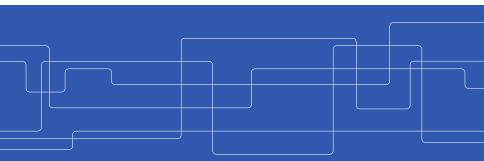


#### ID2214 Programming for Data Science - Evaluating Predictive Models

#### Henrik Boström

Prof. of Computer Science - Data Science Systems Dept. of Software and Computer Systems School of Electrical Engineering and Computer Science KTH Royal Institute of Technology bostromh@kth.se

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Evaluation protocols

Performance Metrics for Classification

Performance Metrics for Regression

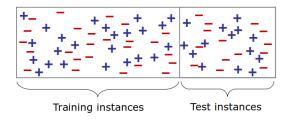
**Empirical Investigations** 



#### Predictive Modeling

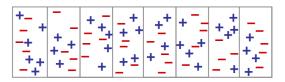
Name	Solu- bility	No. C atoms	Fraction of rotatable bonds	Topol. diam.	Geom. diam.	Log P	No. heavy bonds	
methylpentane	good	6	0.40	4	3.46	2.44	5	
methylcyclohexene	good	7	0	4	3.00	2.51	7	
nonene	med.	9	0.75	8	6.93	3.53	8	
hexadiene	good	6	0.60	5	4.36	2.14	5	
butadiene	good	4	0.33	3	2.65	1.36	3	
naphthalene	good	10	0	5	3.61	2.84	11	
acenaphthylene	good	12	0	5	3.58	3.32	14	
pyrene	poor	16	0	7	5.00	4.58	19	
dimethylanthracene	poor	16	0	7	5.29	4.61	18	
hexahydropyrene	med.	16	0	7	5.00	3.82	19	
triphenylene	poor	18	0	7	5.00	5.15	21	
benzo(e)pyrene	poor	20	0	7	5.29	5.64	24	





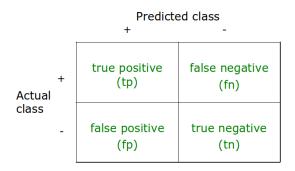
Stratified split = class proportions are approximately the same





- The procedure is called *leave-one-out cross-validation*, if N = no. of instances
- A common choice is N = 10
- Stratification may also be employed here; stratified cross-validation







Accuracy; fraction of correct predictions

$$\textit{Accuracy} = \frac{\textit{tp} + \textit{tn}}{\textit{tp} + \textit{fp} + \textit{tn} + \textit{fn}}$$

Precision; fraction of correct predictions for a class

$$Precision = \frac{tp}{tp + fp}$$

Recall; fraction of certain class correctly predicted

$$\mathit{Recall} = rac{\mathit{tp}}{\mathit{tp} + \mathit{fn}}$$

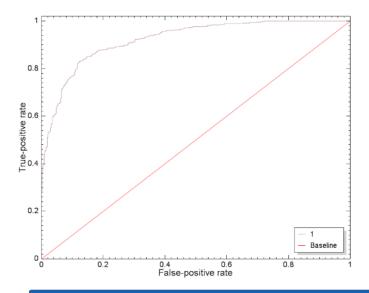


#### Performance Metrics for Classification (cont.)

Correct class			Predict	Predicted class			
	Precision	Recall	good	medium	poor		
good	0.957	0.974	332	1	8		
medium	0.000	0.000	12	0	6		
poor	0.632	0.857	3	1	24		

$$\textit{Accuracy} = \frac{332 + 0 + 24}{341 + 18 + 28} \approx 91.99\%$$

# Receiver Operating Characteristic (ROC) Curve







Plotting a ROC curve

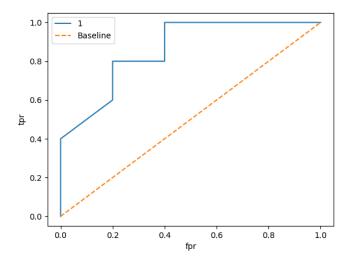
import matplotlib.pyplot as plt

```
pos = [1,1,1,1,0,1,0,0]
neg = [0,0,1,0,1,0,2,1]
```

tpr = [cs/sum(pos) for cs in np.cumsum(pos)]
fpr = [cs/sum(neg) for cs in np.cumsum(neg)]



#### Plotting (cont.)





# Receiver Operating Characteristic (ROC) Curve (cont.)

- The area under the ROC curve (AUC) = the probability of an example belonging to the class being ranked ahead of an example not belonging to the class
- For binary classification tasks, the AUC will be the same for both classes.
- In case there are more than two classes, the resulting AUC may be calculated as the weighted average of the individual AUCs, using relative class frequencies as the weights.

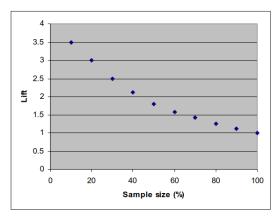
# Calculating the Area under ROC Curve (AUC)

Input: (s\_1,tp\_1,fp\_1), ..., (s\_n,tp\_n,fp\_n) (sorted triples of scores, no. true and false pos. with the scores wrt some class c), Tot\_tp, and Tot\_fp Output: AUC

```
AUC = 0
Cov_tp = 0
for i = 1 to n
    if fp_i = 0 then Cov_tp += tp_i
    else if tp_i = 0 then
        AUC += (Cov_tp/Tot_tp)*(fp_i/Tot_fp)
    else
        AUC += (Cov_tp/Tot_tp)*(fp_i/Tot_fp)+
            (tp_i/Tot_tp)*(fp_i/Tot_fp)/2
        Cov_tp += tp_i
```

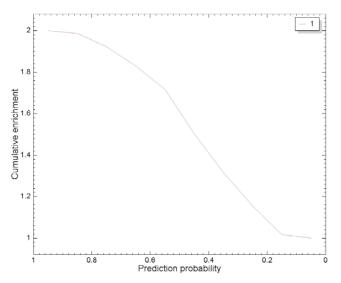


#### Cumulative Lift Chart



$$Lift = \frac{\frac{tp}{tp+fp}}{\frac{TP}{TP+FP}}$$

## Cumulative Lift Chart (alt.)





 Brier score (*quadratic loss*); mean squared error of the predicted probabilities

Brier score 
$$=rac{1}{n}\sum_{i=1}^n(p_i-o_i)^2$$

where  $p_i$  are the predicted and  $o_i$  the actual (observed) probabilities for test instance *i*, where typically all values are zero in  $o_i$ , except one (corresponding to the true class label)



 Log loss (*informational loss*); mean logarithm of the predicted probabilities for the true class labels

$$Log \ loss = -\frac{1}{n} \sum_{i=1}^{n} o_i \ log \ p_i$$

where  $p_i$  are the predicted and  $o_i$  the actual (observed) probabilities for test instance *i*, where typically all values are zero in  $o_i$ , except one (corresponding to the true class label)



Performance Metrics for Regression

$$MSE = \frac{1}{n}\sum_{i=1}^{n}(p_i - o_i)^2$$

Root Mean Squared Error (RMSE)

$$RMSE = \sqrt{\frac{1}{n}\sum_{i=1}^{n}(p_i - o_i)^2}$$

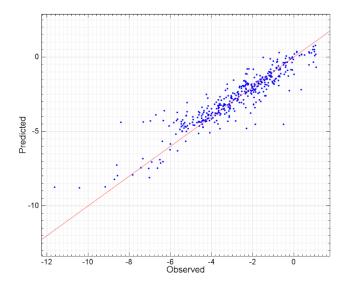
Mean Absolute Error (MAE)

$$MAE = \frac{1}{n} \sum_{i=1}^{n} |p_i - o_i|$$

where  $p_i$  is the predicted and  $o_i$  the actual (observed) target (regression) value for test instance *i* 



#### Predicted vs. Observed Plot





#### Pearson (product moment) correlation coefficient =

$$\frac{\sum p_i o_i - n\bar{p}\bar{o}}{\sqrt{(\sum p_i^2 - n\bar{p}^2)}\sqrt{(\sum o_i^2 - n\bar{o}^2)}}$$

where  $p_i$  is the predicted and  $o_i$  the actual (observed) target (regression) value for test instance i, and  $\bar{p}$  and  $\bar{o}$  are the corresponding averages



What is to be investigated?

- ► What is the predictive performance of *model M* on new (unseen) data?
- ► Is there a (significant) difference between model M<sub>1</sub> and M<sub>2</sub> (or between M<sub>1</sub>, M<sub>2</sub>, M<sub>3</sub>, ...)?
- ► Is there a significant difference between *algorithm A*<sub>1</sub> and *A*<sub>2</sub> (or between *A*<sub>1</sub>, *A*<sub>2</sub>, *A*<sub>3</sub>, ...)?



#### Predictive Performance of a Model

- We assume that we have chosen some performance metric, e.g., accuracy or MSE, for which we can obtain a sample of measurements for the model
- Given the sample, we may infer a *confidence interval*, i.e., by following the procedure we will with high probability obtain an interval that contains the true performance of the model
- Based on the confidence interval, we may perform statistical hypothesis testing, e.g., conclude that the model performance is significantly different from some baseline level, i.e., the baseline falls outside the confidence interval



### Predictive Performance of a Model (cont.)

Questions to ask when evaluating the performance:

- Is the sample representative, i.e., randomly sampled from the target distribution?
- Is it independent, e.g., parameter settings, model choice, etc. were not based on the data?
- ▶ Is the sample size (*N*) large enough?
  - For highly skewed distributions, N needs to be higher than 30, to allow for using the normal distribution when inferring confidence intervals
  - For a proportion P, e.g., accuracy, PN > 10 and (P-1)N > 10
- Is there actually a (single) underlying population from which the sample is drawn?
  - E.g., can the scores obtained from 10-fold cross-validation be considered to be a sample drawn from some population?



- The null hypothesis assumes there is no difference between the models
- A (sufficiently large) random sample is collected, e.g., measured differences in predictive performance
- A significance level is chosen and if measurements of the sample deviate significantly from what can be expected if the null hypothesis is true, then the null hypothesis is rejected, and the alternative hypothesis is accepted



## Statistical Errors

Decision	Truth value for $H_0$			
	True	False		
Reject $H_0$	Error of type I	Correct		
Do not reject $H_0$	Correct	Error of type II		



When performing multiple pairwise comparisons, the elevated risk of type I error may be controlled, by *Bonferroni correction*, i.e., divide the significance level by the number of tested hypotheses

## Comparing Multiple Algorithms

When comparing multiple algorithms over multiple datasets, the standard procedure is to employ a *Friedman test*, followed by some suitable *post hoc* test, e.g., Nemenyi, to reject (some of) the pairwise hypotheses, see e.g.,

Demšar, J., 2006. Statistical comparisons of classifiers over multiple data sets. Journal of Machine learning research, 7, pp.1-30

Python implementation: https://docs.orange.biolab.si/3/datamining-library/reference/evaluation.cd.html

Garcia, S. and Herrera, F., 2008. An extension on "statistical comparisons of classifiers over multiple data sets" for all pairwise comparisons. Journal of Machine Learning Research, 9, pp.2677-2694.

Java implementation: http://sci2s.ugr.es/keel/multipleTest.zip



- The choice of performance metric is (at least) as important as the choice of learning algorithm; careful consideration of what needs to be optimized is required
- Common traps should be avoided, e.g., dependencies between training and test data, over-fitting the test set by repeated experimentation.
- A careful formulation of one or more null hypotheses are needed for empirical investigations; what exactly are to be compared, etc.