Week 5: Cross-validation

MATH-517 Statistical Computation and Visualization

Linda Mhalla

2023-10-20

Motivation

Over the last two lectures, we've covered **KDE** and **non-parametric regression methods**

Both required the choice of a certain tuning parameter

• KDE, h > 0

$$\hat{f}(x) = \frac{1}{nh} \sum_{i=1}^{n} K\left(\frac{X_i - x}{h}\right)$$

• Local Polynomial Regression (with a fixed degree p), h > 0

$$\arg\min\nolimits_{\beta\in\mathbb{R}^{p+1}}\sum_{i=1}^{n}\{Y_{i}-\beta_{0}-\beta_{1}(X_{i}-x)-...-\beta_{p}(X_{i}-x)^{p}\}^{2}K\left(\frac{X_{i}-x}{h}\right)$$

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Motivation

Many other modern methods for regression can be expressed as

penalized regression

$$\label{eq:argmin} \arg\min_{\boldsymbol{\beta}} \sum_{n=1}^{N} \left(y_n - \boldsymbol{x}_n^{\intercal} \boldsymbol{\beta} \right)^2 + \textcolor{red}{\lambda} R(\boldsymbol{\beta}),$$

where R is a penalty, e.g., $\|\cdot\|_2^2$ for ridge regression or $\|\cdot\|_1$ for lasso, or

smoothing splines

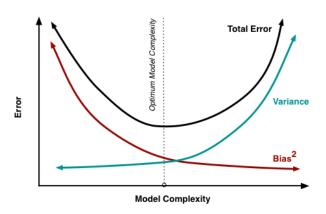
$$\arg\min_{\beta} \sum_{n=1}^N \{y_n - f(x_n)\}^2 + \frac{\lambda}{\lambda} \int \{f''(x)\}^2 dx$$

In all cases, **cross-validation (CV)** can be used to select the tuning parameters

not always straightforward!

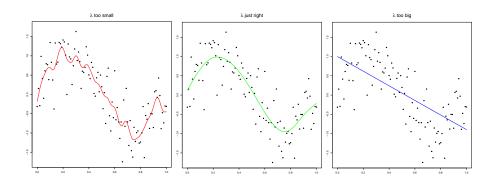
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Bias-variance Trade-off



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Bias-variance Trade-off: Smoothing splines



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Purpose of Cross-validation

Cross-validation (CV) is a very general method for

- tuning the regularization parameter of a method
- estimating the predictive power of a method

Since training an algorithm and evaluating its performance on the same data yields an overoptimistic result, CV fixes the issue by testing the output of a method on (independent) "new data"

CV involves

- splitting a data set into a training data set and a test data set
- fitting the model using the training data set
- using the test data set to evaluate how the model performs (according to a measure of error/risk)
- computing the average over several splits (several splitting strategies exist!)

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Section 1

CV for Supervised Problems

Local Polynomial Regression

 $\begin{array}{l} \textbf{Setup} \colon \mathsf{A} \ \mathsf{sample} \ (x_1,y_1)^\top, \dots, (x_N,y_N)^\top \in \mathbb{R}^2 \ \mathsf{from \ a \ population} \\ Y = m(X) + \epsilon \ \mathsf{with} \ X \perp \!\!\! \perp \epsilon. \ \mathsf{For \ a \ fixed \ bandwidth} \ h, \ \mathsf{we \ estimate} \\ m(x) = \mathbb{E}(Y|X=x) \ \mathsf{as} \ \widehat{m}_h(x) \ \mathsf{by, \ e.g., \ local \ linear \ regression}. \end{array}$

Question: How to choose h? (i.e., how to obtain a good bias-variance trade-off?)

What is the measure of how good our estimator $\widehat{m}_h(x)$ for a given bandwidth is?

$$MISE(\widehat{m}_h) = \int \mathbb{E}\big\{\widehat{m}_h(x) - m(x)\big\}^2 f_X(x) dx,$$

ullet let's choose h that minimizes the (density-weighted) MISE. Here, what matters is to minimize the estimation error on the regions where the density of X is higher

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Local Polynomial Regression

But we don't know m. How about using the average RSS

$$\frac{1}{N} \sum_{n=1}^{N} \left\{ Y_n - \widehat{m}_h(X_n) \right\}^2.$$

as a proxy for the MISE?

That's a bad idea, because $\left\{Y_n-\widehat{m}_h(X_n)\right\}^2 \to 0$ for $h\to 0$

- this is called *overfitting* (useless interpolation)
- the problem lies in validating on data used to fit the model (favours estimates too well-adapted to data and unreasonable for new obs.)

Instead, consider this to approximate MISE:

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where $\widehat{m}_h^{(-n)}(X_n)$ is the model fitted without the n-th observation

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CV for Local Polynomial Regression

$$CV(h) = \frac{1}{N} \sum_{n=1}^{N} \left\{ Y_n - \widehat{m}_h^{(-n)}(X_n) \right\}^2$$

Since $Y = m(X) + \epsilon$, we can write

$$\begin{split} CV(h) &= \frac{1}{N} \sum_{n=1}^{N} \left\{ Y_n - m(X_n) + m(X_n) - \widehat{m}_h^{(-n)}(X_n) \right\}^2 \\ &= \frac{1}{N} \sum_{n=1}^{N} \epsilon_n^2 + \frac{2}{N} \sum_{n=1}^{N} \epsilon_n \left\{ m(X_n) - \widehat{m}_h^{(-n)}(X_n) \right\} \\ &+ \underbrace{\frac{1}{N} \sum_{n=1}^{N} \left\{ m(X_n) - \widehat{m}_h^{(-n)}(X_n) \right\}^2}_{\mathbb{E} \star = MISE\left(\widehat{m}_h\right)}, \end{split}$$

$$MISE(\widehat{m}_h) = \mathbb{E} \int \left\{ \widehat{m}_h(x) - m(x) \right\}^2 f_X(x) dx$$

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CV can be Easy for Prediction

More generally: $(x_1,y_1)^\top,\dots,(x_N,y_N)^\top\in\mathbb{R}^{p+1}$

Model for prediction: $\widehat{Y} = \widehat{m}(X)$

How good is the model: measured by a loss function, e.g., $\mathbb{E}\big\{Y-\widehat{m}(X)\big\}^2$

• other losses possible, e.g., if undershooting better than overshooting

If another data set $(x_1^\star,y_1^\star)^\top,\ldots,(x_M^\star,y_M^\star)^\top$ available (generated by the same process as the original data set), we can approximate loss empirically

$$\frac{1}{M} \sum_{k=1}^{M} \{y_k^{\star} - \widehat{m}(x_k^{\star})\}^2$$

CV is the alternative when no other data set is available:

$$CV(\widehat{m}) := \frac{1}{N} \sum_{n=1}^N \left\{ y_n - \widehat{m}^{(-n)}(x_n) \right\}^2,$$

where $\widehat{m}^{(-n)}$ is the model fitted without the n-th observation

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CV can be Easy for Prediction

It can often be shown (under assumptions!) like in the case of local polynomial regression that

$$CV(\widehat{m}) \to \mathbb{E}\big\{Y - \widehat{m}(X)\big\}^2$$

CV can be used to compare candidate models $\widehat{m}_1, \dots, \widehat{m}_j$

- can be completely different models
 - typically it is the same model with different tuning parameter values
- select the model for which the CV criterion is minimized
- beware: when not in the "vanilla" iid case (e.g. times series, stratified data, etc.), things are not so straightforward...

But there are computational costs. The model has to be re-fitted for

- all the tuning parameter values considered
- every data point left out
 - actually, this might not be necessary...

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Computational Shortcut for Linear Smoothers

If \widehat{m} is a linear smoother, i.e., the predictions $\widehat{y}_n=\widehat{m}(x_n)$ are given all together as

$$\hat{\mathbf{y}} = \mathbf{S}\mathbf{y}$$

where $\mathbf{S} \in \mathbb{R}^{N \times N}$ depends on x's, then re-fitting (leaving out data points one by one) may not be necessary!

Example: Ridge regression is a linear smoother

$$\arg\min \ \sum_{n=1}^N \left(y_n - x_n^\top \boldsymbol{\beta}\right)^2 + \lambda \|\boldsymbol{\beta}\|_2^2.$$

$$\bullet \ \hat{\beta} = (\mathbf{X}^{\top}\mathbf{X} + \lambda I)^{-1}\mathbf{X}^{\top}\mathbf{y}$$

$$\bullet \ \hat{\mathbf{y}} = \underbrace{\mathbf{X}(\mathbf{X}^{\top}\mathbf{X} + \lambda I)^{-1}\mathbf{X}^{\top}}_{=:\mathbf{S}} \mathbf{y}$$

$$CV(\lambda) = \frac{1}{N} \sum_{n=1}^N \left\{ y_n - \mathbf{x}_n^\top \widehat{\beta}^{(-n)} \right\}^2 = \frac{1}{N} \sum_{n=1}^N \left\{ \frac{y_n - \widehat{m}(x_n)}{1 - s_{nn}} \right\}^2$$

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Example: Ridge Regression

Noticing $\hat{\beta}^{(-n)} = (\mathbf{X}^{\top}\mathbf{X} + \lambda I - \mathbf{x}_n\mathbf{x}_n^{\top})^{-1}(\mathbf{X}^{\top}\mathbf{y} - \mathbf{x}_ny_n)$, we can use Sherman-Morrison formula:

- denoting $\mathbf{A} := \mathbf{X}^{\top} \mathbf{X} + \lambda I$
- $\bullet \ \alpha_n := 1 \mathbf{x}_n^{\top} \mathbf{A}^{-1} \mathbf{x}_n$

$$\begin{split} \hat{\beta}^{(-n)} &= \left(\mathbf{A}^{-1} - \frac{\mathbf{A}^{-1}\mathbf{x}_n\mathbf{x}_n^{\intercal}\mathbf{A}^{-1}}{1 - \mathbf{x}_n^{\intercal}\mathbf{A}^{-1}\mathbf{x}_n}\right) (\mathbf{X}^{\intercal}\mathbf{y} - \mathbf{x}_n y_n) \\ &= \hat{\beta} - \frac{1}{\alpha_n} (\mathbf{A}^{-1}\mathbf{x}_n\mathbf{x}_n^{\intercal}\hat{\beta} - \mathbf{A}^{-1}\mathbf{x}_n y_n). \end{split}$$

Plug this back into the general CV formula and do some simple algebra to obtain the last formula on the previous slide

• check out lecture notes for details, if interested

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Computational Shortcut for Linear Smoothers

A similar computational shortcut is possible for

- linear models
- local constant regression
 - what about other polynomial orders?
- ridge regression
- KDE (when working on a grid)

On the other hand, such shortcuts are not possible for

- lasso
- many other penalized or otherwise complicated estimators

When a computational shortcut is impossible, perform K-fold CV instead!

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K-fold CV

Divide the set $\{1,\dots,N\}$ into K subsets (folds) of approximately equal size, J_1,\dots,J_K , such that

• fold $J_k\subset\{1,\dots,N\}$ for $k=1,\dots,K$ such that $J_k\cap J_{k'}=\emptyset$ for $k\neq k'$ and $\bigcup_{k=1}^K J_k=\{1,\dots,N\}$

For k = 1, ..., K:

- \bullet Consider training on (x_i,y_i) , $i\not\in J_k$, and validating on (x_i,y_i) , $i\in J_k$
- Fit the model on the training set and compute the error on the validation set

$$e_k = \sum_{n \in J_k} \{y_n - \widehat{m}^{(-J_k)}(x_n)\}^2$$

where $m^{\left(-J_{k}\right)}$ is the model fitted without the data in the k-th fold J_{k}

Compute the average error over all folds

In practice, choose K=5 or K=10, perform random permutation of indices and split the data

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Instead of the (leave-one-out) CV criterion

$$CV(\widehat{m}) := \frac{1}{N} \sum_{n=1}^N \left\{ y_n - \widehat{m}^{(-n)}(x_n) \right\}^2,$$

use the K-fold CV criterion:

$$CV_K(\widehat{m}) = K^{-1} \sum_{k=1}^K |J_k|^{-1} \sum_{n \in J_k} \big\{ Y_n - \widehat{m}^{(-J_k)}(X_n) \big\}^2.$$

- ullet requires every candidate model to be fitted K-times
- ullet it is difficult to study properties of $CV_K(\widehat{m})$ properly. One usually examines whether leave-one-out CV works and, if yes and if no computational shortcuts available, resorts to K-fold CV for computational reasons

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Section 2

CV for Unsupervised Problems

Sample X_1, \dots, X_N from f, goal is to estimate f(x) by

$$\hat{f}_h(x) = \frac{1}{nh} \sum_{i=1}^N K\left(\frac{X_i - x}{h}\right)$$

• no response here!

A good estimator (a well-chosen h) minimizes

$$\begin{split} MISE(\hat{f}_h) &= \mathbb{E} \int \big\{ \hat{f}_h(x) - f(x) \big\}^2 dx \\ &= \mathbb{E} \underbrace{\int \big\{ \hat{f}_h(x) \big\}^2 dx}_{\|\hat{f}_h(x)\|_2^2} - 2 \underbrace{\mathbb{E} \int \hat{f}_h(x) f(x) dx}_{A(h): \text{ the CV idea?}} + \underbrace{\int \big\{ f(x) \big\}^2 dx}_{\text{no h here}}. \end{split}$$

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Sample X_1,\dots,X_N from f, goal is to estimate f(x) by

$$\hat{f}_h(x) = \frac{1}{nh} \sum_{i=1}^N K\left(\frac{X_i - x}{h}\right)$$

no response here!

A good estimator (a well-chosen h) minimizes

$$\begin{split} MISE(\hat{f}_h) &= \mathbb{E} \int \big\{ \hat{f}_h(x) - f(x) \big\}^2 dx \\ &= \mathbb{E} \underbrace{\int \big\{ \hat{f}_h(x) \big\}^2 dx}_{\|\hat{f}_h(x)\|_2^2} - 2 \underbrace{\mathbb{E} \int \hat{f}_h(x) f(x) dx}_{A(h): \text{ the CV idea?}} + \underbrace{\int \big\{ f(x) \big\}^2 dx}_{\text{no h here}}. \end{split}$$

Let's find an unbiased estimator of A(h)!

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The CV idea: see how your estimator behaves on a left-out datum:

$$\begin{split} \mathbb{E}\widehat{f}_h^{(-n)}(X_n) &= \mathbb{E}\frac{1}{(n-1)h}\sum_{j\neq n}K\left(\frac{X_n-X_j}{h}\right) = \frac{1}{h}\mathbb{E}K\left(\frac{X_1-X_2}{h}\right) \\ &= \int \underbrace{\int \frac{1}{h}K\left(\frac{x-y}{h}\right)f(y)dy}_{\mathbb{E}\widehat{f}_h(x)}f(x)dx = \mathbb{E}\int \widehat{f}_h(x)f(x)dx. \end{split}$$

$$\Rightarrow N^{-1} \sum_{n=1}^N \hat{f}_h^{(-n)}(X_n)$$
 is an unbiased estimator of $\mathbb{E} \int \hat{f}_h(x) f(x) dx$

Thus, up to the constant (not depending on h), an unbiased estimator of

$$MISE(\hat{f}_h) = \mathbb{E} \int \left[\hat{f}_h(x)\right]^2 \! dx - 2\mathbb{E} \int \hat{f}_h(x) f(x) dx + \int \left[f(x)\right]^2 \! dx.$$

is given by the CV

$$CV(h) = \int \big[\hat{f}_h(x)\big]^2 dx - \frac{2}{N} \sum_{n=1}^N \hat{f}_h^{(-n)}(X_n)$$

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The computational formula for CV(h) is given by

$$\frac{1}{n^2 h} \sum_{i=1}^n \sum_{j=1}^n \int K(y) K \bigg(\frac{X_i - X_j}{h} - y \bigg) dy - \frac{2}{n(n-1)h} \sum_{j=1}^n \sum_{i \neq j} K \bigg(\frac{X_i - x_j}{h} \bigg)$$

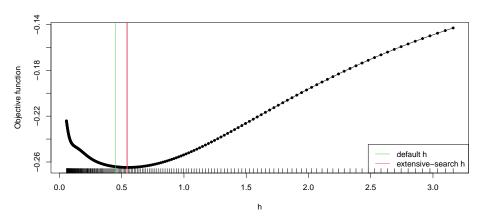
The optimal bandwidth is then given by

$$h_{opt} = \operatorname*{arg\,min}_{h>0} \, CV(h)$$

- Numerical optimisation is required
- \bullet The roughness of the objective function depends on n and $f\Rightarrow$ might have several local minima

 \Rightarrow Always check the solution by plotting CV(h) for a range of h

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CV for PCA

(1) linear combinations with maximal variance (Pearson, 1901)

 $\underset{v^\top v=1}{\operatorname{arg\,max}} \ v^\top \widehat{\Sigma} v$

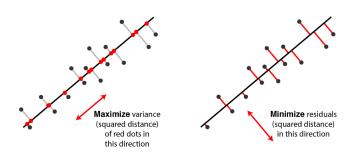
(2) minimum least square error projection into lower dimension (Hotelling, 1933)

 $\underset{V^{\top}V = I_r}{\arg\min} \ \sum_{i=1}^n \|x_i - \mathbf{V}\mathbf{V}^{\top}x_i\|_2^2$

(3) best low-rank matrix approximation (Eckart & Young, 1936)

 $\mathop{\arg\min}_{\mathop{\mathrm{rank}}(\mathbf{L})=r}\|\mathbf{X}-\mathbf{L}\|_2^2$

(1)-(2) Optimisation problems



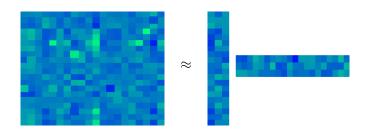
Two equivalent views of principal component analysis.

Source: this blog

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(3) Low-rank Matrix Approximation

Visualization for r=3:



$$\mathbf{X} pprox \mathbf{L} = \mathbf{A} \mathbf{V}^ op = \sum_{i=1}^r \mathbf{a}_i \mathbf{v}_i^ op$$

The tall and skinny matrix A and the short and fat matrix V are obtained by truncating the \mathbf{SVD} decomposition: $\mathbf{X} = \mathbf{UDV}^{\top}$ to keep the r top singular values of X

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CV for PCA

In all formulations, there is a hyperparameter r(< p)!

Let's focus on the third formulation of PCA

$$\operatorname*{arg\,min}_{\mathrm{rank}(\mathbf{L})=r} \|\mathbf{X} - \mathbf{L}\|_2^2$$

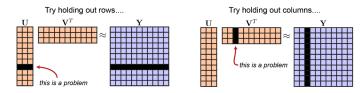
How to choose the rank r? Many people try the following K-fold CV scheme:

- split data into K folds J_1, \dots, J_K
- for k = 1, ..., K
 - ullet solve $\widehat{\mathbf{L}} = rg \min_{\mathrm{rank}(\mathbf{L}) = r} \|\mathbf{X}[J_k^c,] \mathbf{L}\|_2^2$
 - calculate $Err_k(r) = \sum_{n \in L} \|x_n P_{\widehat{L}}x_n\|_2^2$
- end for
- \bullet choose $\hat{r} = \underset{r}{\arg\min} \ \sum_{k=1}^{K} |J_k|^{-1} Err_k(r)$

But this is wrong! (as $r\nearrow$ we have $\|x_j-P_{\widehat{L}}x_j\|\searrow$, so r is overestimated)

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CV for PCA



Not so great ideas for cross-validating matrix factorization.

Source: this blog

Problems with holding out a whole column (or row) of the data matrix are discussed in more detail by Bro et al. (2008) and Owen & Perry (2009)

There are smarter holdout patterns

- Wold hold-out: requires an SVD decomposition with missing data as entries are held-out at random
- Gabriel hold-out: transforms the unsupervised learning problem into a supervised one by holding-out a block of the data matrix

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Intermezzo: Linear Prediction for Gaussian Vectors

For $X \sim \mathcal{N}(\mu, \Sigma)$ split into

$$X = \begin{pmatrix} X_1 \\ X_2 \end{pmatrix}, \qquad \mu = \begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix}, \qquad \Sigma = \begin{pmatrix} \Sigma_{11} & \Sigma_{12} \\ \Sigma_{12} & \Sigma_{22} \end{pmatrix},$$

the conditional expectation of X_1 given X_2 is given by

$$\mathbb{E}_{\mu,\Sigma}\big[X_1 \big| X_2 = \mathbf{x}_2\big] = \mu_1 + \Sigma_{12} \Sigma_{22}^{-1} (\mathbf{x}_2 - \mu_2)$$

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Intermezzo: Linear Prediction for Gaussian Vectors

For $X \sim \mathcal{N}(\mu, \Sigma)$ split into

$$X = \begin{pmatrix} X_1 \\ X_2 \end{pmatrix}, \qquad \mu = \begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix}, \qquad \Sigma = \begin{pmatrix} \Sigma_{11} & \Sigma_{12} \\ \Sigma_{12} & \Sigma_{22} \end{pmatrix},$$

the conditional expectation of X_1 given X_2 is given by

$$\mathbb{E}_{\mu,\Sigma}\big[X_1\big|X_2=\mathbf{x}_2\big] = \mu_1 + \Sigma_{12}\Sigma_{22}^{-1}(\mathbf{x}_2-\mu_2)$$

Assume we have a sample X_1,\dots,X_N from which we obtain estimators $\hat{\mu}$ and $\widehat{\Sigma}$, and a new incomplete observation $X^\star=(X_1^\star,X_2^\star)^\top$, where only X_2^\star is observed. We simply predict the missing part by

$$\widehat{X}_1^\star = \widehat{\mu}_1 + \widehat{\Sigma}_{12} \widehat{\Sigma}_{22}^{-1} (\mathbf{x}_2 - \mu_2)$$

Even without Gaussianity, this is the best linear unbiased predictor (BLUP)

 \bullet The quality of BLUP depends on that of the estimators $\hat{\mu}$ and $\widehat{\Sigma}$

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CV for PCA Repaired

Assume that data $\mathbf{x}_n \in \mathbb{R}^p$ are i.i.d. realizations of $X \sim \mathcal{N}(\mu, \Sigma).$

- \bullet split data into K folds J_1,\dots,J_K
- for $k=1,\ldots,K$
 - estimate μ and Σ empirically using all but the k-th fold J_k , but truncate Σ to be rank-r
 - for $n \in J_k$
 - split ${f x}_n$ into a "missing" part ${f x}^{miss}$ that will be used for validation and an "observed" part ${f x}^{obs}$
 - \bullet predict \mathbf{x}_n^{miss} from \mathbf{x}_n^{obs} as discussed on the previous slide
 - end for
 - calculate $Err_k(r) = \sum_{n \in J_r} \|(\mathbf{x}_n^{obs}, \mathbf{x}_n^{miss})^\top (\mathbf{x}_n^{obs}, \hat{\mathbf{x}}_n^{miss})^\top \|_2^2$
- end for
- \bullet choose $\hat{r} = \mathop{\arg\min}_{r} \; \sum_{k=1}^{K} |J_k|^{-1} Err_k(r)$

Is there a bias-variance trade-off now?

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Assignment 4 [5 %]

Go to Assignment 4 for details.