### PSY9511: Seminar 4

Model selection, validation and testing

Esten H. Leonardsen 23.09.24



#### Outline

- 1. Assignment 3
- 2. Loss functions and 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



# Assigment 3



# Assignment 3





#### Commonalities

- · Allows us to evaluate the performance of a model
- Typically on the form  $f(y, \hat{y})$

#### Loss functions

Tailored specifically for mathematical optimization of models

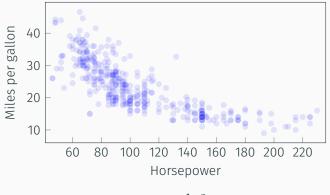
#### Performance metrics

 Tailored specifically for interpretation of model performance by humans



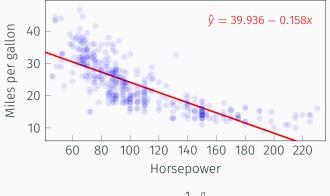
$$mse(y, \hat{y}) = \frac{1}{n} \sum_{i=0}^{n} (y_i - \hat{y}_i)^2$$





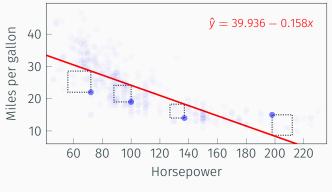
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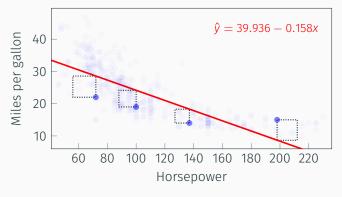
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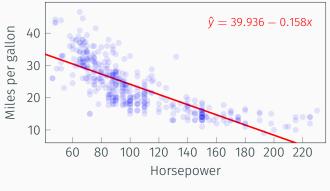
$$mse(y, \hat{y}) = \frac{1}{n} \sum_{i=0}^{n} (y_i - \hat{y}_i)^2$$





$$mse(y, \hat{y}) = \frac{1}{n} \sum_{i=0}^{n} (y_i - \hat{y}_i)^2$$
$$= 23.94$$

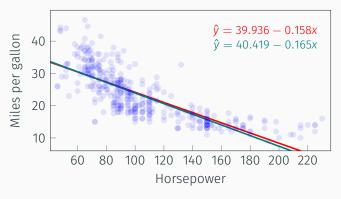




$$mse(y, \hat{y}) = \frac{1}{n} \sum_{i=0}^{n} (y_i - \hat{y}_i)^2$$

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





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

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



#### Loss functions

- Different loss functions measures different properties of the model fit
- Optimizing for them gives different parameter estimates



#### Tolerance-based accuracy:

A prediction is considered correct if it is within a predefined margin of error from the true value

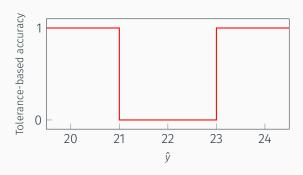
accuracy\*
$$(y, \hat{y}) = \begin{cases} 0 & \text{if } |y - \hat{y}| < \text{tol} \\ 1 & \text{else} \end{cases}$$



mpg	horsepower
22	72

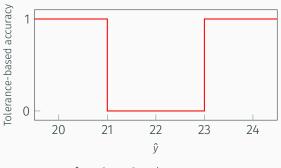


mpg	horsepower
22	72





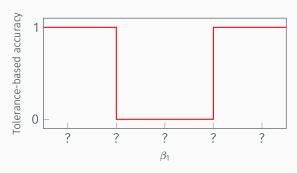
mpg	horsepower
22	72





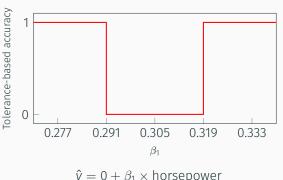


mpg	horsepower
22	72



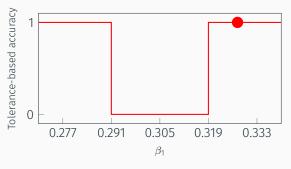
$$\hat{y} = \beta_0 + \beta_1 \times \text{horsepower}$$





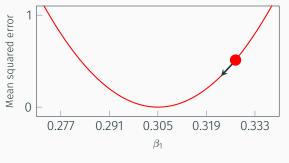
















#### Loss functions

- Different loss functions measures different properties of the model fit
- Optimizing for them gives different parameter estimates
- Must be differentiable to allow for mathematical optimization



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

$$OR$$

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



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

#### Mean squared error (MSE)

- + Can be used as a loss function
- + Widely used
- + Intuitive
- + Penalizes large errors
- ? Interpretation
- Depends on scale



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

#### Root mean squared error (RMSE)

- + Can be used as a loss function
- + Intuitive
- + Penalizes large errors
- + More interpretable than MSE, total loss ≈ individual loss
- Depends on scale



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

#### Mean absolute error (MAE)

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



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

#### Pearson correlation coefficient (r)

- + Scale independent
- ? Captures linear correlation
- Should not be used as a loss function
- Does not care about whether the predictions are close to the true values



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

#### Proportion of variance explained $(r^2)$

- + Scale independent
- + Interpretable
- ? Captures linear correlation
- Should not be used as a loss function
- Does not care about whether the predictions are close to the true values

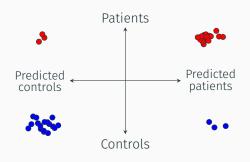






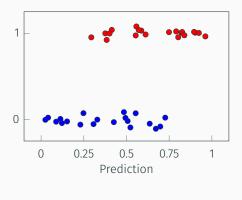
 $y \in \{Patients, Controls\}$ 



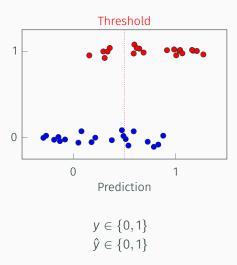


 $y \in \{Patients, Controls\}$  $\hat{y} \in \{Patients, Controls\}$ 

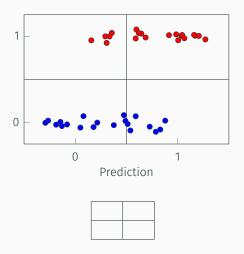




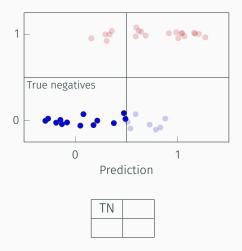
$$y \in \{0, 1\}$$
  
 $\hat{y} \in [0, 1]$ 



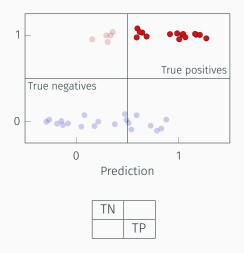




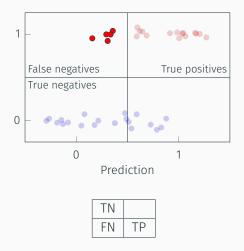




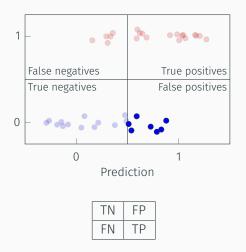














#### Confusion matrix:



#### Binary classification metrics:

- Many metrics rely on thresholding the predictions to obtain binary predictions.
- Although not a metric per se, the confusion matrix is a very useful tool to understand model behaviour, and should always be looked at (and preferably reported).

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

#### Accuracy

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



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

## 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)



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

## 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)

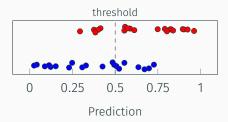


$$\frac{\frac{TP}{TP+FN} + \frac{TN}{TN+FP}}{2}$$

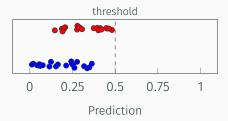
#### Balanced accuracy

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

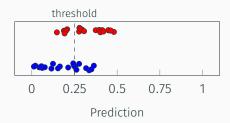




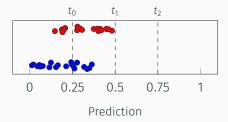




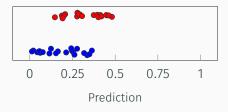




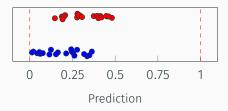






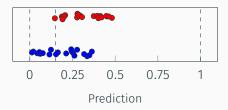


threshold	TPR	FPR

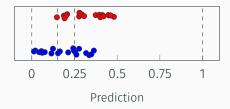


threshold	TPR	FPR
0	1	1
1	0	0



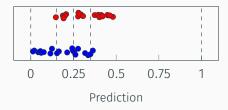


threshold	TPR	FPR
0	1	1
0.15	0.95	0.5
1	0	0



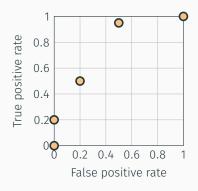
threshold	TPR	FPR
0	1	1
0.15	0.95	0.5
0.25	0.5	0.2
1	0	0





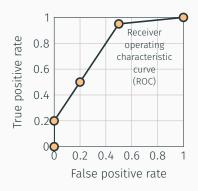
threshold	TPR	FPR
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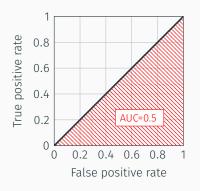
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threshold	TPR	FPR
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0.35	0.2	0.0
1	0	0

#### Area under the receiver operating characteristic curve (AUC/AUROC)

- A performance metric that does not rely on a correct classification threshold
- Measures whether the predictions are ranked correctly (e.g. patients have a higher prediction than controls)
- Handles class imbalance (relatively well) and is commonly reported in the literature



# Loss functions and performance metrics: Summary

# Performance metrics and loss functions measure the performance of a predictive model

- There is a range of alternatives that can be used, each capturing a different aspect of a model's performance
- It is good practice to report more than one metric
- · For regression:
  - MSE is a common loss function with nice mathematical properties.
  - · MAE is an intuitive performance metric
- For classification:
  - Log-loss is the most common loss function for probabilistic classifiers
  - 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



# Loss functions and performance metrics: Summary

http://localhost:8889/notebooks/notebooks%2FClassification%20metrics.ipynb



# Strategies for model evaluation



#### Model evaluation: Rationale

#### Statistical inference:

Goal: In-sample quantification

Predictive modelling:

Goal: Out-of-sample generalization



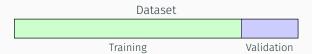
#### Model evaluation: Rationale

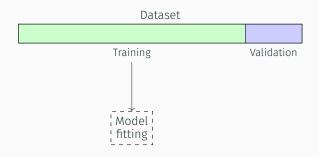
How can we test how good our model is on **unseen data** and **be** certain that performance holds if we present even more new data



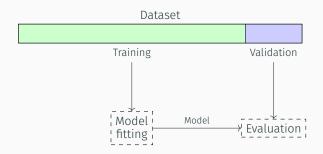




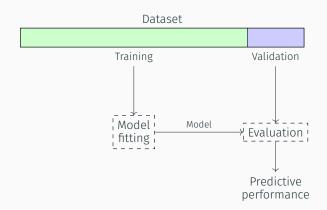










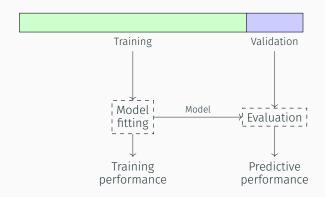




In the validation set approach we split the dataset into two subsets (commonly  $\sim 80\%/20\%$ ), use the first for training the model and the second to test its performance.

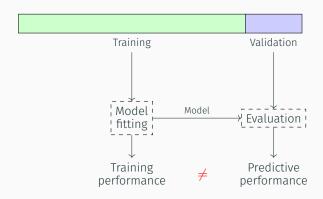
- + Accurate estimate of out-of-sample error
- + Simple
- Variable results depending on the exact split
- Only uses a subset of data for training models
- Gives a point estimate of the error, without confidence intervals

#### Model evaluation: Validation set



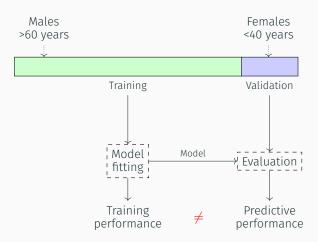


#### Model evaluation: Validation set





#### Model evaluation: Validation set





#### Stratification:

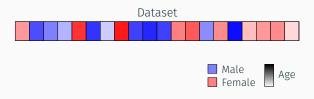
Ensuring all folds of the dataset are similar with respect to some given characteristics.



# Dataset

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





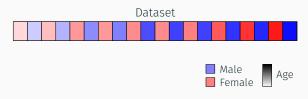
```
In[1]: df = ...
```





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



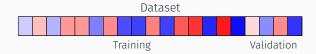


```
In[1]:    df = ...
    df = df.sort_values(['sex', 'age'])
```











#### Stratification:

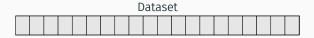
Ensuring all folds of the dataset are similar with respect to some given characteristics.

- Helps alleviate the risk of training performance >> validation performance
- · Always stratify on target variable first

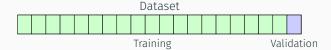
stratified(data, columns, split)

 Also good idea to stratify on other core characteristics, e.g. sex and age

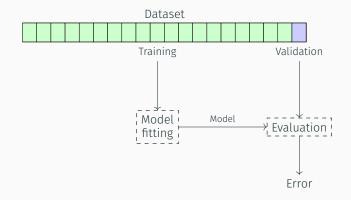




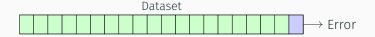




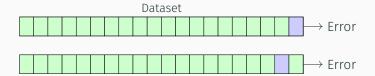




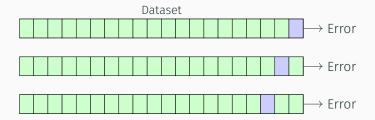


















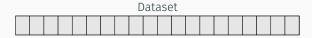




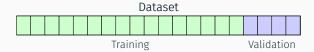
Fits *n* models for *n* datapoints, each time leaving a single datapoint out for testing.

- + 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
  - → highly correlated

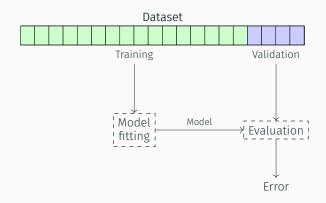




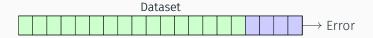




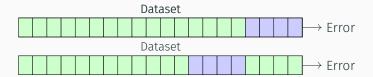




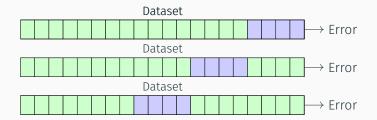




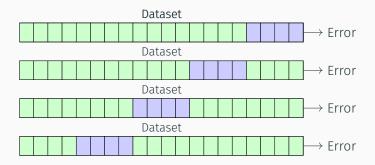




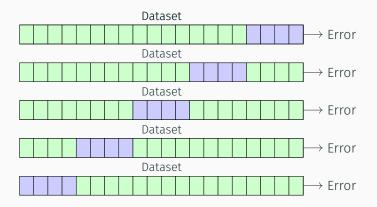


















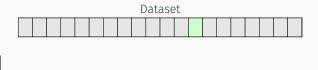
Fits k (usually  $k \in \{5, 10\}$ ) models for n > k datapoints, each leaving n/k datapoints for out-of-sample testing.

- + 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





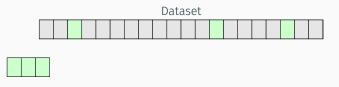




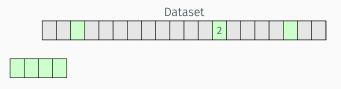








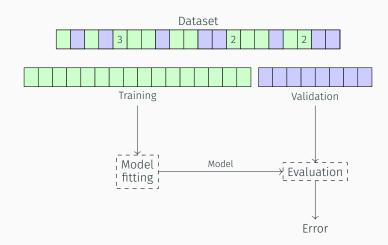




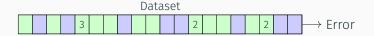




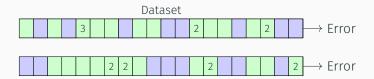




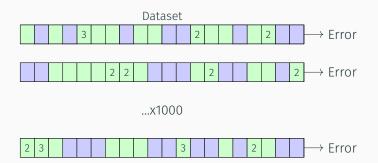




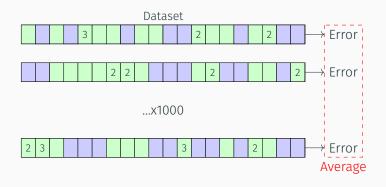










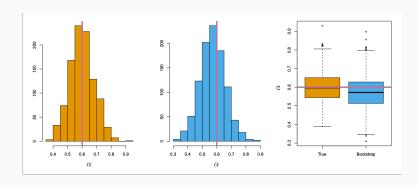




Fits b models with m datapoints (typically m < n), sampled from the original dataset with replacement.

- + 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



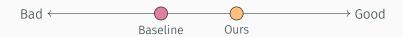




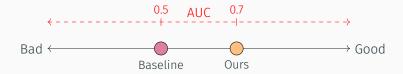
#### Why do we want to evaluate our model?

- 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











http://localhost:8890/notebooks/notebooks%2FModel%20variability.ipynb





There is going to be variability in 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 (with regards to variability that is **unrelated** to efficacy) significantly higher than the point-estimate baseline?

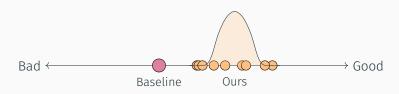




#### Approach 1:

Is the mean of the distribution of performances from our model (with regards to variability that is **unrelated** to efficacy) significantly higher than the point-estimate baseline?





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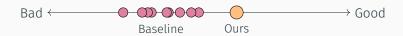




#### Approach 2:

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?



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



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#### 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  $\leq$  k  $\leq$  10 is a good trade-off between bias and variance
- The bootstrap is an effective way of getting confidence intervals for model performance and parameters
- Cross-validation (or bootstrapping) will produce a distribution of model performances (although caution the correlation)
- Permutation testing will produce a distribution of baseline performances
- Compare models across folds using Wilcoxon signed-rank test (ensure the folds are the same!)

