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May 2, 2023

# **Course Information**

- Course web page: https://research.nii.ac.jp/~satoh/utpr/
- Course materials can be found in the above page
- Course videos can be found in ITC-LMS
- Credits will be given based on final report (mandatory)
- Attendance record will not be taken
- Assignments may be imposed (3 out of 7 are mandatory: subject to change)
- The first assignment issued on April 18 will be due on today, the second issued on April 25 will be due on next week, and the third issued today will be due on the next to next week (May 16)
- If you fail to submit minimum 3 assignments and final report, you will not obtain credits.

### Recap

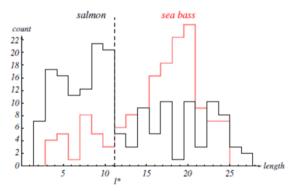
- We visited "parametric" methods so far.
- Probability distribution functions (or equivalently decision boundaries) can be represented by parametric forms.
- e.g., Normal density case: mean and variance (or covariance matrix)
- These methods assume that the underlying probability distribution of the actual observations is known and yields parametric forms.
- However, in many cases this assumption is suspect.

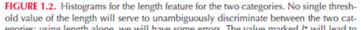
# Today's topics

- Nonparametric Density Estimation Methods
  - Parzen Window
  - k-Nearest Neighbor Estimation

### Nonparametric Methods

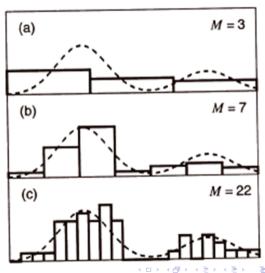
- Simple approach is to compose histogram
- Knowing sample data, we can compose histogram with certain bin size (division of each axis)
- Treat the histogram as probability distribution function





# Nonparametric Methods

- The optimal number of bins M (or bin size) is the issue.
  - If bin width is small (i.e., big M), then the estimated density is very spiky (i.e., noisy).
  - If bin width is large (i.e., small M), then the true structure of the density is smoothed out.
- In practice, we need to find an optimal value for M that compromises between these two issues.
- Also, how we extend to the multidimensional case?



 The probability that a given vector x, drawn from the unknown density p(x), will fall inside some region R in the input space is given by:

$$P = \int_R p(\mathbf{x}') d\mathbf{x}'$$

• If we have *n* data points {*x*<sub>1</sub>, *x*<sub>2</sub>,..., *x<sub>n</sub>*} drawn independently from *p*(**x**), the probability that *k* of them will fall in *R* is given by the binomial law:

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

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• The expected value of k is:

$$E\{k\} = nP$$

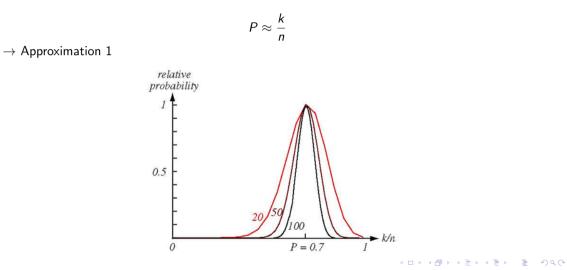
• The expected percentage of points falling in R is:

$$E\{\frac{k}{n}\}=P$$

• The variance is given by:

$$Var\{\frac{k}{n}\} = E\{(\frac{k}{n} - P)^2\} = \frac{P(1 - P)}{n}$$

The distribution is sharply peaked as  $n \rightarrow \inf$ , thus:

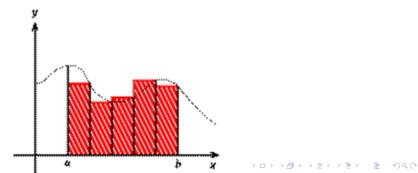


If we assume that  $p(\mathbf{x})$  is continuous and does not vary significantly over the region R, we can approximate P by:

$$P = \int_{R} p(\mathbf{x}') d\mathbf{x}' pprox p(\mathbf{x}) V$$

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where V is the volume enclosed by R.



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Combining these two approximations we have:

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

- The above approximation is based on contradictory assumptions:
  - *R* is relatively large (i.e., it contains many samples so that  $P_k$  is sharply peaked): Approximation 1
  - R is relatively small so that p(x) is approximately constant inside the integration region: Approximation 2
- We need to choose an optimum R in practice ...

- Suppose we form regions  $R_1, R_2, \ldots$  containing **x**.
  - $R_1$  contains  $k_1$  sample,  $R_2$  contains  $k_2$  samples, etc.
  - We assume that each case corresponds to *n* samples.
- $R_i$  has volume  $V_i$  and contains  $k_i$  samples.
- The *n*-th estimate  $p_n(\mathbf{x})$  of  $p(\mathbf{x})$  is given by:

$$p_n(\mathbf{x}) pprox rac{k_n/n}{V_n}$$

The following conditions must be satisfied in order for  $p_n(\mathbf{x})$  to converge to  $p(\mathbf{x})$ :

$$\lim_{n \to \infty} V_n = 0 \quad \text{Approximation 1}$$
$$\lim_{n \to \infty} k_n = \infty \quad \text{Approximation 2}$$
$$\lim_{n \to \infty} \frac{k_n}{n} = 0 \quad \text{to allow } p_n(\mathbf{x}) \text{ to converge}$$

$$p_n(\mathbf{x}) pprox rac{k_n/n}{V_n}$$

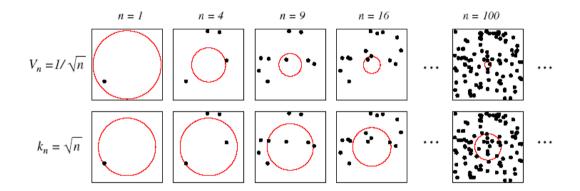
How to choose the optimum values for  $V_n$  and  $k_n$ ? Two leading approaches:

(1) Fix the volume  $V_n$  and determine  $k_n$  from the data (kernel-based density estimation methods), e.g.,

$$V_n = \frac{1}{\sqrt{n}}$$

(2) Fix the value of  $k_n$  and determine the corresponding volume  $V_n$  from the data (k-nearest neighbor method), e.g.,

$$k_n = \sqrt{n}$$



$$p_n(\mathbf{x}) pprox rac{k_n/n}{V_n}$$

- **Problem**: Given a vector  $\mathbf{x}$ , estimate  $p(\mathbf{x})$
- Assume  $R_n$  to be a hypercube with sides of length  $h_n$ , centered on the point **x**:

$$V_n = h_n^{\alpha}$$

• To find an expression for  $k_n$  (i.e., # points in the hypercube) let us define a kernel function:

$$arphi(u) = \left\{ egin{array}{cc} 1 & |u_j| \leq rac{1}{2} & (j=1,\ldots,d) \ 0 & ext{otherwise} \end{array} 
ight.$$

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• The total number of points x<sub>i</sub> falling inside the hypercube is:

$$k_n = \sum_{i=1}^n \varphi(\frac{\mathbf{x} - x_i}{h_n})$$

• Then, the estimate

$$p_n(\mathbf{x}) pprox rac{k_n/n}{V_n}$$

becomes

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

 $\rightarrow$  Parzen windows estimate

• The density estimate is a superposition of kernel functions and the samples  $x_i$ .

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

- $\varphi(u)$  interpolates the density between samples.
- Each sample x<sub>i</sub> contributes to the estimate based on its distance from **x**.

- The kernel function  $\varphi(u)$  can have a more general form (i.e., not just hypercube).
- In order for  $p_n(\mathbf{x})$  to be a legitimate estimate,  $\varphi(u)$  must be a valid density itself:

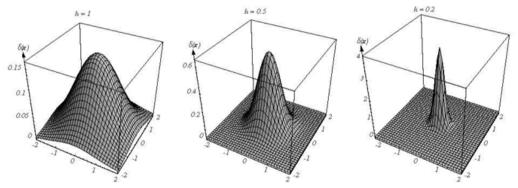
 $arphi(u) \geq 0$   $\int arphi(u) du = 1$ 



The parameter  $h_n$  acts as a smoothing parameter that needs to be optimized.

- When  $h_n$  is too large, the estimated density is over-smoothed (i.e., superposition of "broad" kernel functions).
- When  $h_n$  is too small, the estimate represents the properties of the data rather than the true density (i.e., superposition of "narrow" kernel functions)

 $\varphi(u)$  assuming different  $h_n$  values:



**FIGURE 4.3.** Examples of two-dimensional circularly symmetric normal Parzen windows for three different values of *h*. Note that because the  $\delta(\mathbf{x})$  are normalized, different vertical scales must be used to show their structure. From: Richard O. Duda, Peter E. Hart, and David G. Stork, *Pattern Classification*. Copyright © 2001 by John Wiley & Sons Inc.

#### Example: $p_n(\mathbf{x})$ estimates assuming 5 samples:

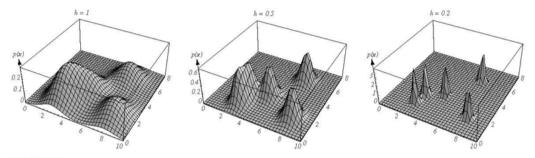
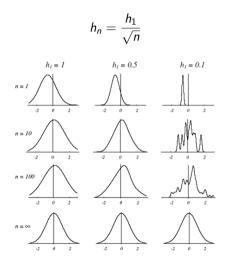


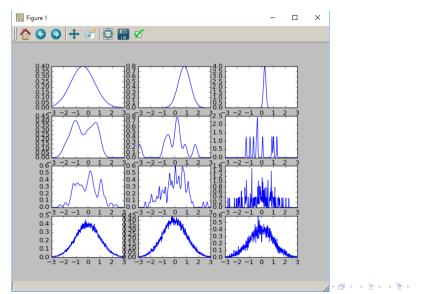
FIGURE 4.4. Three Parzen-window density estimates based on the same set of five samples, using the window functions in Fig. 4.3. As before, the vertical axes have been scaled to show the structure of each distribution. From: Richard O. Duda, Peter E. Hart, and David G. Stork, *Pattern Classification*. Copyright © 2001 by John Wiley & Sons, Inc.

Example: both  $p(\mathbf{x})$  and  $\varphi(u)$  are Gaussian

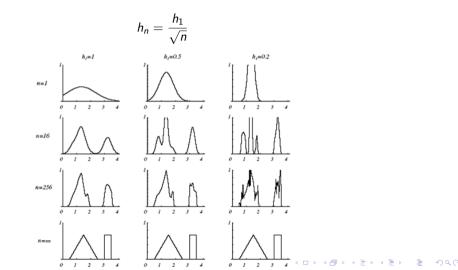


# Exercise (parzeng.py)

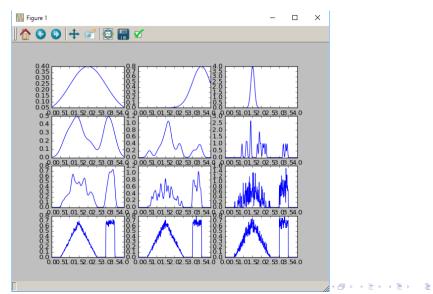
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Example:  $p(\mathbf{x})$  consists of a uniform and triangular density and  $\varphi(u)$  is Gaussian



# Exercise (parzentr.py)



Fix  $k_n$  and allow  $V_n$  to vary:

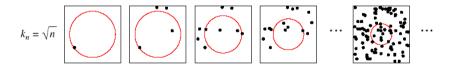
- Consider a hypersphere around **x**.
- Allow the radius of the hypersphere to grow until it contains  $k_n$  data points.

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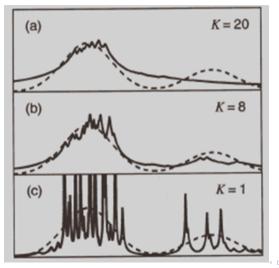
•  $V_n$  is determined by the volume of the hypersphere.

$$p_n(\mathbf{x}) pprox rac{k_n/n}{V_n}$$

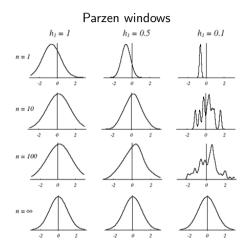
The size depends on the density

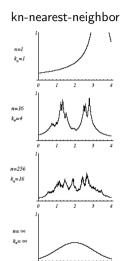


The parameter  $k_n$  acts as a smoothing parameter and needs to be optimized.



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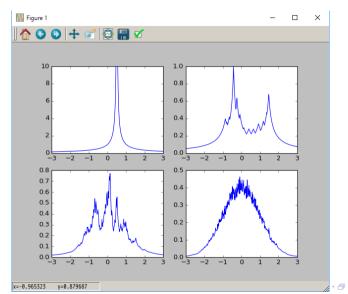




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# Exercise (knng.py)



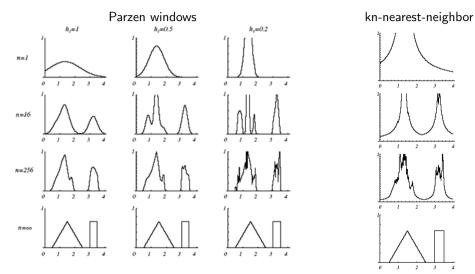
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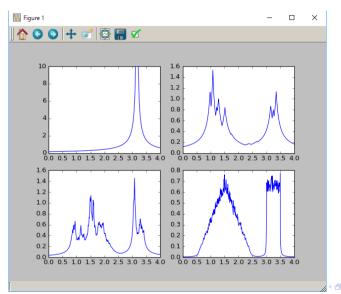
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# Exercise (knntr.py)



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#### Assignment

- Programming project and non-programming project are imposed.
- You are expected to solve either programming project OR non-programming project.
- Programming project is recommended.
- Of course you are most welcomed to solve both.
- Due on May 16.

# Programming project

- Download data file from the course web site. The file contains two variables: x1 and x2.
- (The data is in Matlab format. Use "loadmatfile" for Scilab or "scipy.io.loadmat" for python. Refer to the course material on April 13)
- Assume that they are samples of two classes c1 and c2.
- Plot conditional probability distributions p(x|ci) using Parzen windows (both Gaussian and box functions) and k-NN (with various k).
- Plot posterior probabilities  $P(c_i|x)$  assuming prior probabilities  $P(c_1) = P(c_2) = \frac{1}{2}$ .

#### Non-programming project

- The probability that a given vector  $\mathbf{x}$ , drawn from the unknown density  $p(\mathbf{x})$ , will fall inside some region R in the input space is assumed to be P.
- If we have *n* data points  $\{x_1, x_2, \ldots, x_n\}$  drawn independently from  $p(\mathbf{x})$ , we assume that *k* of the points will fall in *R*.
- Show that the expected value of k is:

$$E\{k\} = nP$$

• Show that the expected percentage of points falling in R is:

$$E\{\frac{k}{n}\}=P$$

• Show that the variance is given by:

$$Var\{\frac{k}{n}\} = E\{(\frac{k}{n} - P)^2\} = \frac{P(1 - P)}{n}$$

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