#### **EMBEDDED VISION DESIGN 3**

# **ML PERFORMANCE**

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# **CONTENTS**

- Confusion matrix
- Evaluating classifiers
- Learning curves

# THE BOY WHO CRIED WOLF

"Wolf" is a **positive class**.

"No wolf" is a negative class

An Aesop's Fable ~620 BCE



Source: Sam Taplin

## **CONFUSION MATRIX**

#### **ACTUAL**

(Type I error)

#### **True Positive (TP)**

Reality: A wolf threatened. Shepherd said: "Wolf."

Outcome: Shepherd is a hero.

### **False Positive (FP)**

Reality: No wolf threatened. Shepherd said: "Wolf."

Outcome: Villagers are angry at shepherd for waking them up.

#### False Negative (FN)

Reality: A wolf threatened. Shepherd said: "No wolf."

Outcome: The wolf ate all the sheep.

Type II error)

#### True Negative (TN)

Reality: No wolf threatened. Shepherd said: "No wolf." Outcome: Everyone is fine.

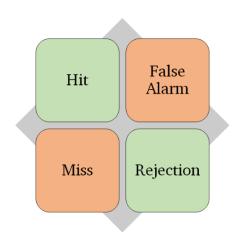
# **ACCURACY**

Fraction of predictions the model got right

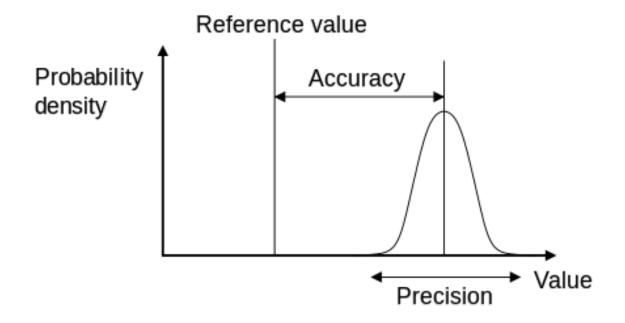
$$\label{eq:accuracy} Accuracy = \frac{Number\ of\ correct\ predictions}{Total\ number\ of\ predictions}$$

For binary classification

$$\text{Accuracy} = \frac{TP + TN}{TP + TN + FP + FN}$$



# **ACCURACY VS PRECISION**



**accuracy** is closeness of the measurements to a specific value, while **precision** is the closeness of the measurements to each other.

## PRECISION AND RECALL

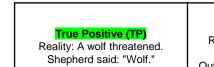
- Precision, fraction of correct positive predictions
  - $\Sigma$  True positive /  $\Sigma$  Predicted condition positive

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

Recall, probability of detection

 $\Sigma$  True positive /  $\Sigma$  Condition positive

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



precision

Outcome: Shepherd is a hero.

#### False Negative (FN) Reality: A wolf threatened.

ecall

Shepherd said: "No wolf." Outcome: The wolf ate all the sheep.

#### False Positive (FP)

Reality: No wolf threatened. Shepherd said: "Wolf." Outcome: Villagers are angry at shepherd for waking them up.

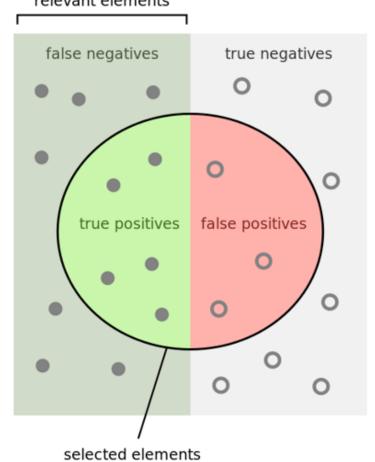
#### True Negative (TN)

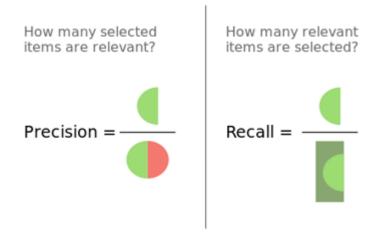
Reality: No wolf threatened. Shepherd said: "No wolf." Outcome: Everyone is fine.

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# PRECISION AND RECALL

#### relevant elements

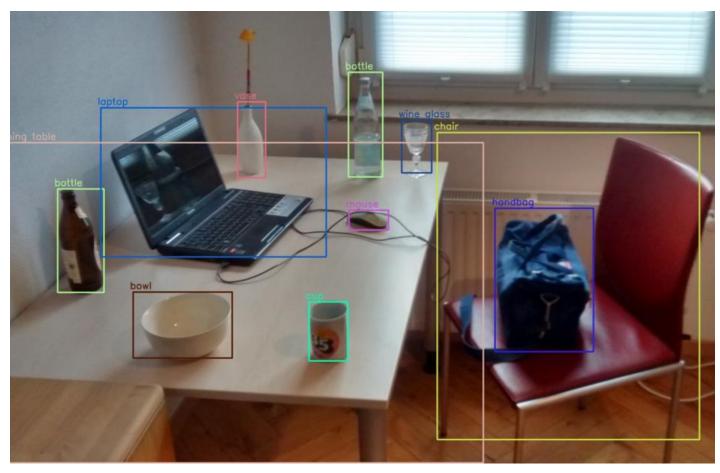




Recall = sensitivity = true positive rate (TPR)

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# LIMITATION OF USING SENSITIVITY AND SPECIFICITY



# **LET'S CHECK A VIDEO CLIP**

https://www.youtube.com/watch?v=TtljAiSojFE

	precision	recall	f1-score
0.0	0.96	0.97	0.97
1.0	0.65	0.58	0.62

### F1 SCORE

- To fully evaluate the effectiveness of a model, you must examine
   both precision and recall
- F1 score is the harmonic mean of precision and recall

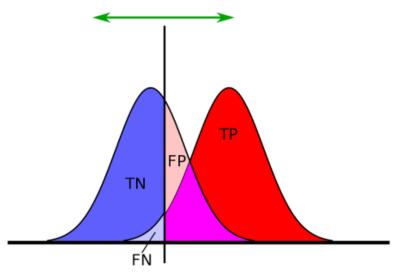
$$F_1 = \frac{2}{\frac{1}{\text{precision}} + \frac{1}{\text{recall}}} = 2 \times \frac{\text{precision} \times \text{recall}}{\text{precision} + \text{recall}} = \frac{TP}{TP + \frac{FN + FP}{2}}$$

# **MANY METRICS**

	veren Alle tests van de deele		Source: https://en.wikipedia.org/wiki/Receiver_operating_characteristic		
	Prediction of the test was a de deelnemende partijen zijn eerst beoordeeld door het RIVM. "We sturen tien monsters, vijf met SARS-CoV-2 in verschillende concentraties, drie met andere coronavirussen, en twee waar niets in zit. De labs weten niet wat waar in zit. Als hun uitslagen kloppen, is de test goed", zegt Reusken.  De testprestaties zullen per lab wat variëren: de specificiteit (de kans dat je een niet-ziek iemand ook als niet-ziek detecteert) en de sensitiviteit (de kans dat je een ziek iemand als ziek detecteert). De E-gen test is gevoeliger dan de RdRP-gen test, maar wat minder specifiek: naar schatting 99 tot 99,5 procent. Tests op meerdere genen zijn iets specifieker. Door de variatie ligt die specificiteit landelijk minimaal op 99,5 procent, zegt Reusken.  4   Wat bepaalt hoeveel foute uitslagen een test geeft?  "De specificiteit hangt af van de test zelf	n tigative	$= \frac{\text{Prevalence}}{\sum \text{Total population}}$	$\frac{\text{Accuracy (ACC)} =}{\sum \text{True positive} + \sum \text{True negative}}{\sum \text{Total population}}$	
Predic		g Zitive, F <sub>ror</sub>	Positive predictive value  (PPV), Precision = $\Sigma$ True positive $\Sigma$ Predicted condition positive	False discovery rate (FDR) =  Σ False positive  Σ Predicted condition positive	
condi		s r ritive	False omission rate (FOR) =  Σ False negative Σ Predicted condition negative	Negative predictive value (NPV) =  Σ True negative  Σ Predicted condition negative	
		e rate -out, -out, -se alarm	Positive likelihood ratio (LR+) = TPR FPR	Diagnostic odds ratio (DOR)	F <sub>1</sub> score = 2 · Precision · Recall Precision + Recall
		True  (TNR)  Jative  negative	Negative likelihood ratio (LR-) $= \frac{FNR}{TNR}$	= <u>LR+</u> LR-	Trocision - Nocul

# PRECISION/RECALL TRADE-OFF

#### Decision threshold



$$precision = \frac{TP}{TP + FP} \qquad recall = \frac{TP}{TP + FN}$$

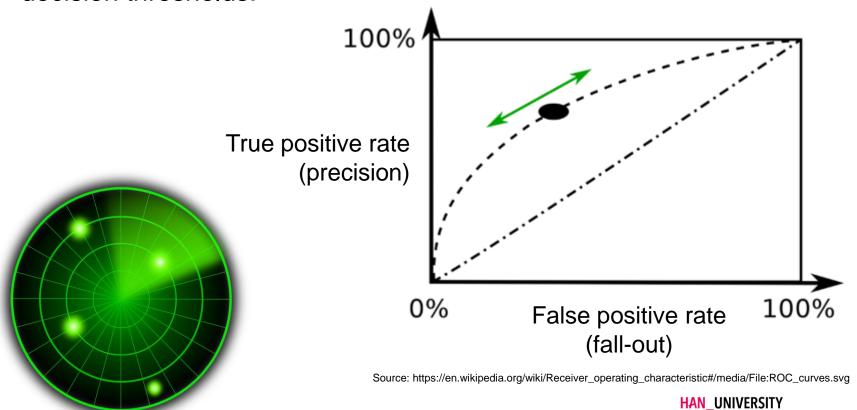


Source: https://en.wikipedia.org/wiki/Tug\_of\_war#/media/File:Touwtrekken.jpg



# **ROC CURVE**

• probability of detection vs probability of false alarm at different decision thresholds.

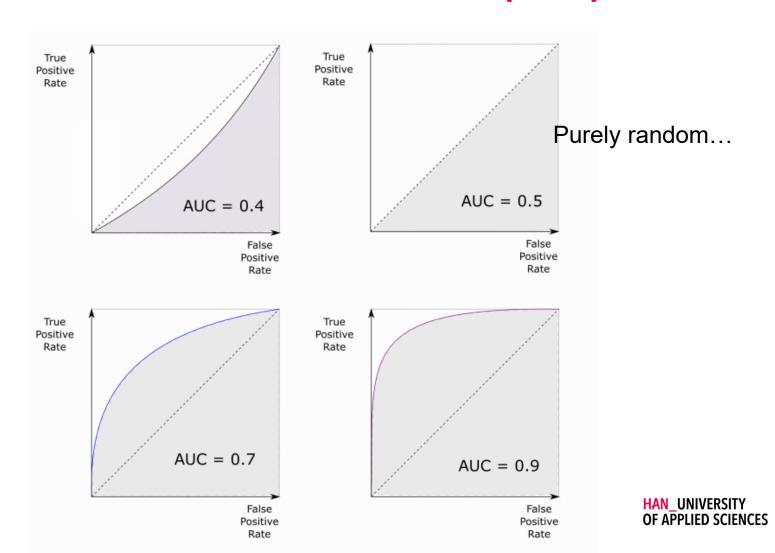


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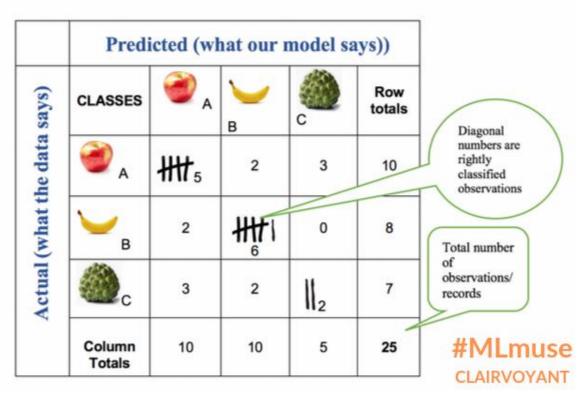
## **COST OF CLASSIFICATION**

- Sometimes false negatives don't hurt as much as false positives Think of a poisonous mushroom detector....
- Use the ROC curve (receiver operating characteristics) to help balance the cost of classification

# **ROC AREA UNDER THE CURVE (AUC)**



# **MULTICLASS CONFUSION MATRIX**



Source: https://miro.medium.com/max/1400/1\*jtoE1zEJaG0JvGIX3jOTFQ.png

## **SPLITTING DATA**

Slice data into three subsets: Training, validation and test data

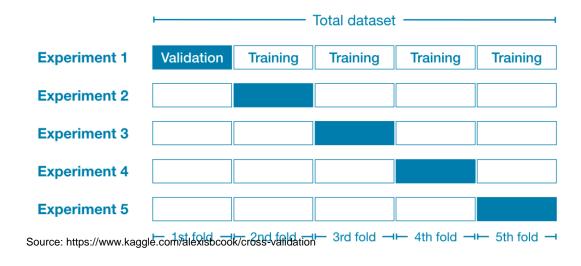


- Make sure that your subsets meet the following conditions:
  - Large enough to yield statistically meaningful results.
  - Representative of the data set as a whole.
     E.g. don't pick a test set with different characteristics than the training set.



# **CROSS-VALIDATION**

- Estimate of a model's generalization performance
- Break the data into folds



 For small datasets, where extra computational burden isn't a big deal, you should run cross-validation.

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# **LEARNING CURVES**

```
from sklearn.model_selection import learning_co
from sklearn.svm import SVC
from sklearn.datasets import load_digits
from matplotlib import pyplot as plt
import numpy as np

X, y = load_digits(return_X_y=True)
estimator = SVC(gamma=0.001)
```

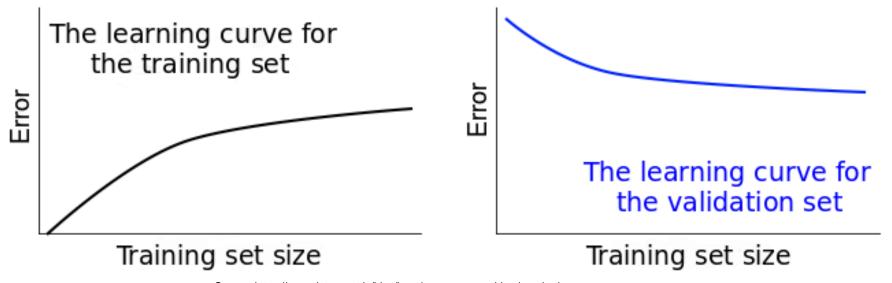
```
1.000
  0.975
  0.950
e 0.925
  0.900
  0.875
  0.850
                                                                     train
                                                                     test
                   400
                          600
                                  800
                                         1000
                                                 1200
           200
                                                         1400
                                                                 1600
                                      data size
```

```
train_sizes, train_scores, test_scores, fit_times, _ = learning_curve(estimator, X, y, cv=30,return_times=True)
```

plt.plot(train\_sizes,np.mean(train\_scores,axis=1))

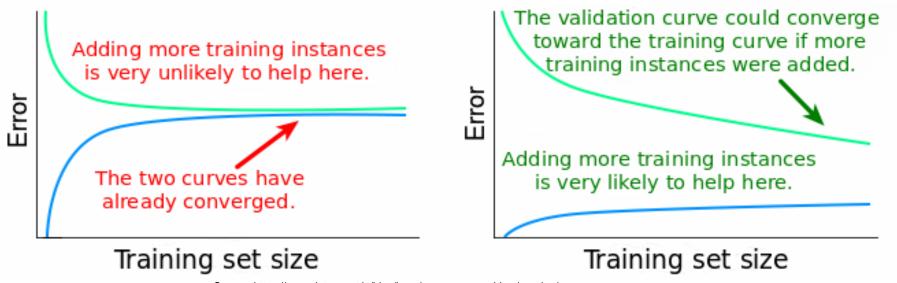
# **LEARNING CURVES**

- Cost as a function of the training set size (or the training iteration)
- Examine evolution of train and validation learning curves



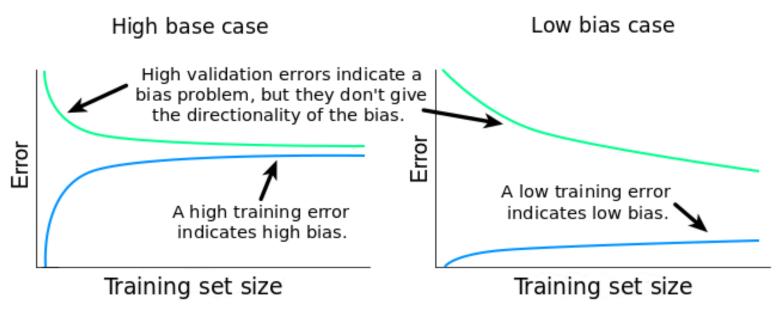
# **LEARNING CURVES**

- Convergence of curves
- Diagnose generalization problems



# **BIAS PROBLEM**

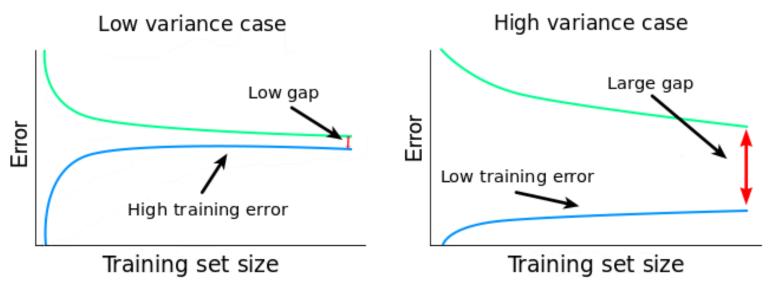
- High validation error indicates a prediction bias problem
- Underfitting usually gives high bias





# **VARIANCE PROBLEM**

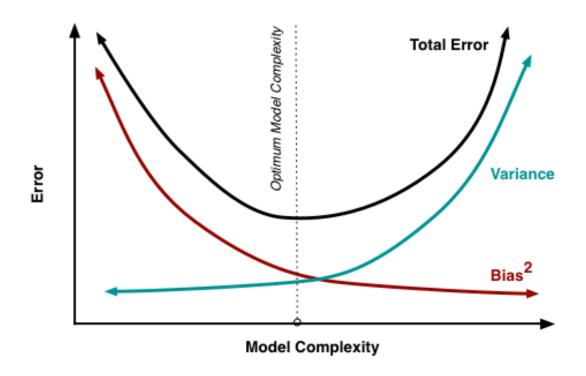
- Low gap indicates low prediction variance
- Overfitting usually gives high variance





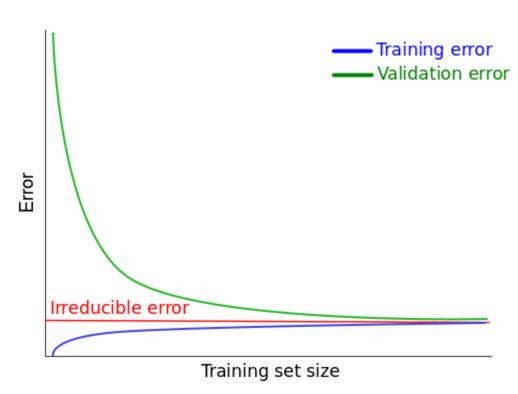
# PREDICTION BIAS-VARIANCE TRADEOFF

Central problem in supervised learning



# **IRREDUCIBLE ERROR**

- Noise
- Cannot be predicted
- Loss cannot be reduced



## **MORE ON PREDICTION BIAS**

- Average of predictions ≈ average of labels in test set
- A significant difference shows there is bias
- Possible causes:
  - Underfitting, e.g. incomplete feature set, overly strong regularization
  - Biased training samples
  - (Noisy data set)

# **QUESTION**

- We know that on average, 1% of all emails are spam.
- My spam filter predicts that 20% of my incoming mail is spam.

What can we say about my spam filter?



# **COMPUTING CROSS-VALIDATED METRICS**

Predefined scoring parameters

Scoring	Function	Comment
Classification		
'accuracy'	metrics.accuracy_score	
'balanced_accuracy'	metrics.balanced_accuracy_score	
'average_precision'	metrics.average_precision_score	
'neg_brier_score'	metrics.brier_score_loss	
'f1'	metrics.f1_score	for binary targets
'f1_micro'	metrics.f1_score	micro-averaged
'f1_macro'	metrics.f1_score	macro-averaged
'f1_weighted'	metrics.f1_score	weighted average
'f1_samples'	metrics.f1_score	by multilabel sample
'neg_log_loss'	metrics.log_loss	requires predict_proba support
'precision' etc.	metrics.precision_score	suffixes apply as with 'f1'
'recall' etc.	metrics.recall_score	suffixes apply as with 'f1'
'jaccard' etc.	metrics.jaccard_score	suffixes apply as with 'f1'
'roc_auc'	metrics.roc_auc_score	
'roc auc ovr'	metrics.roc auc score	

See: https://scikit-learn.org/stable/modules/model\_evaluation.html

