PSY9511: Seminar 4

Testing, resampling, and splitting

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Outline

- 1. Coding tips
 - · Loops
 - Functions
- 2. Performance metrics
- 3. Strategies for model evaluation
 - · Training and validation split
 - (Stratification)
 - (Leave-one-out cross-validation)
 - · Cross-validation
 - Bootstrap
 - Model comparison
- 4. Strategies for model selection and evaluation
 - Train/validation/test split
 - · Nested cross-validation



Coding tips



Coding tips

```
In[1]:
    import numpy as np
    import pandas as pd

df = pd.read_csv('Auto.csv')
    df = df.replace('?', np.nan)
    train = df.iloc(:200].copy()
    test = df.iloc(:300:).copy()

test['cylinders'] = (test['cylinders'] - train['cylinders'].mean()) / train['cylinders'].std()
    train['cylinders'] = (train['cylinders'] - train['cylinders'].mean()) / train['cylinders'].std()
    test['weight'] = (test['weight'] - train['weight'].mean()) / train['weight'].std()
    train['weight'] = (train['weight'] - train['weight'].mean()) / train['weight'].std()
    test['year'] = (train['year'] - train['year'].mean()) / train['year'].std()
    train['year'] = (train['year'] - train['year'].mean()) / train['year'].std()
```



Coding tips: Live coding

Live coding



Coding tips: Python

return train, test

for column in ['cylinders', 'displacement', 'weight']: train, test = standardize(train, test, column=column)

In[1]:

```
import numpy as np
         import pandas as pd
         df = pd.read csv('Auto.csv')
         df = df.replace('?', np.nan)
         train = df.iloc[:200].copy()
         test = df.iloc[300:].copv()
         test['cylinders'] = (test['cylinders'] - train['cylinders'].mean()) / train['cylinders'].std()
         train['cvlinders'] = (train['cvlinders'] - train['cvlinders'].mean()) / train['cvlinders'].std()
         test['weight'] = (test['weight'] - train['weight'].mean()) / train['weight'].std()
         train['weight'] = (train['weight'] - train['weight'].mean()) / train['weight'].std()
         test['vear'] = (test['vear'] - train['vear'].mean()) / train['vear'].std()
         train['year'] = (train['year'] - train['year'].mean()) / train['year'].std()
In[2]:
         import numpy as np
         import pandas as pd
         df = pd.read csv('Auto.csv')
         df = df.replace('?', np.nan)
         train = df.iloc[:200].copy()
         test = df.iloc[300:].copv()
         def standardize(train: pd.DataFrame, test: pd.DataFrame, column: str):
             train = train.copy()
             test = test.copv()
             test[column] = (test[column] - train[column].mean()) / train[column].std()
```



Coding tips: R

```
data <- read.csv('Auto.csv')
data[] <- lapply(data, function(x) replace(x, x == '?', NA))

train <- data[1:200,]
test <- data[2e0:nrow(data),]

test$cylinders <- (testain$cylinders - mean(train$cylinders)) / sd(train$cylinders)
train$cylinders <- (testain$cylinders - mean(train$cylinders)) / sd(train$cylinders)
test$weight <- (test$weight - mean(train$weight)) / sd(train$weight)
train$weight <- (train$weight - mean(train$weight)) / sd(train$weight)
train$yeight <- (train$veight - mean(train$year)) / sd(train$veight)
train$year <- (train$veight - mean(train$year)) / sd(train$year)</pre>
```

```
data <- read.csv('-/Downloads/Auto.csv')
data[] <- lapply(data, function(x) replace(x, x == '?', NA))

train <- data[1:200,]
test <- data[200:nrow(data),]

standardize <- function(train, test, column)
train <- copy(train)
test <- copy(test)

test[,column] <- (test[column] - mean(train[,column])) / sd(train[,column])
train(,column] <- (train[column] - mean(train[,column])) / sd(train[,column]))
return(list(train=train, test=test))

for (column in c('cylinders', 'weight', 'year'))
result <- standardize(train, test, column)
train <- result$train
test <- result$test</pre>
```



Coding tips: Minimal, complete scripts

Ctrl+Shift+Enter



Performance metrics



$$\frac{1}{n} \sum_{i=0}^{n} (y_i - \hat{y}_i)^2$$



$$\frac{1}{n} \sum_{i=0}^{n} (y_i - \hat{y}_i)^2$$

Mean squared error (MSE)

- + Widely used
- + Intuitive
- + Penalizes large errors
- ? Interpretation
- Depends on scale



$$\sqrt{\frac{1}{n}\sum_{i=0}^{n}(y_i-\hat{y}_i)^2}$$



$$\sqrt{\frac{1}{n}\sum_{i=0}^{n}(y_i-\hat{y}_i)^2}$$

Root mean squared error (RMSE)

- + Intuitive
- + Penalizes large errors
- + More interpretable than MSE, total loss ≈ individual loss
- Depends on scale



$$\frac{1}{n}\sum_{i=0}^{n}|y_i-\hat{y}_i|$$

$$\frac{1}{n}\sum_{i=0}^{n}|y_i-\hat{y}_i|$$

Mean absolute error (MAE)

- + More interpretable than MSE/RMSE, total loss = average error
- Feels a bit off
- Depends on scale



$$\frac{\sum_{i=1}^{n} (y_i - \bar{y})(\hat{y}_i - \bar{\hat{y}})}{\sqrt{\sum_{i=1}^{n} (y_i - \bar{y})^2 \sum_{i=1}^{n} (\hat{y}_i - \bar{\hat{y}})^2}}$$



$$\frac{\sum\limits_{i=1}^{n}(y_{i}-\bar{y})(\hat{y}_{i}-\bar{\hat{y}})}{\sqrt{\sum\limits_{i=1}^{n}(y_{i}-\bar{y})^{2}\sum\limits_{i=1}^{n}(\hat{y}_{i}-\bar{\hat{y}})^{2}}}$$

Pearson correlation coefficient (r)

- + Scale independent
- Captures linear correlation
- Does not care about whether the predictions are close to the true values

$$1 - \frac{\sum\limits_{i=1}^{n} (y_i - \hat{y}_i)^2}{\sum\limits_{i=1}^{n} (y_i - \bar{y}_i)^2}$$



$$1 - \frac{\sum_{i=1}^{n} (y_i - \hat{y}_i)^2}{\sum_{i=1}^{n} (y_i - \bar{y}_i)^2}$$

Proportion of variance explained (r^2)

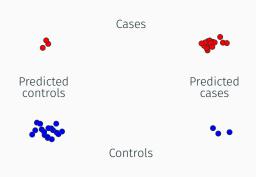
- + Scale independent
- + Interpretable
- Captures linear correlation
- Does not care about whether the predictions are close to the true values



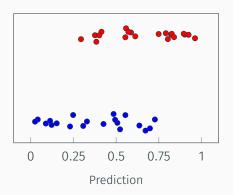


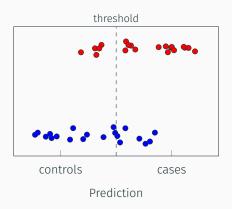


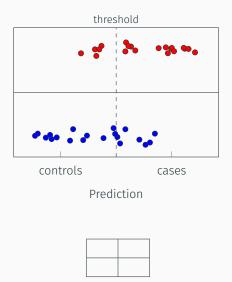


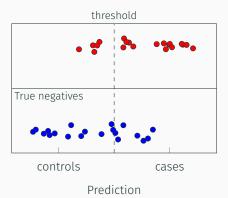




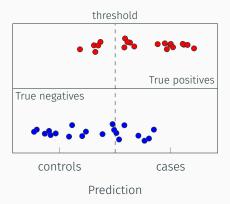




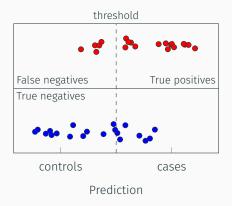




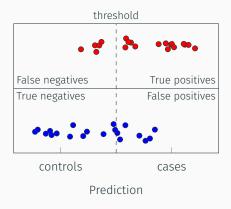
TN



TN	
	TP







TN	FP
FN	TP

$$\frac{\mathit{TP} + \mathit{TN}}{\mathit{TP} + \mathit{TN} + \mathit{FP} + \mathit{FN}}$$



$$\frac{TP+TN}{TP+TN+FP+FN}$$

Accuracy

- + Interpretable
- Does not account for imbalanced classes
- Does not account for different costs of misclassification

$$\frac{TP}{TP+FN}$$



$$\frac{TP}{TP+FN}$$

True positive rate (sensitivity)

- + Interpretable, calculates the proportion of cases that are detected
- + Useful when the cost of false negatives is high (Population-wide screening for severe disease)







$$\frac{TN}{TN+FP}$$

True negative rate (specificity)

- + Interpretable, calculates the proportion of controls that are detected
- Useful when the cost of false positives is high (Intrusive treatment of rare and mild conditions)



$$\frac{TP}{TP+FP}$$



$$\frac{TP}{TP+FP}$$

Positive predictive value (PPV, precision)

- + Interpretable, calculates the proportion of predicted cases that are actually cases
- Useful when the cost of false positives is high (Selection of participants for expensive clinical trials)

$$\frac{\frac{TP}{TP+FN} + \frac{TN}{TN+FP}}{2}$$

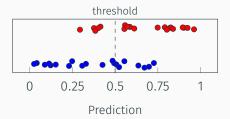


$$\frac{\frac{TP}{TP+FN} + \frac{TN}{TN+FP}}{2}$$

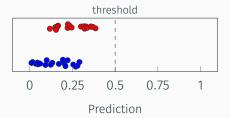
Balanced accuracy

- + Interpretable, behaves similarly to regular accuracy.
- + Takes into account imbalanced classes

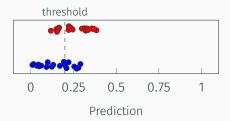




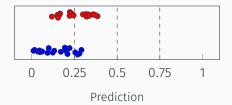


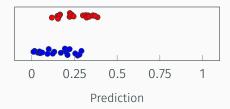




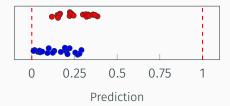




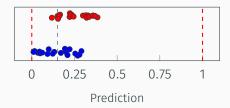




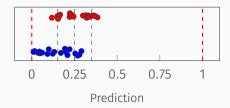
threshold	TPR	FPR



threshold	TPR	FPR
0	1	1
1	0	0

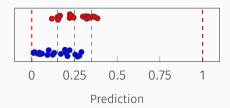


threshold	TPR	FPR
0	1	1
0.15	0.95	0.5
1	0	0

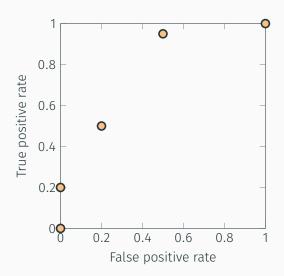


threshold	TPR	FPR
0	1	1
0.15	0.95	0.5
0.25	0.5	0.2
0.35	0.2	0.0
1	0	0

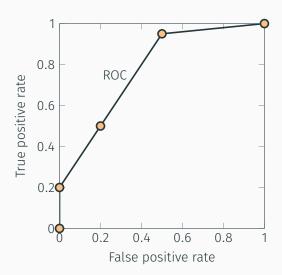




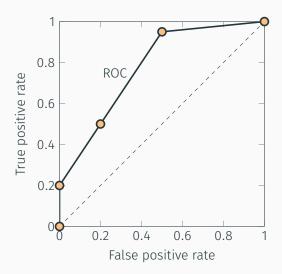
threshold	TPR	FPR
0	1	1
0.15	0.95	0.5
0.25	0.5	0.2
0.35	0.2	0.0
1	0	0



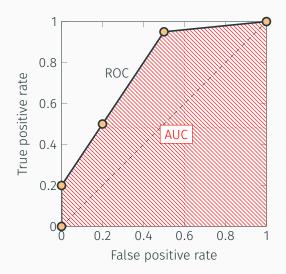




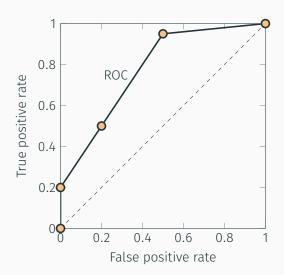




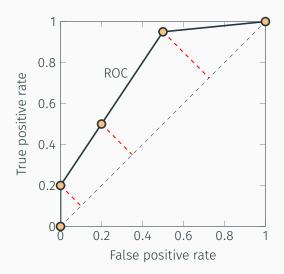


















Performance metrics: Summary

- There is a range of metrics that can be used, each capturing a different aspect of a model's performance
- If possible, (it is my personal preference to) evaluate a model using a different metric than the one that was used for training
- It is good practice to report more than one metric
- For regression, MAE provides a good, intuitive summary of model performance
- For classification, AUC is a widely used metric that is easy to interpret, handles class imbalance (to some degree), and is not reliant on the choice of classification threshold



Strategies for model evaluation



Model evaluation: Rationale

Statistical inference:

Goal: In-sample quantification

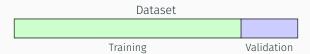
Predictive modelling:

Goal: Out-of-sample generalization

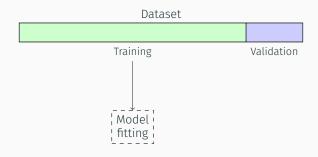




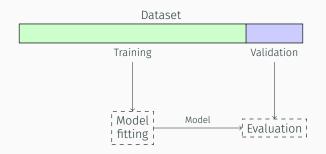




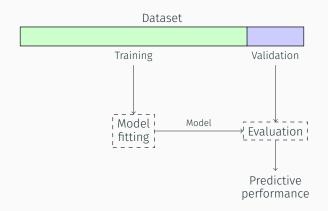










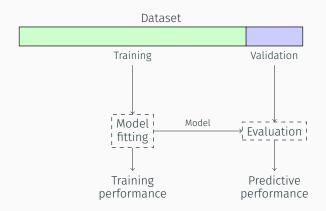




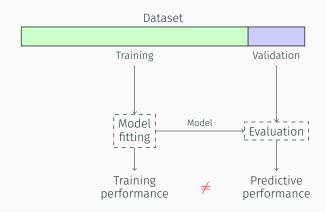
In the validation set approach we split the dataset into two subsets (commonly $\sim 80\%/20\%$), use the first for training the model and the second to test its performance.

- + Accurate estimate of out-of-sample error
- + Simple
- Variable results depending on the exact split
- Only uses a subset of data for training models
- Gives a point estimate of the error, without confidence intervals

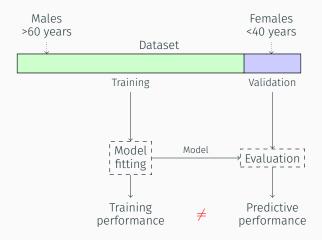














Stratification:

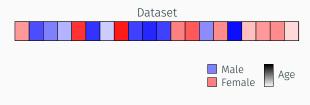
Ensuring all folds of the dataset are similar with respect to some given characteristics.



Dataset

```
In[1]: df = ...
```



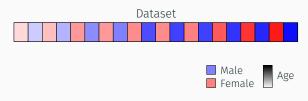


```
In[1]: df = ...
```



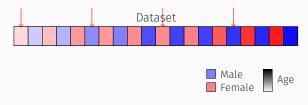


```
In[1]: df = ...
    train = df.iloc[:int(len(df) * 0.8)]
    validation = df.iloc[int(len(df) * 0.8):]
```



```
In[1]: df = ...
    df = df.sort_values(['sex', 'age'])
```



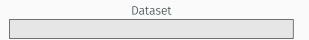


Model evaluation: Stratification

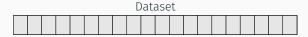
Stratification:

Ensuring all folds of the dataset are similar with respect to some given characteristics.

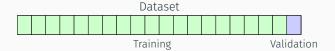
- Helps alleviate the risk of training performance >> validation performance
- · Always stratify on target variable first
- Also good idea to stratify on other core characteristics, e.g. sex and age



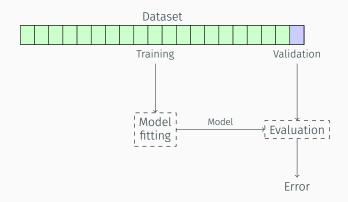




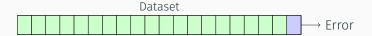




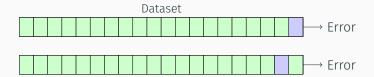


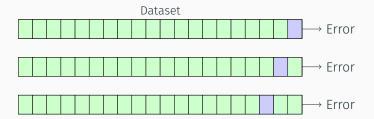








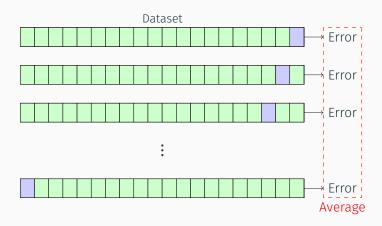










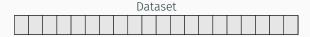




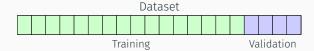
Fits *n* models for *n* datapoints, each leaving a single datapoint out for testing.

- + Uses all data to train models
- + Not dependent on arbitrary data splits
- + Unbiased (with regards to the full dataset)
- Computationally expensive
- Effectively gives a point estimate of the error
- All models are going to be trained on > 99% overlapping data
 - \rightarrow highly correlated

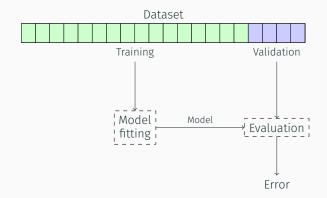




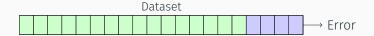




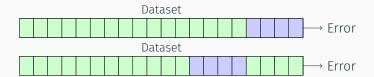




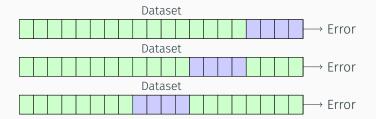




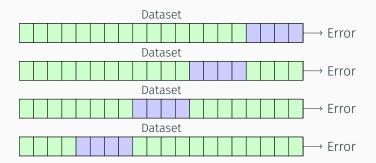




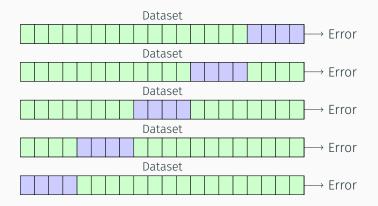




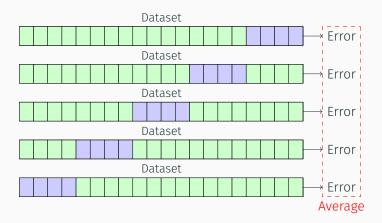










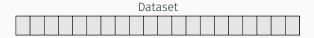


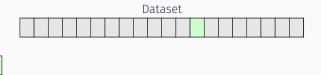


Fits k (usually $k \in \{5, 10\}$) models for n > k datapoints, each leaving n/k datapoints for out-of-sample testing.

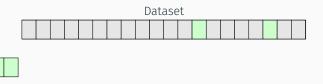
- + Uses all data to train models
- + Yields multiple estimates of out-of-sample error
- Different choices of k (and exact splits) yields different results
- No longer a single model from which information (e.g. parameter estimates and p-values) can be derived



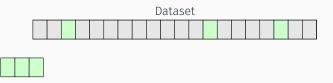




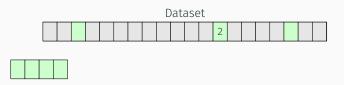






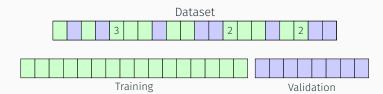




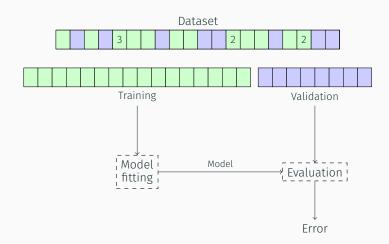




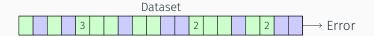




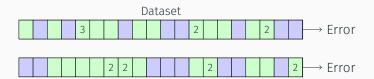


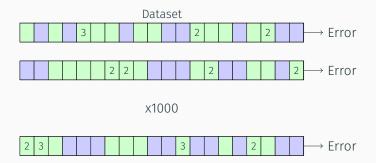




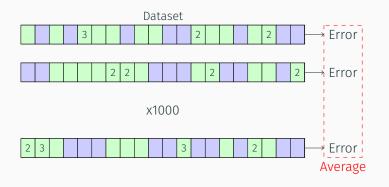








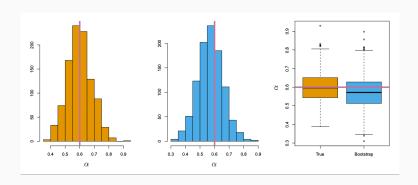






Fits b models with m datapoints (typically m < n), sampled from the original dataset with replacement.

- + Uses all data to train models
- + Provides a dense distribution of model performances
- Versatile: Can be used for other things, e.g. getting a confidence interval for model parameters
- Different choices of b (and exact splits) yields different results





Model evaluation: Comparison

Why do we want to evaluate our model?

- 1. We want to show that our model is better than random guessing
- 2. We want to show that our model is better than another model









There is going to be variability to our model's performance (and possibly the baseline).

Is our model significantly better?





Approach 1:

Is the mean of the distribution of performances from our model significantly higher than the point-estimate baseline?





Approach 1:

Is the mean of the distribution of performances from our model significantly higher than the point-estimate baseline?





Approach 1:

Is the mean of the distribution of performances from our model significantly higher than the point-estimate baseline?



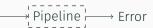


Approach 2:

Is the point-estimate performance of our model significantly higher than the mean of the baseline distribution?



Age	Sex	Feature	Outcome
25	Male	0.53	1
38	Female	-0.76	1
45	Male	0.89	1
33	Female	-0.21	1
29	Male	0.12	1
41	Female	-0.68	0
56	Male	0.45	0
52	Female	-0.32	0
31	Male	0.91	0
48	Female	-0.15	0





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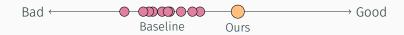




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0.50 → 0.45 0.55



Approach 2:

Is the point-estimate performance of our model significantly higher than the mean of the baseline distribution?

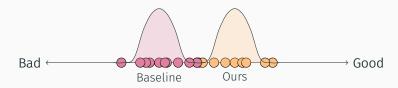




Approach 2:

Is the point-estimate performance of our model significantly higher than the mean of the baseline distribution?





Approach 3:

Is the mean of the distribution of performances from our model significantly higher than the mean of the distribution of baseline performances?



Fold	Ours	Baseline
1	0.75	0.71
2	0.62	0.55
3	0.58	0.57
4	0.87	0.81
5	0.65	0.63
6	0.98	0.97
7	0.55	0.52
8	0.69	0.52
9	0.91	0.85
10	0.88	0.81

The small gain of our model will disappear in the noise between the folds using a non-paired statistical test. Use a paired test, e.g. Wilcoxon signed-rank test

Model evaluation: Summary

- · Model evaluation should always happen out-of-sample
- If n is big (≥ 10000), a single train/validation split is often sufficient
- For smaller samples, k-fold cross-validation with $5 \le k \le 10$ is a good trade-off between bias and variance
- The bootstrap is an effective way of getting confidence intervals for model parameters
- We can use the results from cross-validation (or bootstrapping) to produce a distribution of performances (although caution the correlation)
- We can use a permutation test to produce a distribution of baseline performances
- Compare models across folds using Wilcoxon signed-rank test (ensure the folds are the same!)



Model selection and evaluation



Model selection and evaluation

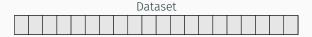
- Model evaluation via cross-validation is sufficient if we want to estimate the out-of-sample error of a known model.
- Very often we want to know whether a set of predictors are informative for an outcome given the best possible model
- In that case, we have to both choose the best model, and estimate its performance
- If we choose the model based on regular cross-validation, the performance estimate will likely be inflated



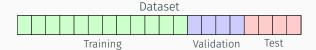
Model selection and evaluation

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- In that case, we have to both choose the best model, and estimate its performance
- If we choose the model based on regular cross-validation, the performance estimate will likely be inflated
- ightarrow We need a more advanced strategy

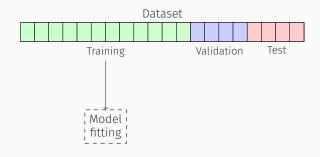




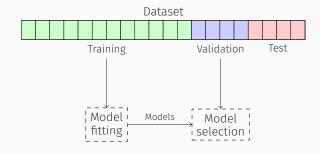




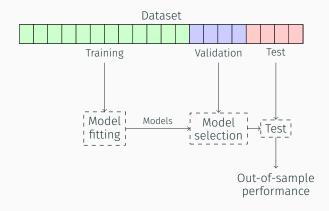




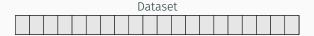






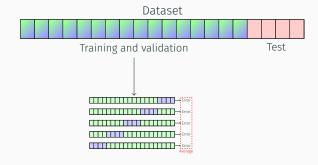




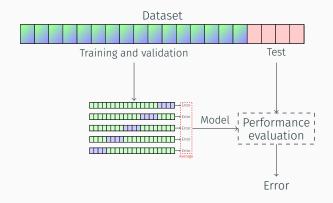




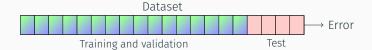




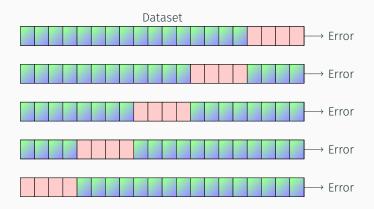




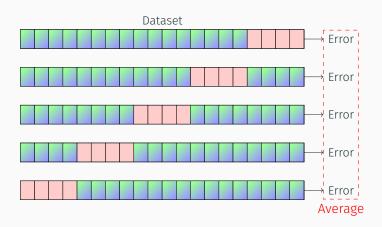














Model selection and evaluation: Summary

- Whenever a choice is made on the basis of performance in a dataset, we have to assume the performance achieved by the chosen model is inflated
- If n is big (≥ 10000), a single train/validation/test split is often sufficient
- When possible, use nested cross-validation to select the best model and estimate the out-of-sample error

