TIARA Summer School on Astrostatistics: Density Estimation and Regression

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Summer 2017

Density Estimation

Density Estimation: Introduction

- ► Recall that a statistical model views the data as random variables X₁, · · · , X_n from an unknown distribution function P(x).
- ► We further assume that such a distribution function has a probability density function (PDF) p(x).

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- ► Recall that a statistical model views the data as random variables X₁, · · · , X_n from an unknown distribution function P(x).
- ► We further assume that such a distribution function has a probability density function (PDF) p(x).
- In most cases, we do not know the PDF p(x) but we want to reconstruct it from the data.
- ► The goal of density estimation is to estimate p(x) using X₁,..., X_n.
- In other words, the parameter of interest is the PDF p(x).

Parametric Approach: Introduction

- The PDF p(x) is a function. It may not be easily estimated.
- A simple approach is to further assume that p(x) = p(x; θ) for some parameter θ.
- Then estimating p can be done by estimating θ .
- Estimating a value is much easier than estimating the entire function.

Parametric Approach: MLE

- We can estimate the parameters by the MLE.
- Here is an example of assuming the data being normally distributed: N(μ, σ²).
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- Here is an example of assuming the data being normally distributed: N(μ, σ²).
- ► In this case, we only need to estimate the two parameters μ, σ^2 .
- The estimators are

$$\widehat{\mu}_{MLE} = \overline{X}_n, \quad \widehat{\sigma}_{MLE}^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \overline{X}_n)^2.$$

Then the estimated density function is

$$\widehat{p}(x) = \frac{1}{\sqrt{2\pi\widehat{\sigma}_{MLE}^2}} e^{-\frac{(x-\widehat{\mu}_{MLE})^2}{2\widehat{\sigma}_{MLE}^2}}$$

Note that MLE is not the only approach to estimate the parameter; one can use method of moments or other approaches.

- Sometimes simple parametric models such as Gaussian, exponential, Gamma distribution are too restrictive to capture the complicated structure of the data.
- For instance, if the distribution has a bimodal density (two local maxima), none of these traditional model is reasonable.
- ▶ In this case, the *mixture model* may be useful.

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- ► For instance, Gaussian mixture model assumes

$$p(x) = \omega_1 \cdot \frac{1}{\sqrt{2\pi\sigma_1^2}} e^{-\frac{(x-\mu_1)^2}{2\sigma_1^2}} + \dots + \omega_K \cdot \frac{1}{\sqrt{2\pi\sigma_K^2}} e^{-\frac{(x-\mu_K)^2}{2\sigma_K^2}},$$

where $\omega_1 + \cdots + \omega_K = 1$ and $\omega_\ell > 0$.

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- The above model assumes that the PDF consists of K components and each component is a Gaussian.
- The quantities ω₁, · · · , ω_K are the mixing proportion of each component.
- In this case, the parameters are

$$\theta = (\omega_1, \mu_1, \sigma_1^2, \cdots, \omega_K, \mu_K, \sigma_K^2).$$

- We can estimate a mixture model using the MLE again.
- However, performing MLE in the mixture model is often computationally difficult and in general, there is no closed form solution to the MLE.
- People often use a method called *EM algorithm* to compute the MLE.

- We can estimate a mixture model using the MLE again.
- However, performing MLE in the mixture model is often computationally difficult and in general, there is no closed form solution to the MLE.
- People often use a method called *EM algorithm* to compute the MLE.
- In additional to the computational challenges, the identifiability is another issue of the mixture model (different parameters lead to the same PDF).

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- Parametric models only require estimating a few parameters to estimate the PDF.
- However, they are either too restrictive to capture the intricate structure of the PDF or computationally infeasible.
- An alternative approach is to estimate the PDF nonparametrically.
- Namely, we directly estimate the PDF without assuming a parametric form of the PDF.

Nonparametric Approach: Histogram

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- To make a histogram a density estimator, we need to rescale the Y-axis a bit.
- Instead of using the count of numbers observations within each bin, we need to divide the count by the total number of observations and the width of the bin.
- ► Assume our histogram has bins B₁, · · · , B_K and all bins have width L.
- ► For a point x within the bin B_ℓ, the density estimated by the histogram is

$$\widehat{p}_{hist}(x) = \frac{\# of X_1, \cdots, X_n \text{ within } B_\ell}{n \cdot L}.$$

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- ► It terms outs that when the size of bin L ≈ 0 and sample size n is large,

$$bias(\widehat{p}_{hist}(x)) = O(L), \quad Var(\widehat{p}_{hist}(x)) = O\left(\frac{1}{nL}\right).$$

Therefore, the MSE of the histogram estimator is

$$\mathsf{MSE}(\widehat{p}_{hist}(x)) = O(L^2) + O\left(\frac{1}{nL}\right).$$

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- Such a tradeoff between bias and variance is known as the bias-variance tradeoff.
- Moreover, it shows that we should choose *L* at the rate of $L \simeq n^{-1/3}$ to minimize the MSE.
- ► This choice leads to the optimal rate of histogram: MSE* $(\hat{p}_{hist}(x)) = O(n^{-2/3}).$

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- ► The KDE estimate the PDF using the following form:

$$\widehat{p}_{\mathsf{KDE}}(x) = \frac{1}{nh} \sum_{i=1}^{n} K\left(\frac{x - X_i}{h}\right),$$

where K(x) is a function called the kernel function and h > 0 is a quantity called smoothing bandwidth that controls the amount of smoothing.

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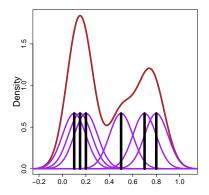
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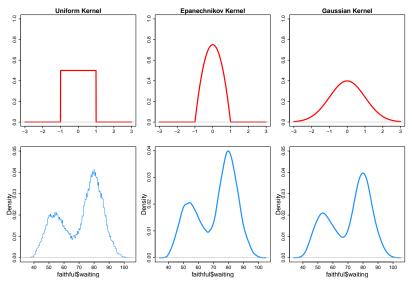
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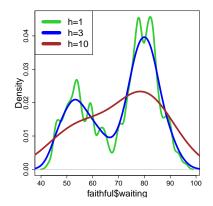
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- The idea of KDE is: we smooth out each data point using the kernel function into small bumps and then we sum over all bumps to obtain a density estimate.



Black dots: locations of observations. Purple bumps: the kernel function at each observation. Brown curve: final density estimate from KDE.



The kernel function generally does not affect the density estimate too much. \$15/52\$



The smoothing bandwidth often has a much stronger effect on the quality of estimation.

- We can also analyze the MSE of the KDE.
- ▶ When the smoothing bandwidth h ≈ 0 and sample size n is large,

$$\mathsf{bias}(\widehat{p}_{\mathsf{KDE}}(x)) = O(h^2), \quad \mathsf{Var}(\widehat{p}_{\mathsf{KDE}}(x)) = O\left(\frac{1}{nh}\right).$$

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Nonparametric Approach: Kernel Density Estimation - 5

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► The optimal choice of h is h ≈ n^{-1/5}, leading to the optimal convergence rate

$$\mathsf{MSE}^*(\widehat{p}_{\mathsf{KDE}}(x)) = O(n^{-4/5}).$$

► Note that this convergence rate is faster than the rate of histogram MSE*(p̂_{hist}(x)) = O(n^{-2/3}).

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- For a given point x (not necessarily an observation), its k-NN are the collection of observations whose distance to x is among the shortest k.
- ▶ Let R_k(x) be the distance from x to its k-th nearest neighbor observation.
- ► The k-NN density estimation uses the following approximation:

$$rac{k}{n} pprox P(X_{new} \in B(x, R_k(x))) pprox C_d R_k^d(x) \cdot p(x),$$

where d is the dimension of the data (often d = 1, 2, 3) and C_d is the size of d-dimensional unit ball and p(x) is the PDF.

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$$\widehat{p}_{knn}(x) = rac{k}{n} \cdot rac{1}{C_d R_k^d(x)}.$$

► When
$$d = 1$$
, $C_d = 2$ so
 $\frac{k}{n} \approx 2R_k(x) \cdot p(x)$, $\widehat{p}_{knn}(x) = \frac{k}{n} \cdot \frac{1}{2R_k(x)}$.
► When $d = 2$, $C_d = \pi$ so
 $\frac{k}{n} \approx \pi R_k^2(x) \cdot p(x)$, $\widehat{p}_{knn}(x) = \frac{k}{n} \cdot \frac{1}{\pi R_k^2(x)}$.
► When $d = 3$, $C_d = \frac{4}{3}\pi$ so
 $\frac{k}{n} \approx \frac{4}{3}\pi R_k^3(x) \cdot p(x)$, $\widehat{p}_{knn}(x) = \frac{k}{n} \cdot \frac{3}{4\pi R_k^3(x)}$.

And again, there will be a bias-variance tradeoff; in the case of d = 1, we have:

$$MSE(\widehat{p}_{knn}(x)) = \underbrace{O\left(\left(\frac{k}{n}\right)^{4}\right)}_{\text{bias}} + \underbrace{O\left(\frac{1}{k}\right)}_{\text{variance}}.$$

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- But note that there are two types of CI for a 'function'.
- Pointwise CI: given a point x and confidence level 1 − α, we construct an interval C_{1−α} = [ℓ_{1−α}, u_{1−α}] from the data such that

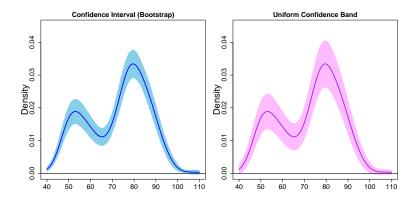
$$P(\ell_{1-\alpha} \leq p(x) \leq u_{1-\alpha}) \approx 1-\alpha.$$

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Simultaneous CB (confidence band): given α, we construct a band C_{1−α}(x) = [L_{1−α}(x), U_{1−α}(x)] from the data such that

$$P(L_{1-\alpha}(x) \le p(x) \le U_{1-\alpha}(x) \text{ for all } x) \approx 1-\alpha.$$



Pointwise CI (left) and simultaneous CB (right)¹.

¹A tutorial on this topic is in: https://arxiv.org/abs/1704.03924

Regression

Regression: Introduction

- Regression is an approach to study the relationship between a response variable Y and a covariate X.
- The covariate is also called a *feature*, a *predictor*, or an *independent variable*.
- ▶ Note that the covariate X can be multivariate.

Regression: Introduction

- Regression is an approach to study the relationship between a response variable Y and a covariate X.
- The covariate is also called a *feature*, a *predictor*, or an *independent variable*.
- Note that the covariate X can be multivariate.
- A traditional way to summarize the relationship via the regression function:

$$r(x) = \mathbb{E}(Y|X=x) = \int y \cdot f(y|x) dy.$$

► The goal of regression is to estimate r(x) using the random sample (X₁, Y₁), · · · , (X_n, Y_n).

Linear regression is a parametric approach that models the function r(x) as a linear function:

$$r(x) = \beta_0 + \beta_1 x.$$

 In many case, we will make further assumption on the noise and rewrite the linear model as

$$Y_i = \underbrace{\beta_0 + \beta_1 X_i}_{\text{signal}} + \underbrace{\epsilon_i}_{\text{noise}},$$

where $\mathbb{E}(\epsilon_i|X_i) = 0$ and $Var(\epsilon_i|X_i) = \sigma^2$.

In the linear regression model, there are two parameters: intercept β₀ and slope β₁.

- In the linear regression model, there are two parameters: intercept β₀ and slope β₁.
- To estimate them, a classical approach is the least squares (LS):

$$(\widehat{\beta}_0, \widehat{\beta}_1) = \operatorname{argmin}_{\beta_0, \beta_1} \sum_{i=1}^n (Y_i - \beta_0 - \beta_1 X_i)^2,$$

where the notation $\text{argmin}_{\beta_0,\beta_1}$ means finding the value of β_0,β_1 that minimizes the followings.

You can solve the above LS criterion and find a closed form solution to the estimate:

$$\widehat{\beta}_1 = \frac{\sum_{i=1}^n (X_i - \bar{X}_n)(Y_i - \bar{Y}_n)}{\sum_{i=1}^n (X_i - \bar{X}_n)^2}, \quad \widehat{\beta}_0 = \bar{Y}_n - \widehat{\beta}_1 \bar{X}_n.$$

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The difference between predicted and observed value is called residual

$$e_i = Y_i - \widehat{Y}_i = Y_i - \widehat{\beta}_0 - \widehat{\beta}_1 X_i.$$

► The residual sums of squares RSS = ∑ⁿ_{i=1} e²_i measures how our estimate fits the data.

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- ► The residual sums of squares RSS = ∑ⁿ_{i=1} e²_i measures how our estimate fits the data.
- You can interpret the LS approach as finding the best linear model to minimize RSS.
- ▶ Note that the noise level σ^2 can be estimated by $\widehat{\sigma^2} = \frac{1}{n-2} \sum_{i=1}^n e_i^2$.

The LSE has nice theoretical properties:

$$bias(\widehat{\beta}_0|X_1, \cdots, X_n) = 0, \quad bias(\widehat{\beta}_1|X_1, \cdots, X_n) = 0$$
$$Var(\widehat{\beta}_0|X_1, \cdots, X_n) = \frac{\sigma^2}{ns_X^2} \frac{1}{n} \sum_{i=1}^n X_i^2$$
$$Var(\widehat{\beta}_1|X_1, \cdots, X_n) = \frac{\sigma^2}{ns_X^2},$$

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where $s_X^2 = \frac{1}{n} \sum_{i=1}^n (X_i - \bar{X}_n)^2$.

- Moreover, central limit theorem implies that the LSE converges to a normal distribution under appropriate conditions.
- ► Thus, we can construct CI for β_0 and β_1 using the standard errors of $\hat{\beta}_0, \hat{\beta}_1$.

$$\operatorname{Var}(\widehat{\beta}_{0}|X_{1},\cdots,X_{n}) = \frac{\sigma^{2}}{ns_{X}^{2}}\frac{1}{n}\sum_{i=1}^{n}X_{i}^{2}$$
$$\longrightarrow SE(\widehat{\beta}_{0}) = \frac{\widehat{\sigma}}{s_{X}\sqrt{n}}\sqrt{\frac{\sum_{i=1}^{n}X_{i}^{2}}{n}}$$
$$\operatorname{Var}(\widehat{\beta}_{1}|X_{1},\cdots,X_{n}) = \frac{\sigma^{2}}{ns_{X}^{2}}$$
$$\longrightarrow SE(\widehat{\beta}_{1}) = \frac{\widehat{\sigma}}{s_{X}\sqrt{n}}.$$

Thus, a $1-\alpha$ CI will be

$$\widehat{\beta}_0 \pm z_{\alpha/2} SE(\widehat{\beta}_0), \quad \widehat{\beta}_1 \pm z_{\alpha/2} SE(\widehat{\beta}_1)$$

for β_0 and β_1 respectively.

Linear Regression: Multiple Covariates - 1

► In the case of the multiple covariates x = (x₁, · · · , x_p), the linear regression can be easily extended:

$$r(x) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \cdots + \beta_p x_p.$$

• Let $Y = (Y_1, \cdots, Y_n)$ be the vector of responses and

$$\mathbb{X} = \begin{pmatrix} 1 & X_{1,1} & \cdots & X_{1,p} \\ 1 & X_{2,1} & \cdots & X_{2,p} \\ \cdots & \cdots & \cdots & \cdots \\ 1 & X_{n,1} & \cdots & X_{n,p} \end{pmatrix}$$

be the $n \times (p + 1)$ data matrix (each row is an observation). The multiple linear regression can be written as the follows:

$$Y = \mathbb{X}\beta + \epsilon,$$

where $\beta = (\beta_0, \dots, \beta_p)$ is the parameter vector and $\epsilon = (\epsilon_1, \dots, \epsilon_n)$ is the noise.

Linear Regression: Multiple Covariates - 2

The LS method is to find

$$\widehat{\boldsymbol{\beta}} = \mathrm{argmax}_{\boldsymbol{\beta}} \| \boldsymbol{Y} - \mathbb{X} \boldsymbol{\beta} \|^2$$

And it has a closed form solution:

$$\widehat{\boldsymbol{\beta}} = \left(\mathbb{X}^{\mathsf{T}} \mathbb{X} \right)^{-1} \mathbb{X}^{\mathsf{T}} \boldsymbol{Y}.$$

Linear Regression: Multiple Covariates - 2

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$$\widehat{\beta} = \left(\mathbb{X}^T \mathbb{X} \right)^{-1} \mathbb{X}^T Y.$$

The LSE has a nice property that

$$\widehat{\boldsymbol{\beta}} \approx \boldsymbol{N} \left(\boldsymbol{\beta}, \sigma^2 \left(\boldsymbol{\mathbb{X}}^T \boldsymbol{\mathbb{X}} \right)^{-1} \right).$$

Actually, you can show that the LSE is an unbiased estimator and the variance is σ² (X^TX)⁻¹. The above expression further suggests that we can use it to construct a CI for β.

Linear Regression: Remarks

- The linear regression is an important topic in statistics. It can be a course for an entire semester!
- You can search online to learn more about it.
- Here are a few key words related to it: ANOVA, R², outliers, leverage points.
- Note that the idea of LS approach can be applied to 'non-linear' model as well. For instance, we can model

$$r(x) = \beta_0 + \beta_1 x + \beta_2 x^2 + \beta_3 exp(-\beta_4 x)$$

and apply LS approach to find the parameters.

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- If we naively model it as a linear function, then we may obtain a negative probability or a probability greater than 1, both are not reasonable.
- The logistic regression uses a smart way to model such a probability.

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The logistic regression models the log odds as a linear function of x:

$$\ell(x)=\beta_0+\beta_1x.$$

The model

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leads to the following form of r(x):

$$r(x) = P(Y = 1 | X = x) = \frac{e^{\beta_0 + \beta_1 x}}{1 + e^{\beta_0 + \beta_1 x}}.$$

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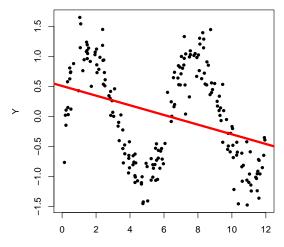
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- Using this probability model, we can then apply the MLE to find β₀ and β₁.
- Note that the MLE does not have a closed form solution but one can find it using numerical methods such a gradient descent approach.

Nonparametric Regression

- A problem of parametric regression is: the actual regression function may not have the desired form.
- When the parametric form is mis-specified, the result can be very bad.
- Nonparametric regression attempts to directly estimate the regression function without assuming a parametric form of it.
- We will talk about three popular methods: regressogram (binning), kernel regression, and spline approach.

Nonparametric Regression: Example



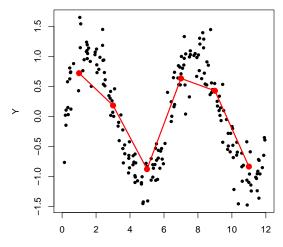
Regressogram (Binning)

- The regressogram (binning) might be one of the most popular regression approach but very few people know its name.
- ► The regressogram = regression + histogram.

Regressogram (Binning)

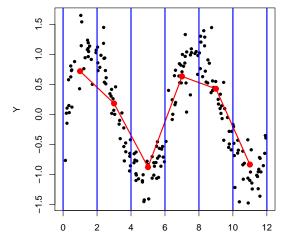
- The regressogram (binning) might be one of the most popular regression approach but very few people know its name.
- ► The regressogram = regression + histogram.
- The idea is: we bin the range of covariates into several intervals.
- We then use the average of the responses for observations within the same interval as the estimated value.

Regressogram: Example - 1

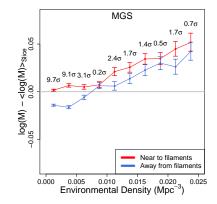


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Regressogram: Example - 1



Regressogram: Example - 2



- The kernel regression is another nonparametric regression estimator.
- ► The kernel regression uses an estimator of the form

$$\widehat{r}_{ker}(x) = \sum_{i=1}^{n} W_i(x) Y_i$$
$$= \frac{\sum_{i=1}^{n} K\left(\frac{X_i - x}{h}\right) Y_i}{\sum_{j=1}^{n} K\left(\frac{X_j - x}{h}\right)}$$

where

$$W_i(x) = rac{K\left(rac{X_i-x}{h}
ight)}{\sum_{j=1}^n K\left(rac{X_j-x}{h}
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► The function K(x) is again the kernel function we talk about in the KDE.

The quantity

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- The estimator $\hat{r}_{ker}(x) = \sum_{i=1}^{n} W_i(x) Y_i$ can be interpreted as follows.
- ► To estimate the regression function at X = x, we use a weighted average of all responses such that observations close to x will be given a higher weight (W_i(x) will be large if X_i is close to x).

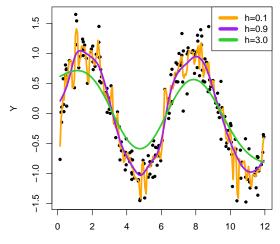
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- ► To estimate the regression function at X = x, we use a weighted average of all responses such that observations close to x will be given a higher weight (W_i(x) will be large if X_i is close to x).
- The kernel function determines how we are going to give weights to the nearby points.
- The smoothing bandwidth h controls the range of influence from each observation (the degree of smoothing).

Kernel Regression: Example



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$$R = \mathbb{E}(|Y_{new} - \widehat{m}(X_{new})|^2),$$

where (X_{new}, Y_{new}) is a new observation.

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We want to pick the smoothing bandwidth

$$h^* = \operatorname{argmin}_h R(h).$$

- ► The quantity R(h) = E(|Y_{new} m̂_{ker}(X_{new})|²) is unknown to us we need to estimate it.
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- This idea is called data splitting.
- ▶ The *cross-validation* is a modified approach of data splitting that repeat the splitting procedure multiple times and then use the average as the final estimate of *R*(*h*).

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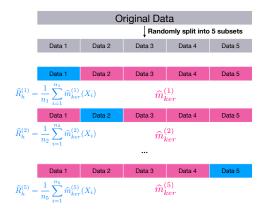
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- We often repeat the above procedure several times and take the total average as the the final risk estimate.
- ▶ Note: if we split the data into *k* subset, we call this approach the k-fold cross validation.

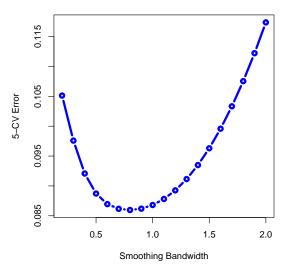
Cross-Validation Approach: 5-fold CV

► Here is an illustration for 5-fold CV:



- ► Validation Set. Training Set.
- We use the average $\widehat{R}(h) = \frac{1}{5} \sum_{\ell=1}^{5} \widehat{R}^{(\ell)}(h)$ as a risk estimate.
- In practice, we repeat this procedure for several times and take the total average of them.

5-fold Cross-Validation: Example



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- As an extreme example: consider h ≈ 0, then the kernel regression passes every data point. If we use the training set as the validation set, this leads to a prediction risk = 0!
- Note that: an opposite case is called underfitting you fit a too easy model so it cannot capture the complicated structure of the data. When we apply the linear regression to the example of a wave-form data, we suffer from underfitting.

- Spline approach is a *penalized regression* method.
- The goal is to find a function f such that it fits the data well and f is smooth.

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- ▶ In more details, the spline approach attempts to find \hat{f}_{sp} such that

$$\widehat{f}_{sp} = \operatorname{argmin}_{f} \frac{1}{n} \sum_{i=1}^{n} (Y_i - f(X_i))^2 + \lambda \int_{X_{min}}^{X_{max}} |f''(s)|^2 ds,$$

where $\lambda > 0$ is a parameter determines how smooth we want.

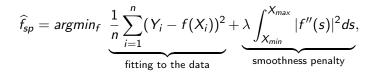
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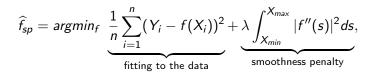
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where $\lambda > 0$ is a parameter determines how smooth we want. • There are some smart ways² to find such a minimal function \hat{f}_{sp} .

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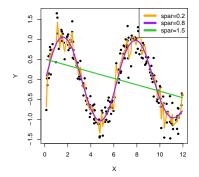


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- A large λ leads to a smooth function $\widehat{f_{sp}}$.
- A small λ yields a more wiggly function.
- The choice of λ determines how we want to weight the fitting quality and smoothness.
- We often use cross-validation to choose λ .

Spline Approach: Example



spar: a quantity in R related to λ .

Useful References

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