## Non-parametric Methods

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CS 551, Fall 2015

- 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 x<sub>1</sub>,..., x<sub>n</sub> are drawn i.i.d. according to the distribution p(x).
- The probability P that a vector x will fall in a region R is given by

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

► The probability that k of the n will fall in 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  $\hat{P} = \frac{k}{2}$ 



If we assume that p(x) is continuous and R is small enough so that p(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  $\mathbf{x}$  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, R<sub>n</sub> be the region used with n samples, V<sub>n</sub> be the volume of R<sub>n</sub>, k<sub>n</sub> be the number of samples falling in R<sub>n</sub>, and p<sub>n</sub>(x) = <sup>k<sub>n</sub>/n</sup>/<sub>V<sub>n</sub></sub> be the estimate for p(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.

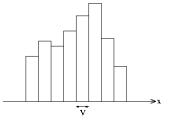


Figure 1: Histogram in one dimension.

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. 6/25 CS 551, Fall 2015

- 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.

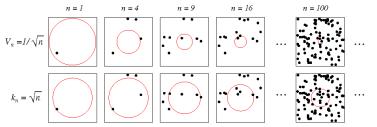


Figure 2: Methods for estimating the density at a point, here at the center of each square.



 Suppose that φ 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

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

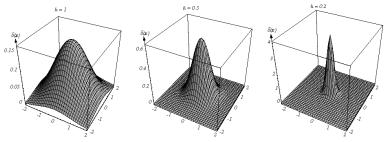


Figure 3: Examples of two-dimensional circularly symmetric Parzen windows functions for three different values of  $h_n$ . The value of  $h_n$  affects both the amplitude and the width of  $\delta_n(\mathbf{x})$ .

- ► If h<sub>n</sub> is very large, p<sub>n</sub>(x) is the superposition of n broad functions, and is a smooth "out-of-focus" estimate of p(x).
- ► If *h<sub>n</sub>* is very small, *p<sub>n</sub>*(**x**) is the superposition of *n* sharp pulses centered at the samples, and is a "noisy" estimate of *p*(**x**).
- As h<sub>n</sub> approaches zero, δ<sub>n</sub>(x − x<sub>i</sub>) approaches a Dirac delta function centered at x<sub>i</sub>, and p<sub>n</sub>(x) is a superposition of delta functions.

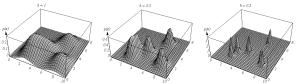


Figure 4: Parzen window density estimates based on the same set of five samples using the window functions in the previous figure.

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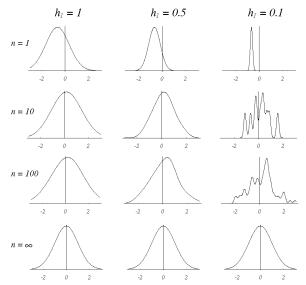


Figure 5: Parzen window estimates of a univariate Gaussian density using different window widths and numbers of samples where  $\varphi(u) = N(0, 1)$  and  $h_n = h_1/\sqrt{n}$ .

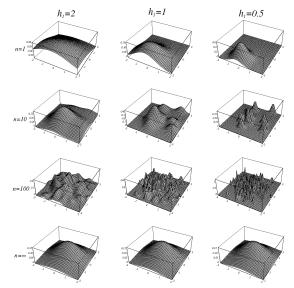


Figure 6: Parzen window estimates of a bivariate Gaussian density using different window widths and numbers of samples where  $\varphi(\mathbf{u}) = N(\mathbf{0}, \mathbf{I})$  and  $h_n = h_1/\sqrt{n}$ .

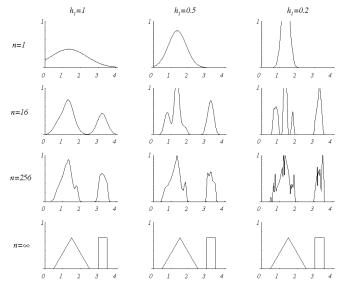


Figure 7: Estimates of a mixture of a uniform and a triangle density using different window widths and numbers of samples where  $\varphi(u) = N(0, 1)$  and  $h_n = h_1/\sqrt{n}$ .

- 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.

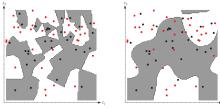


Figure 8: Decision boundaries in 2-D. The left figure uses a small window width and the right figure uses a larger window width.

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- 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.
- To estimate p(x) from n samples, we can center a volume about x and let it grow until it captures k<sub>n</sub> samples, where k<sub>n</sub> 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.



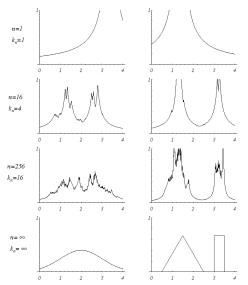


Figure 9: *k*-nearest neighbor estimates of two 1-D densities: a Gaussian and a bimodal distribution.



## 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 x includes k samples, k<sub>i</sub> of which are labeled as belonging to class w<sub>i</sub>.
- As estimate for the joint probability  $p(\mathbf{x}, w_i)$  becomes

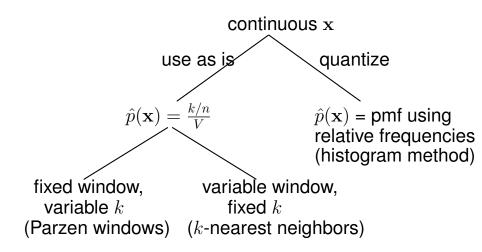
$$p_n(\mathbf{x}, w_i) = \frac{k_i/n}{V}$$

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.



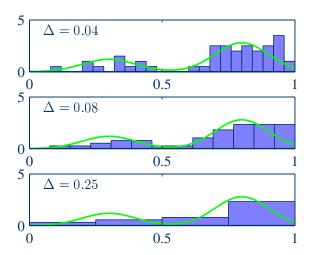


Figure 10: An illustration of the histogram approach to density estimation, in which a data set of 50 points is generated from the distribution shown by the green curve. Histogram density estimates are shown for various values of the cell volume ( $\Delta$ ).



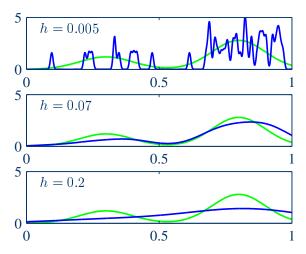


Figure 11: Illustration of the Parzen density model. The window width (h) acts as a smoothing parameter. If it is set too small (top), the result is a very noisy density model. If it is set too large (bottom), the bimodal nature of the underlying distribution is washed out. An intermediate value (middle)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



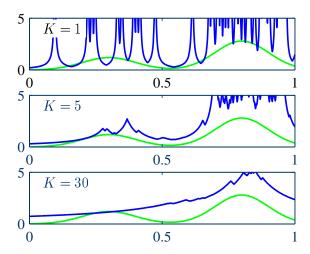


Figure 12: Illustration of the k-nearest neighbor density model. The parameter k governs the degree of smoothing. A small value of k (top) leads to a very noisy density model. A large value (bottom) smoothes out the bimodal nature of the true distribution.



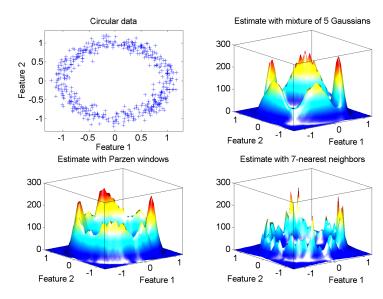
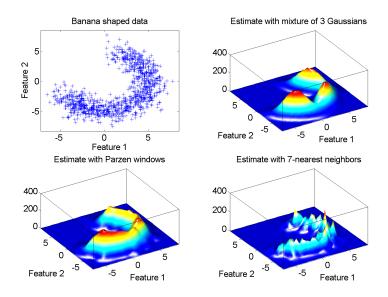


Figure 13: Density estimation examples for 2-D circular data.





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



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