

ML2 – the basics

Machine Learning – Tools and applications for policy – Lecture 3

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DNB Data Science Hub



ML2 – the basics

1. How do I assess success?

- Confusion matrix, Receiver Operator Characteristic (ROC)

2. What are overfitting, bias and variance?

- L2-regularisation

3. Support Vector Machines (SVM) and k nearest neighbours (KNN)

ML2 – the basics

- Measuring success

- Overfitting, bias and variance

- Fixing overfitting

- Classification: Support Vector Machines (SVM)

- K-nearest neighbors (KNN)

- Evaluation: Simple Measures for Classification (CompStat Munich) ([link](#))
Focus on 3.12-6.10 for the explanation on using a **cost function**. The remainder of the clip talks about other cost functions (Brier etc.) – advanced topic.
- Evaluation: Measures for Binary Classification: ROC Measures (CompStat Munich) ([link](#))
Focus on **imbalanced classes** at the start (e.g. defaults) and confusing naming (11.39-13.00)
- Evaluation: Measures for Binary Classification: ROC visualization (CompStat Munich) ([link](#))
If you are interested in deriving ROC/AUC from first principles
- How do Support Vector Machines work? (Brandon Rohrer) ([link](#))
- Decision Boundary (Andrew Ng) ([link](#))
- To discuss in class: [Decision tree animation](#)
- To discuss in class: [Bias Variance](#)

ML2 – the basics

Measuring success

Overfitting, bias and variance

Fixing overfitting

Classification: Support Vector Machines (SVM)

K-nearest neighbors (KNN)



- So far we've mainly been focussed on **accuracy**
- Accuracy can be misleading for **imbalanced datasets**
 - If 99.9% of all days it does not rain, then it will be difficult to beat the very simple predictor: it will be dry tomorrow
- We need to compute the performance for the off-diagonal
 - Is it bad if I **predict sunshine** but it **rains**? Is it bad if I **predict rain** but it **is sunny**?
 - Is it bad if you test **positive for Covid** but **you are healthy**? Is it bad if I **say you are healthy** but you **have Covid**?
- **Confusion matrix** helps visualize different types of errors a classifier can make by reporting the counts of these errors
 - true positive (TP), true negative (TN), false positive (FP), and false negative (FN) predictions

Accuracy

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

Error

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

Precision

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

Recall

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

F1 score

$$F1 = 2 \times \frac{Prec \times Rec}{Prec + Rec}$$

		Prediction		Tot.
		pos	neg	
Actual value	pos'	True positive (TP)	False negative (FN)	TP + FN
	neg'	False positive (FP)	True negative (TN)	FP + TN
Tot.		TP + FP	FN + TN	

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		pos	neg	Tot.
Actual value	pos'	True positive (TP)	False negative (FN)	TP + FN
	neg'	False positive (FP)	True negative (TN)	FP + TN
Tot.		TP + FP	FN + TN	

- The costs for the 'boxes' / categories might differ
- What if the cost of missing a sick person are much higher than missing a healthy person?
 - If 0=Healthy, 1=sick \rightarrow missing sick = 2x healthy \rightarrow FN == 2x FP
- **precision** is for the scared bureaucrat, embarrassed to get a wrong prediction
- **recall** is the superhero that needs to defuse all the bombs and not miss one
- See CompStat Munich clip ([link](#)). Focus on 3.12-6.10 for the explanation on using a **cost function**. The remainder of the clip talks about other cost functions (Brier etc.) – advanced topic
- Also be mindful of **imbalanced classes** (See the start of the [CompStat Munich link](#))

Question: 0.1% of the population has a disease, and a test detects it 99% of the time but falsely identifies 5% of healthy people as sick. What is the likelihood of a positive test result being accurate?

Each format below conveys the same fundamental information about risk structure.

A Single-event probability format

Bayes' theorem is necessary (and difficult) when using single-event probabilities to calculate the probability of a hypothesis (having the disease) given the evidence for it (a positive test result).

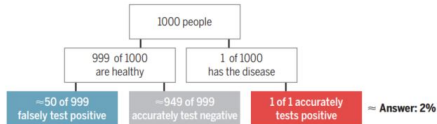
$$\begin{aligned}P(\text{disease}) &= 0.1\% \text{ prevalence of disease} \\P(\text{positive test}|\text{disease}) &= 99\% \text{ true positive rate} \\P(\text{positive test}|\text{no disease}) &= 5\% \text{ false positive rate}\end{aligned}$$

$$P(\text{disease}|\text{positive test}) = \frac{0.1\% \times 99\%}{(0.1\% \times 99\%) + (99.9\% \times 5\%)} = 1.94\% \approx \text{Answer: 2\%}$$

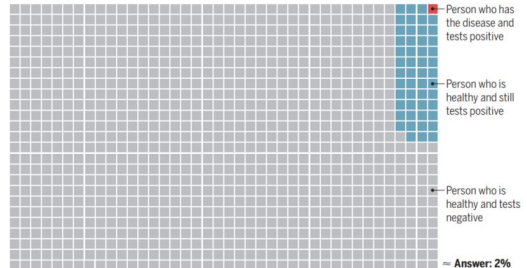
B Nested set format

This view facilitates accurate judgment because it represents base rates (prevalence) and reference class size (1 of 1000) without having to multiply a conditional probability by the base rate.

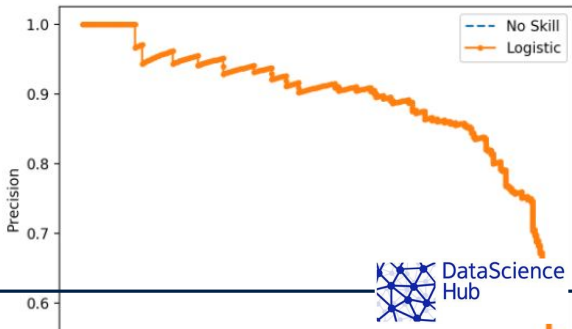
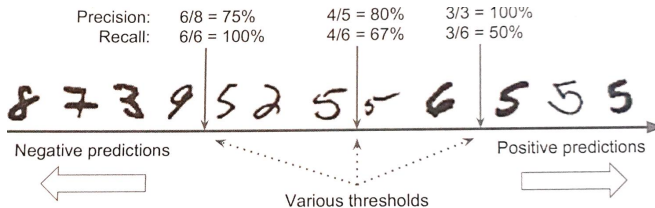
$$P(\text{disease}|\text{positive test}) = \frac{P(\text{disease AND positive test})}{P(\text{positive test})} = \frac{1}{50 + 1} = 1.96\% \approx \text{Answer: 2\%}$$



C Pictograph format



- Choosing the right threshold can lead to any level of precision
- we can see the trade-off if we plot **recall** against **precision**
- More common **Receiver Operating Curve** or **ROC**
 - True Positive Rate == Recall
 - False Positive Rate == $FP / (FP + TN)$ == 1 - Specificity



ML2 – the basics

Measuring success

Overfitting, bias and variance

Fixing overfitting

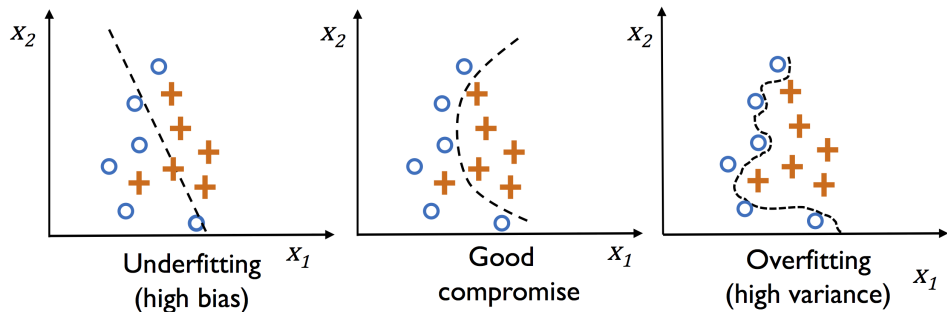
Classification: Support Vector Machines (SVM)

K-nearest neighbors (KNN)



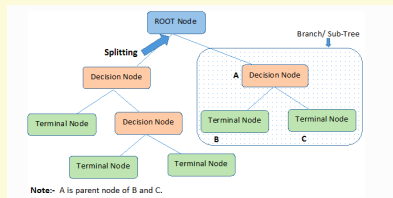
- **Overfitting**: sometimes **model performs well** on **training data** → **low bias** ...
... but does **not generalize well** to unseen data (**test data**)
 - If a model suffers from overfitting, the model has a **high variance**
 - This is often caused by a model that's too complex
- **Underfitting** can also occur (**high bias**)
 - Underfitting is caused by a model's not being complex enough
- Both suffer from low performance on unseen data

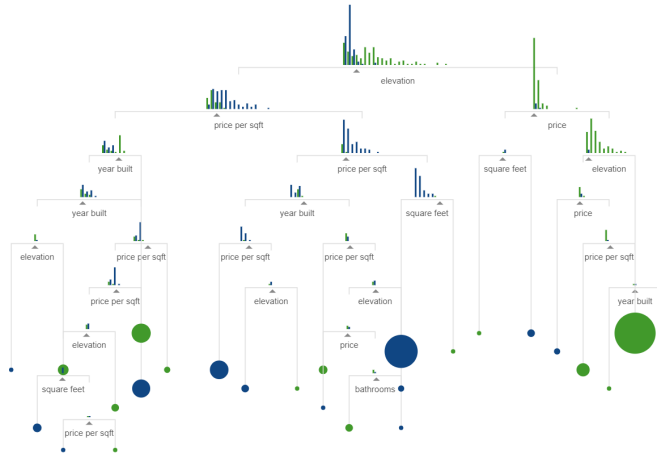
Seen as classification

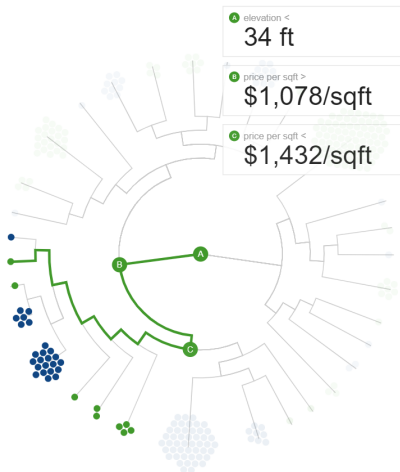


Seen as regression

- **Root Node**: the entire population which gets divided into two or more homogeneous sets
- **Splitting**: It is a process of dividing a node into two or more sub-nodes
- **Decision Node**: When a sub-node splits into further sub-nodes
- **Leaf/Terminal Node**: Nodes that do not split
- **Pruning**: When we remove sub-nodes of a decision node. The opposite process of splitting
- **Branch/Sub-Tree**: Subsection of the entire tree
- **Parent/Child Node**: A node, which is divided into sub-nodes is a parent node of sub-nodes. Sub-nodes are the children







ML2 – the basics

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- Regularization is a way to tune the complexity of the model
- Regularization helps to filter out noise from training data
- As a result, regularization prevents overfitting
- There are two main forms of regularization
 1. **L1 regularization**: covered in Lecture *ML3 – dimensionality and assessment*
 2. **L2 regularization**: weight decay

The most common form of regularization is the so-called **L2 regularization** (sometimes also called **L2 shrinkage** or **weight decay**):

$$\frac{\lambda}{2} \|\mathbf{w}\|^2 = \frac{\lambda}{2} \sum_{j=1}^m w_j^2$$

Where λ is the so-called **regularization parameter**. To apply regularization, we add the regularization term to the cost function, which shrinks the weights:

$$J(\mathbf{w}) = \sum_{i=1}^n \left[-y^{(i)} \log(\phi(z^{(i)})) - (1 - y^{(i)}) \log(1 - \phi(z^{(i)})) \right] + \frac{\lambda}{2} \|\mathbf{w}\|^2$$



- We control how well we fit the training data via the regularization parameter λ
- By increasing λ , we increase the strength of regularization
- Sometimes (e.g. in scikit-learn), Support Vector Machine (SVM) terminology is used

$$C = \frac{1}{\lambda}$$

- I.e. we rewrite the regularized cost function of logistic regression as:

$$C \left[\sum_{i=1}^n \left(-y^{(i)} \log(\phi(z^{(i)})) - (1 - y^{(i)}) \log(1 - \phi(z^{(i)})) \right) \right] + \frac{1}{2} \|\mathbf{w}\|^2$$

ML2 – the basics

Measuring success

Overfitting, bias and variance

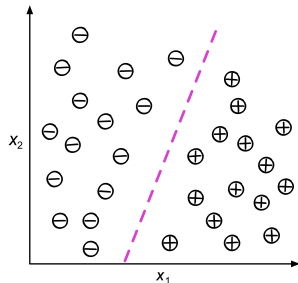
Fixing overfitting

Classification: Support Vector Machines (SVM)

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Recap:

- Predict categorical class labels based on past observations
- Class labels are discrete unordered values which cannot be ordered.
- That is, each element of Y represents a class label and output Y consists of a discrete set of outcomes
- Examples
 - **Binary**: Email spam classification example or unemployment status
 - **Multi-class**: Handwritten digit classification example



- In a p -dimensional space, it's a flat affine subspace of dimension $p-1$.
 - Example: in a **2-dimensional space**, it's a 1-dimensional **line**.
 - Example: in a 3-dimensional space, it's a 2-dimensional plane.
- Formal definition:
 - For a 2-dimensional hyperplane: $\beta_0 + \beta_1 X_1 + \beta_2 X_2 = 0$
 - For a p -dimensional hyperplane: $\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_p X_p = 0$
 - Any point that satisfies the equation lies on the hyperplane.
- Suppose that X instead satisfies
$$\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_p X_p > 0 \rightarrow \text{lies "above" the hyperplane}$$
$$\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_p X_p < 0 \rightarrow \text{lies "below" the hyperplane}$$

- A **separating hyperplane** then has the properties:

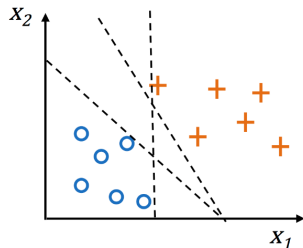
$$\beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots + \beta_p x_{ip} > 0, \text{ if } y_i = 1$$

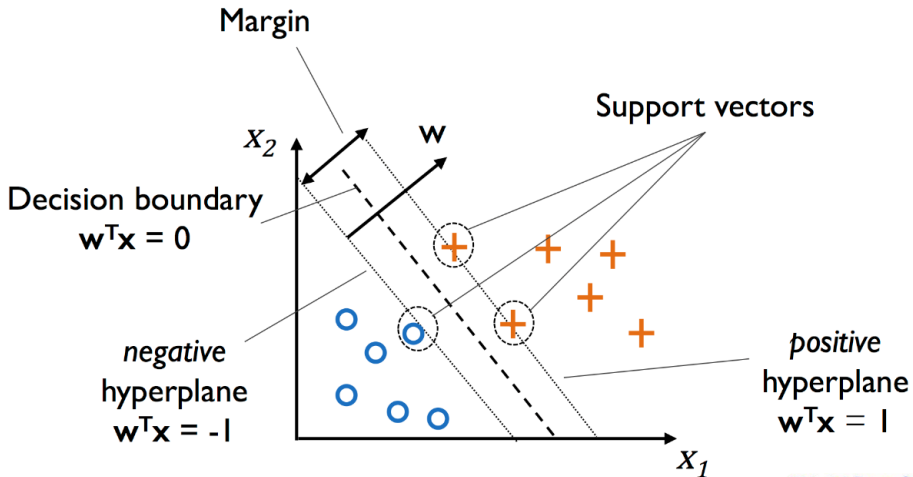
$$\beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots + \beta_p x_{ip} < 0, \text{ if } y_i = -1$$

- Or, put differently:

$$y_i (\beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \dots + \beta_p x_{ip}) > 0 \text{ for all } i$$

- This gives an **infinite number of planes**
- But which plane is the optimal one?
→ **maximum margin classification**



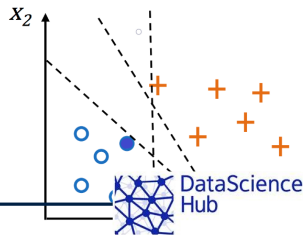
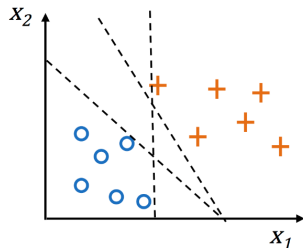


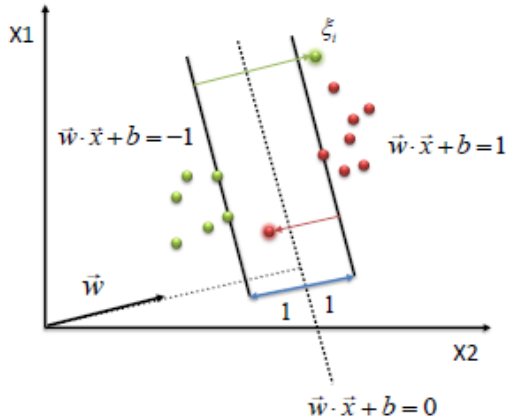
- In SVMs, the optimization objective is to maximize the **margin**
- The margin is defined as the distance between the separating hyperplane and the training samples that are closest to this hyperplane (**support vectors**)
- Intuitively, the **larger the margin**, the **lower generalization error** (variance)
- Models with small margin are prone to overfitting

See “How do Support Vector Machines work?” (Brandon Rohrer) in [Knowledge clips](#).

1. It is very sensitive to small changes in the training set, especially the supports
 - What if the blue dot is removed?
2. It assumes that a separating hyperplane exists

Solution: introduce a **slack variable** ξ_i that allows some instances to 'cross' the margin but then penalize this





slack variable:

$$\xi_i$$

Allow some instances to fall off the margin, but penalize them

- To ensure convergence in presence of misclassifications we need to relax the linear constraints
- Introduce **slack variables** ξ

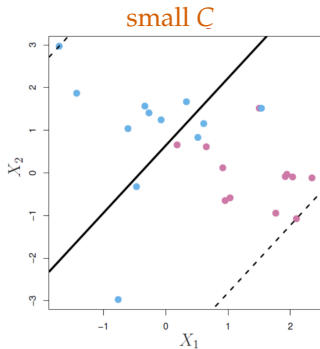
$$\mathbf{w}^T \mathbf{x}^{(i)} \geq 1 - \xi^{(i)} \text{ if } y^{(i)} = 1$$

$$\mathbf{w}^T \mathbf{x}^{(i)} < -1 + \xi^{(i)} \text{ if } y^{(i)} = -1$$

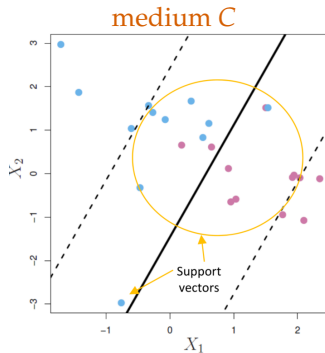
- New objective to be minimized: $\frac{1}{2} \|\mathbf{w}\|^2 + C \left(\sum_i \xi^{(i)} \right)$
- C controls width of the margin with **large values of $C \equiv$ large error penalties**
- C is a way to do **regularization** in SVMs

What is the effect of different values of C ?

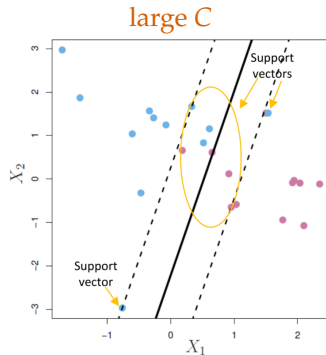
35



- All observations are support vectors
- High bias, low variance

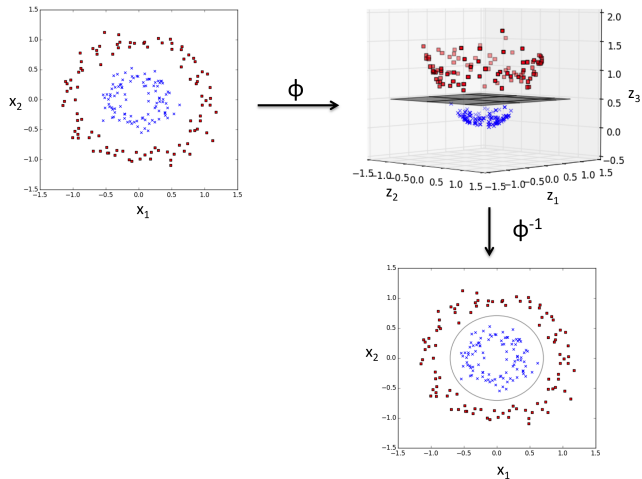


- Most observations are support vectors here



- Few support vectors
- Low bias, high variance

- With non-linear relationships, a support vector classifier is problematic
- One way to add non-linearity is to add higher order terms (x_i^2, x_i^3, \dots) into the separating hyperplane, but this is often computationally impractical.
- A popular alternative is using a **kernel**
 - **linear** kernel $K(x_i, x_{i'}) = \sum_{j=1}^p x_{ij} x_{i'j}$
 - **polynomial** kernel $K(x_i, x_{i'}) = \left(1 + \sum_{j=1}^p x_{ij} x_{i'j}\right)^d$
 - **radial** kernel $K(x_i, x_{i'}) = \exp\left(-\gamma \sum_{j=1}^p (x_{ij} - x_{i'j})^2\right)$
- General blueprint:
 - Transform training data into a higher dimensional space via a mapping function $\phi(\cdot)$
 - Train a linear SVM to classify the data in the new feature space
 - Use the same mapping function $\phi(\cdot)$ to transform new (unseen) data
 - Classify unseen data using the linear SVM model



- A separating hyperplane approach **does not work for multiple classes**

Workarounds:

1. **One-versus-One** classification approach

- Take all possible pairs, compare them one-versus-one, and create several classifiers. e.g., if 3 classes (A,B,C), then compare A with B, A with C, and B with C
- Then, for a given test observation, run it through all K classifiers and tally the number of times it has been assigned to any of the classes
- Assign it the class to which it was predicted to belong the most often

2. **One-versus-All** classification approach

- Fit K SVMs, each time comparing one of the K classes to the remaining $K-1$ classes (treated as if they were one combined class)
- For a given test observation, run it through all K classifiers and assign it to the class for which it was most frequently predicted

ML2 – the basics

- Measuring success

- Overfitting, bias and variance

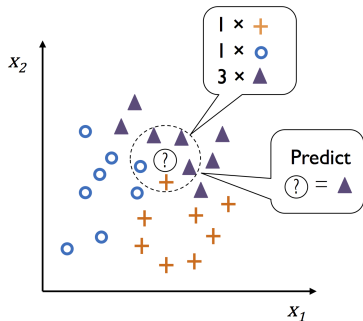
- Fixing overfitting

- Classification: Support Vector Machines (SVM)

- K-nearest neighbors (KNN)



- KNN is an example of a **non-parametric model**
 - Parametric models learn parameters from training data
 - Once training done, the training set not required
- KNN is an **instance-based** or **lazy** learner
 - so needs all the data, all the time
- Basic KNN algorithm
 1. Choose k and a **distance metric**
 2. Find k nearest neighbors of the sample to be classified
 3. Assign the class label by **majority vote**



- Distance metrics:

$$d(\mathbf{x}^{(i)}, \mathbf{x}^{(j)}) = \sqrt[p]{\sum_k |x_k^{(i)} - x_k^{(j)}|^p}$$

Euclidean distance if we set the parameter $p = 2$

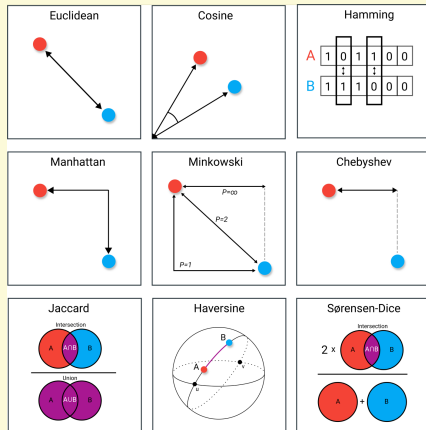
Manhattan distance if we set the parameter $p = 1$

(cf. Hamming distance and Minkowski distance)

- Classifier immediately adapts as we receive new training examples ...
- ... so computational complexity grows linearly with the number of samples
- Need for efficient data structures such as **KD-trees**

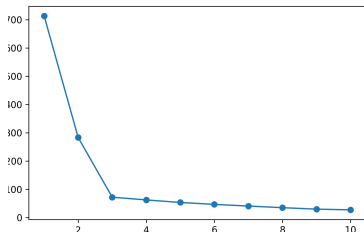
What is the right distance?

42



Source: [Maarten Grootendorst on Medium](#)

- As you've seen before, there is no easy way to answer this.
- Increasing k will lead to high bias but low variance etc. etc.
- Playing around with k will show you the best performance in your model
- Maybe an **elbow graph** can be of use
 - Pick the point where increasing k does not improve the model much
 - See a similar discussion for K -means



Number of neighbors

Visually similar images



- Credit ratings, financial institutes will predict the credit rating of customers
- In loan disbursement, banks predict whether the loan is safe or risky.
- In a regression framework: KNN can be used to predict financial time series
 - stock markets, FX rates, etc., etc.

In this lecture we covered:

1. The balance between **bias** and **variance**
2. Looked at various measures of success in classification
 - **Confusion matrix, Accuracy, Precision, Recall**
3. Discussed how **Support Vector Machines** and **K Nearest Neighbours** work

$$\begin{aligned}\frac{\partial J}{\partial w_j} &= \frac{\partial}{\partial w_j} \frac{1}{2} \sum_i \left(y^{(i)} - \phi(z^{(i)}) \right)^2 \\&= \frac{1}{2} \frac{\partial}{\partial w_j} \sum_i \left(y^{(i)} - \phi(z^{(i)}) \right)^2 \\&= \frac{1}{2} \sum_i 2(y^{(i)} - \phi(z^{(i)})) \frac{\partial}{\partial w_j} (y^{(i)} - \phi(z^{(i)})) \\&= \sum_i (y^{(i)} - \phi(z^{(i)})) \frac{\partial}{\partial w_j} \left(y^{(i)} - \sum_i (w_j^{(i)} x_j^{(i)}) \right) \\&= \sum_i \left(y^{(i)} - \phi(z^{(i)}) \right) \left(-x_j^{(i)} \right) \\&= - \sum_i \left(y^{(i)} - \phi(z^{(i)}) \right) x_j^{(i)}\end{aligned}$$

Calculate the partial derivative of the log-likelihood function with respect to the j th weight:

$$\frac{\partial}{\partial w_j} l(\mathbf{w}) = \left(y \frac{1}{\phi(z)} - (1 - y) \frac{1}{1 - \phi(z)} \right) \frac{\partial}{\partial w_j} \phi(z)$$

Partial derivative of the sigmoid function:

$$\begin{aligned} \frac{\partial}{\partial z} \phi(z) &= \frac{\partial}{\partial z} \frac{1}{1 + e^{-z}} = \frac{1}{(1 + e^{-z})^2} e^{-z} = \frac{1}{1 + e^{-z}} \left(1 - \frac{1}{1 + e^{-z}} \right) \\ &= \phi(z)(1 - \phi(z)). \end{aligned}$$

Resubstitute $\frac{\partial}{\partial z}\phi(z) = \phi(z)(1 - \phi(z))$ to obtain:

$$\begin{aligned} & \left(y \frac{1}{\phi(z)} - (1 - y) \frac{1}{1 - \phi(z)} \right) \frac{\partial}{\partial w_j} \phi(z) \\ &= \left(y \frac{1}{\phi(z)} - (1 - y) \frac{1}{1 - \phi(z)} \right) \phi(z)(1 - \phi(z)) \frac{\partial}{\partial w_j} z \\ &= \left(y(1 - \phi(z)) - (1 - y)\phi(z) \right) x_j \\ &= (y - \phi(z))x_j \end{aligned}$$



Operskalski, J. T., & Barbey, A. K. (2016). Risk literacy in medical decision-making. Science, 352(6284), 413–414.

