{V_n} \, \varphi\left(\frac{\mathbf{x}}{h_n}\right).$$

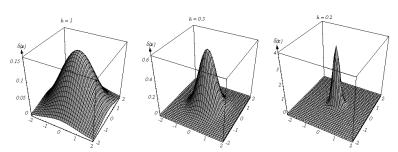


Figure 3: Examples of two-dimensional circularly symmetric Parzen windows functions for three different values of  $h_n$ . The value of  $h_n$  affects both the amplitude and the width of  $\delta_n(\mathbf{x})$ .

- ▶ If  $h_n$  is very large,  $p_n(\mathbf{x})$  is the superposition of n broad functions, and is a smooth "out-of-focus" estimate of  $p(\mathbf{x})$ .
- ▶ If  $h_n$  is very small,  $p_n(\mathbf{x})$  is the superposition of n sharp pulses centered at the samples, and is a "noisy" estimate of  $p(\mathbf{x})$ .
- As  $h_n$  approaches zero,  $\delta_n(\mathbf{x} \mathbf{x}_i)$  approaches a Dirac delta function centered at  $\mathbf{x}_i$ , and  $p_n(\mathbf{x})$  is a superposition of delta functions.

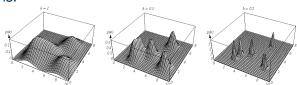


Figure 4: Parzen window density estimates based on the same set of five samples using the window functions in the previous figure.

11 / 25

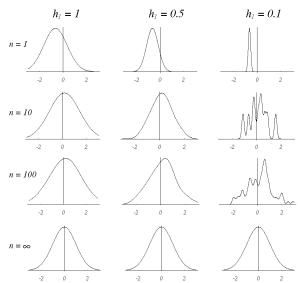


Figure 5: Parzen window estimates of a univariate Gaussian density using different window widths and numbers of samples where  $\varphi(u)=N(0,1)$  and  $h_n=h_1/\sqrt{n}$ .

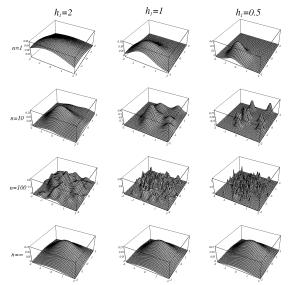


Figure 6: Parzen window estimates of a bivariate Gaussian density using different window widths and numbers of samples where  $\varphi(\mathbf{u}) = N(\mathbf{0}, \mathbf{I})$  and  $h_n = h_1/\sqrt{n}$ .

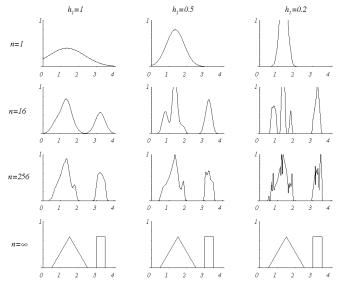
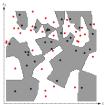


Figure 7: Estimates of a mixture of a uniform and a triangle density using different window widths and numbers of samples where  $\varphi(u)=N(0,1)$  and  $h_n=h_1/\sqrt{n}$ .

- Densities estimated using Parzen windows can be used with the Bayesian decision rule for classification.
- ► The training error can be made arbitrarily low by making the window width sufficiently small.
- ► However, the goal is to classify novel patterns so the window width cannot be made too small.



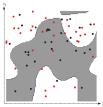


Figure 8: Decision boundaries in 2-D. The left figure uses a small window width and the right figure uses a larger window width.

## k-Nearest Neighbors

- A potential remedy for the problem of the unknown "best" window function is to let the estimation volume be a function of the training data, rather than some arbitrary function of the overall number of samples.
- ightharpoonup To estimate  $p(\mathbf{x})$  from n samples, we can center a volume about x and let it grow until it captures  $k_n$  samples, where  $k_n$  is some function of n.
- ▶ These samples are called the *k*-nearest neighbors of x.
- ▶ If the density is high near x, the volume will be relatively small. If the density is low, the volume will grow large.

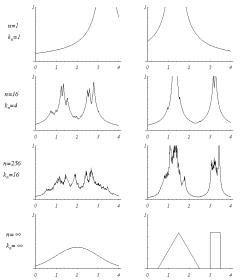


Figure 9: *k*-nearest neighbor estimates of two 1-D densities: a Gaussian and a bimodal distribution.

### k-Nearest Neighbors

- ▶ Posterior probabilities can be estimated from a set of n labeled samples and can be used with the Bayesian decision rule for classification.
- ▶ Suppose that a volume V around  $\mathbf{x}$  includes k samples,  $k_i$  of which are labeled as belonging to class  $w_i$ .
- lacktriangle As estimate for the joint probability  $p(\mathbf{x}, w_i)$  becomes

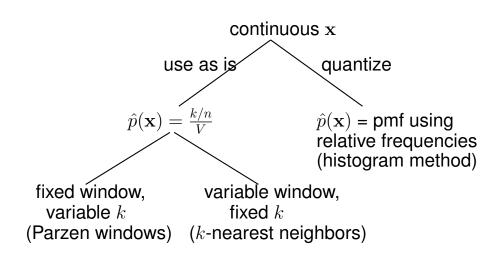
$$p_n(\mathbf{x}, w_i) = \frac{k_i/n}{V}$$

and gives an estimate for the posterior probability

$$P_n(w_i|\mathbf{x}) = \frac{p_n(\mathbf{x}, w_i)}{\sum_{j=1}^c p_n(\mathbf{x}, w_j)} = \frac{k_i}{k}.$$



# Non-parametric Methods





### Non-parametric Methods

#### Advantages:

- No assumptions are needed about the distributions ahead of time (generality).
- With enough samples, convergence to an arbitrarily complicated target density can be obtained.

#### Disadvantages:

- The number of samples needed may be very large (number grows exponentially with the dimensionality of the feature space).
- There may be severe requirements for computation time and storage.

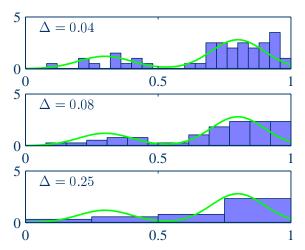


Figure 10: An illustration of the histogram approach to density estimation, in which a data set of 50 points is generated from the distribution shown by the green curve. Histogram density estimates are shown for various values of the cell volume  $(\Delta)$ .

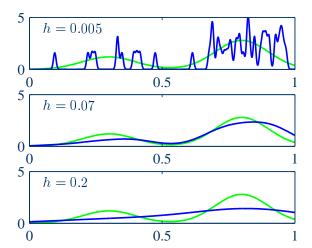


Figure 11: Illustration of the Parzen density model. The window width (h) acts as a smoothing parameter. If it is set too small (top), the result is a very noisy density model. If it is set too large (bottom), the bimodal nature of the underlying distribution is washed out. An intermediate value (middle) gives a good estimate.

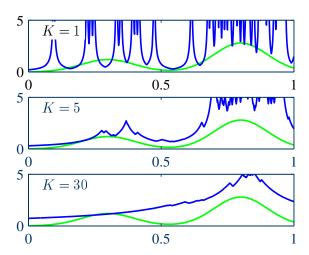


Figure 12: Illustration of the k-nearest neighbor density model. The parameter k governs the degree of smoothing. A small value of k (top) leads to a very noisy density model. A large value (bottom) smoothes out the bimodal nature of the true distribution.

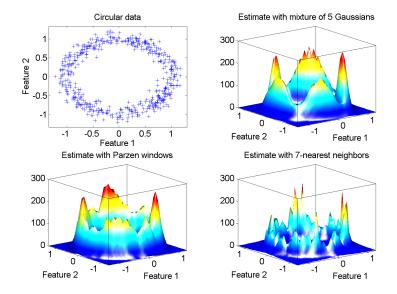


Figure 13: Density estimation examples for 2-D circular data.



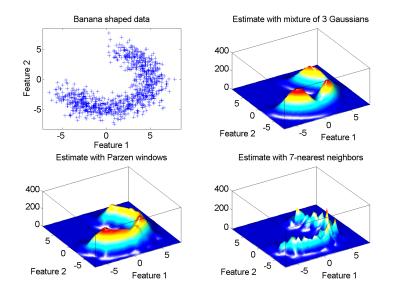


Figure 14: Density estimation examples for 2-D banana shaped data.