

Advanced statistics and modelling

2023 spring

Statistical models

Regression

Point estimation
and bias

Mean squared
error

Confidence
interval

Empirical CDF

Empirical PDF
Cross validation

Kernel density
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MODELS, INFERENCE, LEARNING

The basis statistical inference problem

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- The basic statistical inference problem is the following:
 - We observe $X_1, X_2, \dots, X_n \sim F$.
 - Based on the observations we would like to infer (or estimate or learn) F or some feature of F (such as e.g., its mean or variance).
- In computer science statistical inference is usually called as "learning".

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Statistical model

- A **statistical model** is a set of distributions \mathcal{F} .

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Statistical model

- A **statistical model** is a set of distributions \mathfrak{F} .
- A **parametric model** is a set of distributions that can be parametrised with a finite number of parameters,

$$\mathfrak{F} = \{\rho(x \mid \theta); \theta \in \Theta\},$$

where θ is an unknown parameter (or a vector of parameters) that can take values in the parameter space Θ .

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E.g., if the data comes from a Normal distribution, then

$$\mathfrak{F} = \left\{ \rho(x \mid \mu, \sigma) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x-\mu)^2}{2\sigma^2}}, \mu \in \mathbb{R}, \sigma > 0 \right\}$$

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- If θ is a vector, but we are interested in only a part of the parameters, then the remaining parameters are called **nuisance parameters**.

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- If θ is a vector, but we are interested in only a part of the parameters, then the remaining parameters are called **nuisance parameters**.
- A **non-parametric model** is a set \mathfrak{F} that cannot be parametrised by a finite number of parameters. E.g., the set of all possible CDF-s, $\mathfrak{F} = \{ \text{all CDF's} \}$ is a non-parametric model.

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- Example for parametric estimation: Let x_1, x_2, \dots, x_n be independent Bernoulli(p) observations. The problem is to estimate p .

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- Example for non-parametric estimation: Let x_1, x_2, \dots, x_n be independent observations, and we would like to estimate the PDF assuming that $\mathcal{F} = \{ \text{all PDF's} \}$.

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- If $\mathfrak{F} = \{ \rho(x | \theta) : \theta \in \Theta \}$ is a parametric model, then e.g., $P_\theta(X \in A)$ or $\mathbb{E}_\theta(f(X))$ mean that

$$P_\theta(X \in A) = \int_A \rho(x | \theta) dx, \quad \mathbb{E}_\theta(f(x)) = \int f(x) \rho(x | \theta) dx,$$

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thus, the subscript θ indicates that the given probability or expectation is depending on θ and was taken with respect to $\rho(x | \theta)$, and it does NOT indicate that we were averaging over θ !

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Suppose we observe pairs of data $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$. E.g., X corresponds to the blood pressure of patients, and Y is how long they live.

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- The variables X and Y are called

$$X : \begin{cases} \text{predictor,} \\ \text{regressor,} \\ \text{feature variable,} \\ \text{independent variable} \end{cases}$$
$$Y : \begin{cases} \text{outcome,} \\ \text{response variable,} \\ \text{dependent variable} \end{cases}$$

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- The **regression function** is given by the conditional expectation of Y given X , written as

$$r(x) = \mathbb{E}(Y \mid X = x).$$

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- If $r(x) \in \mathfrak{F}$ where \mathfrak{F} can be parametrised by a finite number of parameters (e.g., all possible straight lines), then we have a **parametric regression model**.

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- If the goal is to predict Y for new patients based on their X value, that is called **prediction**.

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- If Y is discrete (e.g., live or die), then the problem is called **classification**.

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- If the goal is to predict Y for new patients based on their X value, that is called **prediction**.
- If Y is discrete (e.g., live or die), then the problem is called **classification**.
- If the goal is to estimate the curve $r(x)$, then this is called **curve estimation** or **regression**.

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Regression models can always be written in the form of

$$Y = r(x) + \epsilon,$$

where $\mathbb{E}(\epsilon) = 0$.

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Regression models can always be written in the form of

$$Y = r(x) + \epsilon,$$

where $\mathbb{E}(\epsilon) = 0$.

Proof:

Since $\epsilon = Y - r(x) = Y - \mathbb{E}(Y | X)$ we can write

$$\mathbb{E}(\epsilon) = \mathbb{E}[Y - \mathbb{E}(Y | X)] = \mathbb{E}(Y) - \mathbb{E}[\mathbb{E}(Y | X)] = \mathbb{E}(Y) - \mathbb{E}(Y) = 0.$$

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- **Point estimation** refers to providing a single "best guess" of some quantity.

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- **Point estimation** refers to providing a single "best guess" of some quantity.
- This quantity can be a parameter in the model, a CDF or PDF, a regression function $r(x)$, a prediction for a future value of Y , etc.

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- By convention we denote the point estimate of θ by $\hat{\theta}$.

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Note that

θ : FIXED unknown quantity,

$\hat{\theta}$: RANDOM VARIABLE, that depends on the data.

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Point estimation and bias

- Formally, if X_1, X_2, \dots, X_n are IID data points from some distribution F , then a **point estimator** $\widehat{\theta}$ of a parameter θ is some function of X_1, X_2, \dots, X_n :

$$\widehat{\theta} = g(X_1, X_2, \dots, X_n).$$

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$$\hat{\theta} = g(X_1, X_2, \dots, X_n).$$

- The **bias** corresponds to the difference between the mean of $\hat{\theta}$ and the true value of the parameter,

$$\text{bias}(\hat{\theta}) = \mathbb{E}_{\theta}(\hat{\theta}) - \theta.$$

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- A point estimator $\hat{\theta}$ is **unbiased** if $\text{bias}(\hat{\theta}) = 0$.

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- The distribution of $\hat{\theta}$ is called the **sampling distribution**.

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- The distribution of $\hat{\theta}$ is called the **sampling distribution**.
- The standard deviation of $\hat{\theta}$ is called as the **standard error**,

$$\text{se}(\hat{\theta}) = \sqrt{\mathbb{V}_{\theta}(\hat{\theta})}.$$

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- The distribution of $\hat{\theta}$ is called the **sampling distribution**.
- The standard deviation of $\hat{\theta}$ is called as the **standard error**,

$$\text{se}(\hat{\theta}) = \sqrt{\mathbb{V}_{\theta}(\hat{\theta})}.$$

- Often, it is not possible to compute $\text{se}(\hat{\theta})$. However, usually we can estimate it, and the estimated standard error is denoted by $\widehat{\text{se}}(\hat{\theta})$.

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Example:

- Assume $X_1, X_2, \dots, X_n \sim \text{Bernoulli}(p)$.

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Example:

- Assume $X_1, X_2, \dots, X_n \sim \text{Bernoulli}(p)$.

$$\rightarrow \hat{p} = \overline{X_n} = \frac{1}{n} \sum_{i=1}^n x_i.$$

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$$\rightarrow \hat{p} = \overline{X_n} = \frac{1}{n} \sum_{i=1}^n x_i.$$

\rightarrow Since $\mathbb{E}(\hat{p}) = \frac{1}{n} \sum_{i=1}^n \mathbb{E}(X_i) = p$, the \hat{p} is unbiased.

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\rightarrow Since $\mathbb{E}(\hat{p}) = \frac{1}{n} \sum_{i=1}^n \mathbb{E}(X_i) = p$, the \hat{p} is unbiased.

\rightarrow The standard error: $\text{se}(\hat{p}) = \sqrt{\mathbb{V}(\hat{p})} = \sqrt{p(1-p)/n}$.

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$$\rightarrow \text{Since } \mathbb{E}(\hat{p}) = \frac{1}{n} \sum_{i=1}^n \mathbb{E}(X_i) = p, \text{ the } \hat{p} \text{ is unbiased.}$$

$$\rightarrow \text{The standard error: } \text{se}(\hat{p}) = \sqrt{\mathbb{V}(\hat{p})} = \sqrt{p(1-p)/n}.$$

$$\rightarrow \text{The estimated standard error: } \widehat{\text{se}}(\hat{p}) = \sqrt{\hat{p}(1-\hat{p})/n}.$$

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Mean squared error

The quality of a point estimate is often measured by the **mean squared error**, defined as

$$\text{MSE}(\widehat{\theta}) = \mathbb{E}_{\theta} [(\widehat{\theta} - \theta)^2].$$

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The quality of a point estimate is often measured by the **mean squared error**, defined as

$$\text{MSE}(\widehat{\theta}) = \mathbb{E}_{\theta} [(\widehat{\theta} - \theta)^2].$$

Note that $\mathbb{E}_{\theta}(\cdot)$ refers to the expectation with respect to

$$\rho(x_1, x_2, \dots, x_n \mid \theta) = \prod_{i=1}^n \rho(x_i \mid \theta)$$

that generated the data, and it does NOT mean that we are averaging with respect to some density of θ .

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MSE and bias

The MSE can always be written as

$$\text{MSE} = [\text{bias}]^2 + \mathbb{V}.$$

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Proof:

Let us denote the mean of $\hat{\theta}$ as $\mathbb{E}_{\theta}(\hat{\theta}) = \bar{\theta}$.

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Proof:

Let us denote the mean of $\widehat{\theta}$ as $\mathbb{E}_{\theta}(\widehat{\theta}) = \bar{\theta}$. Based on that

$$\begin{aligned}\text{MSE}_{\theta}(\widehat{\theta}) &= \mathbb{E}_{\theta}[(\widehat{\theta} - \theta)^2] = \mathbb{E}_{\theta}[(\widehat{\theta} - \bar{\theta} + \bar{\theta} - \theta)^2] = \\ &\underbrace{\mathbb{E}_{\theta}[(\widehat{\theta} - \bar{\theta})^2]}_{\mathbb{V}(\widehat{\theta})} + 2(\bar{\theta} - \theta) \underbrace{\mathbb{E}_{\theta}(\widehat{\theta} - \bar{\theta})}_0 + \underbrace{\mathbb{E}_{\theta}[(\bar{\theta} - \theta)^2]}_{(\bar{\theta} - \theta)^2} = \\ &\mathbb{V}_{\theta}(\widehat{\theta}) + (\bar{\theta} - \theta)^2 = \mathbb{V}_{\theta}(\widehat{\theta}) + [\text{bias}(\widehat{\theta})]^2.\end{aligned}$$

MSE and bias

MSE and bias

The MSE can always be written as

$$\text{MSE} = [\text{bias}]^2 + \mathbb{V}.$$

Proof:

Let us denote the mean of $\widehat{\theta}$ as $\mathbb{E}_{\theta}(\widehat{\theta}) = \bar{\theta}$. Based on that

$$\begin{aligned}\text{MSE}_{\theta}(\widehat{\theta}) &= \mathbb{E}_{\theta}[(\widehat{\theta} - \theta)^2] = \mathbb{E}_{\theta}[(\widehat{\theta} - \bar{\theta} + \bar{\theta} - \theta)^2] = \\ &= \underbrace{\mathbb{E}_{\theta}[(\widehat{\theta} - \bar{\theta})^2]}_{\mathbb{V}(\widehat{\theta})} + \underbrace{2(\bar{\theta} - \theta)\mathbb{E}_{\theta}(\widehat{\theta} - \bar{\theta})}_0 + \underbrace{\mathbb{E}_{\theta}[(\bar{\theta} - \theta)^2]}_{(\bar{\theta} - \theta)^2} = \\ &= \mathbb{V}_{\theta}(\widehat{\theta}) + (\bar{\theta} - \theta)^2 = \mathbb{V}_{\theta}(\widehat{\theta}) + [\text{bias}(\widehat{\theta})]^2.\end{aligned}$$

Consequence:

If $\widehat{\theta}$ is unbiased, the MSE is simply the variance of $\widehat{\theta}$.

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Asymptotically normal estimator

The estimator $\hat{\theta}$ is asymptotically normal if

$$\frac{\hat{\theta} - \theta}{\text{se}} \xrightarrow{d} N(0, 1).$$

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Confidence interval

A $1 - \alpha$ **confidence interval** for a parameter θ is an interval $C = [a, b]$ where $a = a(X_1, X_2, \dots, X_n)$ and $b = b(X_1, X_2, \dots, X_n)$ are functions of the data such that

$$P_{\theta}(\theta \in C) \geq 1 - \alpha, \quad \forall \theta \in \Theta.$$

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$$P_{\theta}(\theta \in C) \geq 1 - \alpha, \quad \forall \theta \in \Theta.$$

- The **coverage** of the confidence interval is $1 - \alpha$.

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$$P_{\theta}(\theta \in C) \geq 1 - \alpha, \quad \forall \theta \in \Theta.$$

- The **coverage** of the confidence interval is $1 - \alpha$.
- Note: θ is fixed, and C_n is random.

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Example:

- Assume a coin tossing experiment, where $\hat{p} = \overline{X_n} = \frac{1}{n} \sum_{i=1}^n x_i$, and let the confidence interval be $[\hat{p} - \epsilon, \hat{p} + \epsilon]$. How should we choose ϵ for a given value of α ?

Confidence interval

Example:

- Assume a coin tossing experiment, where $\hat{p} = \overline{X_n} = \frac{1}{n} \sum_{i=1}^n x_i$, and let the confidence interval be $[\hat{p} - \epsilon, \hat{p} + \epsilon]$. How should we choose ϵ for a given value of α ?
- According to Hoeffding's inequality:
- If $X_1, X_2, \dots, X_n \sim \text{Bernoulli}(p)$, then for the sample mean $\overline{X_n}$ we can write

$$P(|\overline{X_n} - p| \geq \epsilon) \leq 2e^{-2n\epsilon^2}.$$

Confidence interval

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Thus, if we choose

$$\epsilon = \sqrt{\frac{1}{2n} \ln\left(\frac{2}{\alpha}\right)},$$

Confidence interval

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Thus, if we choose

$$\epsilon = \sqrt{\frac{1}{2n} \ln\left(\frac{2}{\alpha}\right)},$$

then according to the inequality

$$P(|\widehat{p} - p| > \epsilon) \leq 2e^{-2n\epsilon^2} = \alpha.$$

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Normal-based confidence interval

If we have a point estimator $\hat{\theta}$ with limiting normal distribution, then we can assume that (based on the finite data we have) $\hat{\theta} \approx N(\theta, \sigma^2)$.

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Based on that, let us choose the confidence interval as

$$C = [\hat{\theta} - z_{\alpha/2} \widehat{se}, \hat{\theta} + z_{\alpha/2} \widehat{se}],$$

where $z_{\alpha/2} = \Phi^{-1}(1 - \alpha/2)$ with $\Phi(z)$ denoting the CDF of the standard Normal distribution.

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where $z_{\alpha/2} = \Phi^{-1}(1 - \alpha/2)$ with $\Phi(z)$ denoting the CDF of the standard Normal distribution.

This way $P(-z_{\alpha/2} < z < z_{\alpha/2}) = 1 - \alpha$ and for $\hat{\theta}$ we can write

$$P(\theta \in C) \rightarrow 1 - \alpha.$$

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This means that $C = [\hat{\theta} - z_{\alpha/2} \widehat{\text{se}}, \hat{\theta} + z_{\alpha/2} \widehat{\text{se}}]$ provides an **approximate** confidence interval.

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where $z_{\alpha/2} = \Phi^{-1}(1 - \alpha/2)$ with $\Phi(z)$ denoting the CDF of the standard Normal distribution.

This way $P(-z_{\alpha/2} < Z < z_{\alpha/2}) = 1 - \alpha$ and for $\hat{\theta}$ we can write

$$P(\theta \in C) \rightarrow 1 - \alpha.$$

This means that $C = [\hat{\theta} - z_{\alpha/2} \widehat{\text{se}}, \hat{\theta} + z_{\alpha/2} \widehat{\text{se}}]$ provides an **approximate** confidence interval.

Proof:

Let us define $Z = (\hat{\theta} - \theta) / \widehat{\text{se}}$. Assuming $Z \xrightarrow{d} N(0, 1)$, we have

$$P_{\theta}(\theta \in C) = P_{\theta}(\hat{\theta} - z_{\alpha/2} \widehat{\text{se}} < \theta < \hat{\theta} + z_{\alpha/2} \widehat{\text{se}}) =$$

$$P_{\theta} \left(-z_{\alpha/2} < \frac{\hat{\theta} - \theta}{\widehat{\text{se}}} < z_{\alpha/2} \right) \longrightarrow P(-z_{\alpha/2} < Z < z_{\alpha/2}) = 1 - \alpha$$

Confidence interval

Example:

- Assuming $X_1, X_2, \dots, X_n \sim \text{Bernoulli}(p)$ we have

$$\hat{p} = \frac{1}{n} \sum_{i=1}^n x_i, \quad \mathbb{V}(\hat{p}) = \frac{1}{n^2} \sum_{i=1}^n \mathbb{V}(X_i) = \frac{1}{n^2} \sum_{i=1}^n p(1-p) = \frac{p(1-p)}{n}.$$

$$\text{se} = \sqrt{\frac{p(1-p)}{n}} \quad \hat{\text{se}} = \sqrt{\frac{\hat{p}(1-\hat{p})}{n}}.$$

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- Based on the Central Limit Theorem we can assume $\widehat{p} \approx N(p, \widehat{\text{se}}^2)$, thus, an approximate $1 - \alpha$ confidence interval can be given as

$$[\widehat{p} - z_{\alpha/2} \widehat{\text{se}}, \widehat{p} + z_{\alpha/2} \widehat{\text{se}}] = \left[\widehat{p} - z_{\alpha/2} \sqrt{\frac{\widehat{p}(1-\widehat{p})}{n}}, \widehat{p} + z_{\alpha/2} \sqrt{\frac{\widehat{p}(1-\widehat{p})}{n}} \right]$$

Confidence interval

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- This interval is shorter compared to the one given based on Hoeffding's inequality.

Confidence interval

Example:

- Assuming $X_1, X_2, \dots, X_n \sim \text{Bernoulli}(p)$ we have

$$\begin{aligned}\widehat{p} &= \frac{1}{n} \sum_{i=1}^n x_i, & \mathbb{V}(\widehat{p}) &= \frac{1}{n^2} \sum_{i=1}^n \mathbb{V}(X_i) = \frac{1}{n^2} \sum_{i=1}^n p(1-p) = \frac{p(1-p)}{n}. \\ \text{se} &= \sqrt{\frac{p(1-p)}{n}} & \widehat{\text{se}} &= \sqrt{\frac{\widehat{p}(1-\widehat{p})}{n}}.\end{aligned}$$

- Based on the Central Limit Theorem we can assume $\widehat{p} \approx N(p, \widehat{\text{se}}^2)$, thus, an approximate $1 - \alpha$ confidence interval can be given as

$$[\widehat{p} - z_{\alpha/2} \widehat{\text{se}}, \widehat{p} + z_{\alpha/2} \widehat{\text{se}}] = \left[\widehat{p} - z_{\alpha/2} \sqrt{\frac{\widehat{p}(1-\widehat{p})}{n}}, \widehat{p} + z_{\alpha/2} \sqrt{\frac{\widehat{p}(1-\widehat{p})}{n}} \right]$$

- This interval is shorter compared to the one given based on Hoeffding's inequality.
- However, this is only an **approximate** confidence interval, whereas in case of Hoeffding's inequality we have guarantee for smaller sample sizes as well.

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Empirical CDF

Based on IID variables X_1, X_2, \dots, X_n , the empirical distribution function $\widehat{F}_n(x)$ is defined as the fraction of observed data points falling below x , which can be formulated e.g., as

$$\widehat{F}(x) = \frac{1}{n} |\{x_i \mid x_i < x\}| = \frac{1}{n} \sum_{i=1}^n I(X_i < x),$$

where $I(X_i < x) = 1$ if $x_i < x$ and $I(X_i < x) = 0$ otherwise.

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where $I(X_i < x) = 1$ if $x_i < x$ and $I(X_i < x) = 0$ otherwise.

Glivenko-Cantelli theorem

Let $X_1, X_2, \dots, X_n \sim F$ be IID variables. Then

$$\sup_x |\widehat{F}_n(x) - F(x)| \xrightarrow{\text{a.s.}} 0.$$

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$$\widehat{F}_n(x) = \frac{1}{n} |\{x_i \mid x_i < x\}| = \frac{1}{n} \sum_{i=1}^n I(X_i < x),$$

where $I(X_i < x) = 1$ if $x_i > x$ and $I(X_i < x) = 0$ otherwise.

Glivenko-Cantelli theorem

Let $X_1, X_2, \dots, X_n \sim F$ be IID variables. Then

$$\sup_x |\widehat{F}_n(x) - F(x)| \xrightarrow{\text{a.s.}} 0.$$

Note that for any fixed x , the sequence of $\widehat{F}_n(x)$ is a sequence of random variables that is converging to $F(x)$ according to the law of large numbers. The theorem above strengthens this to the uniform convergence of \widehat{F}_n to F .

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Dvoretzky-Kiefer-Wolfowitz inequality

Let $X_1, X_2, \dots, X_n \sim F$ be IID variables. Then for any $\epsilon > 0$

$$P\left(\sup_x |\widehat{F}_n(x) - F(x)| > \epsilon\right) \leq 2e^{-2n\epsilon^2}.$$

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Dvoretzky-Kiefer-Wolfowitz inequality

Let $X_1, X_2, \dots, X_n \sim F$ be IID variables. Then for any $\epsilon > 0$

$$P\left(\sup_x |\widehat{F}_n(x) - F(x)| > \epsilon\right) \leq 2e^{-2n\epsilon^2}.$$

Note that for any fixed x , the sequence of $\widehat{F}_n(x)$ is a sequence of bounded random variables for which we can apply Hoeffding's inequality. The above theorem strengthens this by providing a uniform bound.

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Application:

We can use the DKW inequality for providing a **confidence band** for the empirical CDF.

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Application:

We can use the DKW inequality for providing a **confidence band** for the empirical CDF.

- Let us choose first an α value, and the goal is to give a confidence band (confidence set) C_n such that

$$P(F \in C_n) \geq 1 - \alpha.$$

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Application:

We can use the DKW inequality for providing a **confidence band** for the empirical CDF.

- Let us choose first an α value, and the goal is to give a confidence band (confidence set) C_n such that

$$P(F \in C_n) \geq 1 - \alpha.$$

- According to the above theorem, for any x the lower bound can be given as $L(x) = \widehat{F}_n(x) - \epsilon_n$ and the upper bound can be given as $U(x) = \widehat{F}_n(x) + \epsilon_n$ where

$$\epsilon_n = \sqrt{\frac{1}{2n} \ln \left(\frac{2}{\alpha} \right)}.$$

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Empirical PDF (histogram estimator)

Given $X_1, X_2, \dots, X_n \sim F$ IID variables the **empirical PDF** $\hat{\rho}_n(x)$ provides a simple estimate for the PDF $\rho(x)$.

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Empirical PDF (histogram estimator)

Given $X_1, X_2, \dots, X_n \sim F$ IID variables the **empirical PDF** $\hat{\rho}_n(x)$ provides a simple estimate for the PDF $\rho(x)$.

- Let's assume altogether m bins of equal bin width h .

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- Let's assume altogether m bins of equal bin width h .
- If the number of observations in bin B_i is ν_i , then the estimate for the probability of this bin is $\hat{p}_i = \nu_i/n$, (whereas the true probability is $p_i = \int_{B_i} \rho(x)dx$).

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- If the number of observations in bin B_i is ν_i , then the estimate for the probability of this bin is $\hat{p}_i = \nu_i/n$, (whereas the true probability is $p_i = \int_{B_i} \rho(x)dx$).
- Based on that, the empirical PDF can be given as

$$\hat{\rho}_n(x) = \sum_{i=1}^m \frac{\hat{p}_i}{h} I(x \in B_i),$$

where $I(x \in B_i)$ is the indicator function of B_i , (i.e., $I(x \in B_i) = 1$ if $x \in B_i$ and $I(x \in B_i) = 0$ otherwise.)

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- How to measure the goodness of the fit between an estimate $\widehat{\rho}_n(x)$ of the PDF and the true PDF $\rho(x)$ itself?

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- How to measure the goodness of the fit between an estimate $\widehat{\rho}_n(x)$ of the PDF and the true PDF $\rho(x)$ itself?
- In general, we can use the **integrated squared error**.

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- How to measure the goodness of the fit between an estimate $\widehat{\rho}_n(x)$ of the PDF and the true PDF $\rho(x)$ itself?
- In general, we can use the **integrated squared error**.

Risk (MISE)

For a statistical estimate $\widehat{\rho}_n(x)$ of the PDF $\rho(x)$ we can define the integrated squared error simply as

$$L(\widehat{\rho}_n, \rho) = \int (\widehat{\rho}_n(x) - \rho(x))^2 dx.$$

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- How to measure the goodness of the fit between an estimate $\widehat{\rho}_n(x)$ of the PDF and the true PDF $\rho(x)$ itself?
- In general, we can use the **integrated squared error**.

Risk (MISE)

For a statistical estimate $\widehat{\rho}_n(x)$ of the PDF $\rho(x)$ we can define the integrated squared error simply as

$$L(\widehat{\rho}_n, \rho) = \int (\widehat{\rho}_n(x) - \rho(x))^2 dx.$$

The expected value of this is the **mean integrated squared error** (MISE) or **risk**:

$$R(\widehat{\rho}_n, \rho) = \mathbb{E} (L(\widehat{\rho}_n, \rho)) .$$

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- The **bias** of $\widehat{\rho}_n(x)$ at a given x can be defined as

$$b(x) = \mathbb{E}(\widehat{\rho}_n(x)) - \rho(x).$$

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- The **bias** of $\widehat{\rho}_n(x)$ at a given x can be defined as

$$b(x) = \mathbb{E}(\widehat{\rho}_n(x)) - \rho(x).$$

- The **variation** of $\widehat{\rho}_n(x)$ at a given x can be given as

$$v(x) = \mathbb{V}(\widehat{\rho}_n(x)) = \mathbb{E}[(\widehat{\rho}_n(x) - \mathbb{E}(\widehat{\rho}_n(x)))^2].$$

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→ The MISE (or risk) can be written as

$$\begin{aligned} R(\widehat{\rho}_n, \rho) &= \mathbb{E}\left(\int (\mathbb{E}(\widehat{\rho}_n(x)) - \rho(x))^2\right) = \dots \\ &= \int b^2(x)dx + \int v(x)dx. \end{aligned}$$

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Thus, in other words,

$$\text{RISK} = \text{BIAS}^2 + \text{VARIANCE}.$$

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Let's calculate the bias and the variation for the histogram estimator.

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Let's calculate the bias and the variation for the histogram estimator.

- For a given x , let's assume that x is in bin B_j . Then

$$\mathbb{E}(\widehat{\rho}_n(x)) = \frac{p_j}{h} = \frac{1}{h} \int_{B_j} \rho(x) dx,$$

$$\mathbb{V}(\widehat{\rho}_n(x)) = \frac{p_j(1-p_j)}{nh^2},$$

since the number of observations ν_j follows a binomial distribution with parameter p_j .

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since the number of observations ν_j follows a binomial distribution with parameter p_j .

- Let's take another point u in the same bin. Since bins are usually small, we can approximate the true ρ at this point as

$$\rho(u) \approx \rho(x) + (u-x)\rho'(x).$$

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$$\rho(u) \approx \rho(x) + (u-x)\rho'(x).$$

Based on that, the probability p_j can be approximated as

$$p_j = \int_{x_{B_j}}^{x_{B_j}+h} \rho(u) du \approx \int_{x_{B_j}}^{x_{B_j}+h} (\rho(x) + (u-x)\rho'(x)) du =$$

$$h\rho(x) - h x \rho'(x) + \rho'(x) \left[u^2/2 \right]_{x_{B_j}}^{x_{B_j}+h} = h\rho(x) + h\rho'(x) \left[\frac{h}{2} + x_{B_j} - x \right]$$

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- Based on that, bias can be given as

$$b(x) = \mathbb{E}(\widehat{\rho}_n(x)) - \rho(x) = \frac{p_j}{h} - \rho(x) \approx \frac{h\rho(x) + h\rho'(x) \left[\frac{h}{2} + x_{B_j} - x \right]}{h} - \rho(x) = \rho'(x) \left[\frac{h}{2} + x_{B_j} - x \right]$$

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- The integral of $b^2(x)$ over the bin can be approximated as

$$\int_{x_{B_j}}^{x_{B_j}+h} b^2(x) dx \approx \int_{x_{B_j}}^{x_{B_j}+h} [\rho'(x)]^2 \left(\frac{h}{2} + x_{B_j} - x\right)^2 dx = \dots = [\rho'(x)]^2 \frac{h^3}{12}.$$

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$$\int_{x_{B_j}}^{x_{B_j}+h} b^2(x) dx \approx \int_{x_{B_j}}^{x_{B_j}+h} [\rho'(x)]^2 \left(\frac{h}{2} + x_{B_j} - x\right)^2 dx = \dots = [\rho'(x)]^2 \frac{h^3}{12}.$$

- Thus, the total contribution to the MISE from bias² is

$$\int b^2(x) dx \approx \sum_{j=1}^m [\rho'(x_{B_j} + h/2)]^2 \frac{h^3}{12} = \frac{h^2}{12} \sum_{j=1}^m h [\rho'(x_{B_j} + h/2)]^2 \approx \frac{h^2}{12} \int [\rho'(x)]^2 dx.$$

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- In case of the variance, let's assume that h is small, and thus, p_j is close to 0, and therefore, $1 - p_j \approx 1$. Based on that, at a given x falling into bin B_j , the $v(x)$ can be written as

$$v(x) = \frac{p_j(1 - p_j)}{nh^2} \approx \frac{p_j}{nh^2} \approx \frac{h\rho(x) + h\rho'(x) \left[\frac{h}{2} + x_j - x \right]}{nh^2}.$$

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By keeping only the dominant term we obtain

$$v(x) \approx \frac{\rho(x)}{nh}.$$

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- The integral of this over the entire x range yields

$$\int v(x)dx \approx \frac{1}{nh} \int \rho(x)dx = \frac{1}{nh}.$$

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→ The MISE (or risk) can be approximated as

$$R(\widehat{\rho}_n, \rho) \approx \frac{h^2}{12} \int [\rho'(x)]^2 dx + \frac{1}{nh}.$$

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MISE of the histogram estimator

- According to the previous calculations, given a PDF $\rho(x)$ where $\int [\rho'(x)]^2 dx < \infty$, the risk of the corresponding histogram estimator $\widehat{\rho}_n(x)$ with uniform bin width h can be written as

$$R(\widehat{\rho}_n, \rho) \approx \frac{h^2}{12} \int [\rho'(x)]^2 dx + \frac{1}{nh}.$$

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$$R(\widehat{\rho}_n, \rho) \approx \frac{h^2}{12} \int [\rho'(x)]^2 dx + \frac{1}{nh}.$$

- The first term comes from the bias², and is increasing as a function of h , whereas the second term is coming from the variation, and is decreasing as a function of h .

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- The first term comes from the bias², and is increasing as a function of h , whereas the second term is coming from the variation, and is decreasing as a function of h .
- The h^* minimising the risk is

$$h^* = \left(\frac{1}{6n} \int [\rho'(x)]^2 dx \right)^{\frac{1}{3}},$$

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and with this choice

$$R(\widehat{\rho}_n, \rho) \approx \frac{C}{n^{2/3}}, \quad C = \left(\frac{3}{4} \right)^{\frac{2}{3}} \left(\int [\rho'(x)]^2 dx \right)^{\frac{1}{3}}$$

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- Although the decay of $R(\widehat{\rho}_n, \rho)$ as $n^{-2/3}$ is nice from a theoretical point of view, the formula for the optimal bin width h^* is "useless" from a practical point of view, since we have to know the true $\rho(x)$ in order to calculate it...

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- How to choose the optimal h in practice?

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 - How to choose the optimal h in practice?
- Let's write the loss function which we want to minimise as

$$\begin{aligned} R(\widehat{\rho}_n, \rho) &= \int (\widehat{\rho}_n(x) - \rho(x))^2 dx = \\ &\int \widehat{\rho}_n^2(x) dx - 2 \int \widehat{\rho}_n(x) \rho(x) dx + \int \rho^2(x) dx. \end{aligned}$$

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The last term does not depend on the estimator, so we have to minimise only

$$J(h) = \int \widehat{\rho}_n^2(x) dx - 2 \int \widehat{\rho}_n(x) \rho(x) dx,$$

(where the h dependence is implicit in this form through $\widehat{\rho}_n(x)$).

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- To obtain an estimator for $J(h)$, we can use the concept of **cross validation**.

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- To obtain an estimator for $J(h)$, we can use the concept of **cross validation**.
- Probably the most simple case for cross validation in general is the **Jackknife** estimate or Jackknife resampling technique:

- To obtain an estimator for $J(h)$, we can use the concept of **cross validation**.
- Probably the most simple case for cross validation in general is the **Jackknife** estimate or Jackknife resampling technique:

The basic idea is to make n "replicas" of the original sample by always leaving out one data point, and based on these replicas we can get reasonable estimates for the **variance** and **bias** of an estimator of interest in a very simple way. (The name "Jackknife" refers to the simple, all around nature of the technique).

- To obtain an estimator for $J(h)$, we can use the concept of **cross validation**.
- Probably the most simple case for cross validation in general is the **Jackknife** estimate or Jackknife resampling technique:

The basic idea is to make n "replicas" of the original sample by always leaving out one data point, and based on these replicas we can get reasonable estimates for the **variance** and **bias** of an estimator of interest in a very simple way. (The name "Jackknife" refers to the simple, all around nature of the technique).

- As a side track, let us quickly have an overview of Jackknife, and then return to the problem of estimating $J(h)$.

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- Based on our original data $\{x_1, x_2, \dots, x_n\}$ we can define n Jackknife samples by always **leaving out one data point**, thus, the i^{th} Jackknife sample is $X_{[i]} = \{x_1, x_2, \dots, x_{i-1}, x_{i+1}, \dots, x_n\}$.

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- Any statistics or estimator can be evaluated on these samples as if we would on the whole data; let us denote the result of the estimator $s(\cdot)$ of interest on the i^{th} Jackknife sample as

$$\theta_{(i)} = s(X_{[i]}).$$

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- The empirical average of the Jackknife replicas is simply

$$\bar{\theta} = \frac{1}{n} \sum_{i=1}^n \theta_{(i)}.$$

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$$\bar{\theta} = \frac{1}{n} \sum_{i=1}^n \theta_{(i)}.$$

- However, the Jackknife estimate of the variance of $s(\cdot)$ is

$$\mathbb{V}_{\text{jack}}(\theta) = \frac{n-1}{n} \sum_{i=1}^n (\theta_{(i)} - \bar{\theta})^2.$$

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$$\mathbb{V}_{\text{jack}}(\theta) = \frac{n-1}{n} \sum_{i=1}^n (\theta_{(i)} - \bar{\theta})^2.$$

→ Where does the factor $(n-1)/n$ come from?

To illustrate that $(n - 1)/n$ is the correct prefactor, let us consider the case where $s(\cdot)$ is the sample mean, $s(\cdot) = \bar{X}_n = \frac{1}{n} \sum_{i=1}^n x_i$.

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- We know actually, that the variation of \bar{X}_n is simply σ^2/n .

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- We know actually, that the variation of \bar{X}_n is simply σ^2/n .
- Let's consider the inner term in $\mathbb{V}_{\text{jack}}(\theta)$:

$$\theta_{(i)} - \bar{\theta} = \frac{n\bar{X}_n - x_i}{n-1} - \frac{1}{n} \sum_{i=1}^n \bar{X}_{(i)},$$

where $\bar{X}_{(i)}$ denotes the sample mean calculated based on $X_{(i)}$ (leaving out x_i).

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where $\bar{X}_{(i)}$ denotes the sample mean calculated based on $X_{(i)}$ (leaving out x_i). The above expression can be also written as

$$\begin{aligned} \theta_{(i)} - \bar{\theta} &= \frac{1}{n-1} (n\bar{X}_n - x_i) - \frac{1}{n} \sum_{i=1}^n \frac{1}{n-1} \sum_{j=1, j \neq i}^n x_j = \\ &= \frac{1}{n-1} (n\bar{X}_n - x_i) - \frac{1}{n} \sum_{i=1}^n \frac{1}{n-1} (n\bar{X}_n - x_i) = \\ &= \frac{1}{n-1} \left(n\bar{X}_n - x_i - \frac{1}{n} \sum_{i=1}^n (n\bar{X}_n - x_i) \right) = \frac{1}{n-1} (\bar{X}_n - x_i). \end{aligned}$$

- By squaring, summing and applying the prefactor $(n-1)/n$ we obtain

$$\frac{n-1}{n} \sum_{i=1}^n (\theta_{(i)} - \bar{\theta})^2 = \frac{1}{n(n-1)} \sum_{i=1}^n (x_i - \bar{X}_n)^2,$$

which is an unbiased estimator of the variance of the sample mean.

- The bias of a general statistic $s(\cdot)$ can be estimated based on Jackknife as

$$\text{bias}_{\text{jack}}(\theta) = (n - 1) (s(\cdot) - \bar{\theta}) .$$

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- Consequently, the expected value of the average over the Jackknife replicas is

$$\mathbb{E}(\bar{\theta}) = \frac{1}{n} \sum_{i=1}^n \mathbb{E}(\theta_{(i)}) = \theta + \frac{b_1}{n - 1}$$

(since the Jackknife replicas have only $n - 1$ data points).

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(since the Jackknife replicas have only $n - 1$ data points). Based on that, the bias of the Jackknife replicates estimator is

$$\mathbb{E}(s(\cdot) - \bar{\theta}) = \theta + \frac{b_1}{n} - \theta - \frac{b_1}{n - 1} = \frac{b_1}{n(n - 1)}.$$

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$$\mathbb{E}(s(\cdot) - \bar{\theta}) = \theta + \frac{b_1}{n} - \theta - \frac{b_1}{n - 1} = \frac{b_1}{n(n - 1)}.$$

- By multiplying with $(n - 1)$ we obtain that the expected value of $\text{bias}_{\text{jack}}$ is equal to b_1/n , which is the bias of $s(\cdot)$.

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Let us now return to the problem of the empirical PDF, where we would like to obtain an estimate for the loss function

$$J(h) = \int \widehat{\rho}_n^2(x) dx - 2 \int \widehat{\rho}_n(x) \rho(x) dx.$$

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- In the spirit of Jackknife replicas, we can consider the following estimator:

$$\widehat{J}(h) = \int \widehat{\rho}_n^2(x) dx - \frac{2}{n} \sum_{i=1}^n \widehat{\rho}_{(i)}(x = x_i),$$

where $\widehat{\rho}_{(i)}(x = x_i)$ is the histogram estimator obtained by excluding data point i from the sample evaluated at $x = x_i$.

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- The expected value of the second term is

$$\mathbb{E} \left(\frac{2}{n} \sum_{i=1}^n \widehat{\rho}_{(i)}(x_i) \right) = \frac{2}{n} \sum_{i=1}^n \mathbb{E} (\widehat{\rho}_{(i)}(x_i)) \approx 2 \int \widehat{\rho}_n(x) \rho(x) dx,$$

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thus, we obtained a nearly unbiased estimator of the loss function,

$$\mathbb{E}(\widehat{J}(h)) \approx \mathbb{E}(J(h)).$$

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Calculation of $\widehat{J}(h)$:

- At first sight, searching for the optimal h by minimising $\widehat{J}(h)$ seems painful, since at every examined value of h , we have to prepare n Jackknife replicas, calculate $\widehat{\rho}_{(i)}$ for every replica, and then evaluate the sum defining $J(h)$ as

$$\widehat{J}(h) = \int \widehat{\rho}_n^2(x) dx - \frac{2}{n} \sum_{i=1}^n \widehat{\rho}_{(i)}(x = x_i) = \sum_{j=1}^m \widehat{\rho}_n^2(x_j) h - \frac{2}{n} \sum_{i=1}^n \widehat{\rho}_{(i)}(x = x_i),$$

where we used that $\widehat{\rho}_n(x)$ is constant within a given bin.

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- Luckily, there is a faster way, since $\widehat{J}(h)$ can also be formulated simply based on the original \widehat{p}_j of the bins at a given bin width h .

Calculation of $\widehat{\mathcal{J}}(h)$:

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- Luckily, there is a faster way, since $\widehat{\mathcal{J}}(h)$ can also be formulated simply based on the original \widehat{p}_j of the bins at a given bin width h .
- Let us first express the first term in $\widehat{\mathcal{J}}(h)$ based on \widehat{p}_j :

$$\sum_{j=1}^m \widehat{\rho}_n^2(x_j) h = \sum_{j=1}^m \frac{\widehat{p}_j^2}{h^2} h = \sum_{j=1}^m \frac{\widehat{p}_j^2}{h}.$$

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- The second term is a bit more tricky, since the $\widehat{\rho}_{(i)}(x)$ is based on the data obtained by removing x_i .

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- The second term is a bit more tricky, since the $\widehat{\rho}_{(i)}(x)$ is based on the data obtained by removing x_i .
- However, according to the above $\widehat{\rho}_{(i)}(x)$ is taken at $x = x_i$, corresponding to the actual bin where x_i is missing from. Thus,

$$\widehat{\rho}_{(i)}(x = x_i) = \frac{1}{h} \frac{\nu_j - 1}{n - 1} = \frac{1}{h} \frac{n\widehat{p}_j - 1}{n - 1},$$

where we denoted the bin of x_i as j , and expressed the number of elements in the original data in this bin as $\nu_j = n\widehat{p}_j$.

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where we denoted the bin of x_i as j , and expressed the number of elements in the original data in this bin as $\nu_j = n\widehat{p}_j$.

- We can also regroup the summation over the individual data points i to summation over the bins j , taking into account that the number of data point falling into bin j is again $\nu_j = n\widehat{p}_j$:

$$\begin{aligned} -\frac{2}{n} \sum_{i=1}^n \widehat{\rho}_{(i)}(x = x_i) &= -\frac{2}{n} \sum_{j=1}^m \frac{1}{h} \frac{n\widehat{p}_j - 1}{n - 1} n\widehat{p}_j = -\frac{2}{hn} \sum_{j=1}^m \frac{n^2 \widehat{p}_j^2}{n - 1} + \frac{2}{hn} \sum_{j=1}^m \frac{n\widehat{p}_j}{n - 1} = \\ &= -\frac{2n}{h(n-1)} \sum_{j=1}^m \widehat{p}_j^2 + \frac{2}{h(n-1)} \underbrace{\sum_{j=1}^m \widehat{p}_j}_1. \end{aligned}$$

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- Putting it all together we obtain

$$\begin{aligned}\hat{J}(h) &= \sum_{j=1}^m \frac{\hat{p}_j^2}{h} - \frac{2n}{h(n-1)} \sum_{j=1}^m \hat{p}_j^2 + \frac{2}{h(n-1)} = \\ &= \underbrace{\frac{1}{h} \left[1 - \frac{2n}{n-1} \right]}_{\frac{-n-1}{n-1}} \sum_{j=1}^m \hat{p}_j^2 + \frac{2}{h(n-1)} = \\ &= \frac{2}{h(n-1)} - \frac{n+1}{h(n-1)} \sum_{j=1}^m \hat{p}_j^2.\end{aligned}$$

- Putting it all together we obtain

$$\begin{aligned}\widehat{\mathcal{J}}(h) &= \sum_{j=1}^m \frac{\widehat{p}_j^2}{h} - \frac{2n}{h(n-1)} \sum_{j=1}^m \widehat{p}_j^2 + \frac{2}{h(n-1)} = \\ &= \underbrace{\frac{1}{h} \left[1 - \frac{2n}{n-1} \right]}_{\frac{-n-1}{n-1}} \sum_{j=1}^m \widehat{p}_j^2 + \frac{2}{h(n-1)} = \\ &= \frac{2}{h(n-1)} - \frac{n+1}{h(n-1)} \sum_{j=1}^m \widehat{p}_j^2.\end{aligned}$$

- Thus, we do not have to actually generate/evaluate anything related to the individual Jackknife replicas, we can calculate $\widehat{\mathcal{J}}(h)$ for a given h straight away based on simply the original \widehat{p}_j values.

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Optimal histogram based estimator

- For a given h bin width the empirical PDF is constructed the usual way,

$$\widehat{\rho}_n(x) = \sum_{j=1}^m \frac{\widehat{p}_j}{h} I(x \in B_j) = \sum_{j=1}^m \frac{\nu_j}{nh} I(x \in B_j).$$

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- The risk (up to an additive constant independent of h) can be estimated by

$$\widehat{J}(h) = \int \widehat{\rho}_n^2(x) dx - \frac{2}{n} \sum_{i=1}^n \widehat{\rho}_{(i)}(x = x_i).$$

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- By changing the value of h from low to high we locate the optimal h , minimising the above function.
- Luckily, the cross-validation terms do not have to be evaluated individually, since $\hat{J}(h)$ can equally be formulated as

$$\hat{J}(h) = \frac{2}{h(n-1)} - \frac{n+1}{h(n-1)} \sum_{j=1}^m \hat{p}_j^2 = \frac{2}{h(n-1)} - \frac{n+1}{h(n-1)n^2} \sum_{j=1}^m \nu_j^2$$

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- The basic idea of kernel density estimation is to "smudge" the data points, and obtain the estimation of the PDF based on the sum of these.

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- The basic idea of kernel density estimation is to "smudge" the data points, and obtain the estimation of the PDF based on the sum of these.
- A bit more precisely: we represent each data point by a unimodal decaying function (given by the kernel) whose peak is centred on the data point, and the estimate of the PDF at a given x is the sum over these functions.

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Kernel density estimation

- Let the kernel $K(x)$ be a smooth function with the following properties:
 - $K(x) \geq 0$,
 - $\int K(x)dx = 1$,
 - $\int xK(x)dx = 0$, and $\int x^2K(x)dx > 0$.

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(In other words, $K(x)$ can be viewed as the PDF of some probability distribution with 0 mean and a larger than 0 variance).

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- $\int K(x)dx = 1$,
- $\int xK(x)dx = 0$, and $\int x^2K(x)dx > 0$.

(In other words, $K(x)$ can be viewed as the PDF of some probability distribution with 0 mean and a larger than 0 variance).

- Based on $K(x)$, the kernel density estimator of the PDF at a fixed bandwidth h is given by

$$\widehat{\rho}_n(x) = \frac{1}{n} \sum_{i=1}^n \frac{1}{h} K\left(\frac{x - x_i}{h}\right).$$

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Most widely used kernels:

- Gaussian kernel:

$$K(x) = \frac{1}{\sqrt{2\pi}} e^{-\frac{x^2}{2}}.$$

- Epanechnikov kernel:

$$K(x) = \begin{cases} \frac{3}{4\sqrt{5}} \left(1 - \frac{x^2}{5}\right) & \text{if } |x| < \sqrt{5}, \\ 0 & \text{if } |x| \geq \sqrt{5}. \end{cases}$$

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Risk for kernel density estimation

- Under weak assumptions on $\rho(x)$ and $K(x)$, the MISE can be given as

$$R(\widehat{\rho}_n, \rho) \approx \frac{\sigma_K^4 h^4}{4} \int [\rho''(x)]^2 dx + \frac{1}{nh} \int K^2(x) dx,$$

where $\sigma_K^2 = \int x^2 K(x) dx$.

- The optimal bandwidth is

$$h^* = \frac{c_1^{-2/5} c_2^{1/5} c_3^{-1/5}}{n^{1/5}},$$

where $c_1 = \int x^2 K(x) dx$, $c_2 = \int K^2(x) dx$ and $c_3 = \int [\rho''(x)]^2 dx$.

- With this choice of the bandwidth,

$$R(\widehat{\rho}_n, \rho) \approx \frac{c_4}{n^{4/5}},$$

where c_4 is some further constant.

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- According to the above, the empirical PDF based on kernel density estimation is converging faster with n compared to the histogram estimator.
 - However, similarly to the histogram estimator, the previous formulation is not usefully in practise, because it needs the knowledge of the true PDF ρ .
- Luckily, the cross validation approach works here as well.
- The MISE (up to constant independent from the bin width h) can be estimated by

$$\widehat{J}(h) = \int \widehat{\rho}_n(x) dx - \frac{2}{n} \sum_{i=1}^n \widehat{\rho}_{(i)}(x = x_i),$$

where $\widehat{\rho}_{(i)}$ is the kernel density estimator obtained after removing x_i from the data set.

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estimation

- Finally, $\widehat{J}(h)$ based on cross validation can be evaluated at a given h simply as

$$\widehat{J}(h) \approx \frac{1}{hn^2} \sum_{i=1}^n \sum_{j=1}^n K^* \left(\frac{x_i - x_j}{h} \right) + \frac{2}{nh} K(0),$$

where $K^*(x) = \int K(x-y)K(y)dy - 2K(x)$.