k-Nearest Neighbor Learning

DS2500: Intermediate Programming with Data

What is machine learning?

- Can we really make our machines (computers) learn?
- "Secret sauce" is data, and lots of it
- Rather than programming expertise into our applications, we program them to learn from data
- Build working machine-learning models then use them to make remarkably accurate predictions

Prediction

- Improve weather forecasting to save lives, minimize injuries and property damage
- Improve cancer diagnoses and treatment regimens to save lives
- Improve business forecasts to maximize profits and secure people's jobs
- Detect fraudulent credit-card purchases and insurance claims
- Predict customer "churn", what prices houses are likely to sell for, ticket sales of new movies, and anticipated revenue of new products and services
- Predict the best strategies for coaches and players to use to win more games and championships
- All of these kinds of predictions are happening today with machine learning.



Machine Learning Approaches

Classification

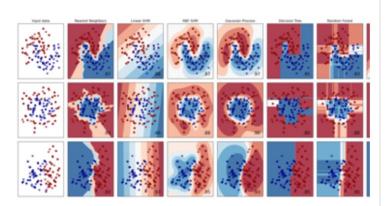
Identifying which category an object belongs to.

Applications: Spam detection, image

recognition.

Algorithms: SVM, nearest neighbors,

random forest, and more...



Regression

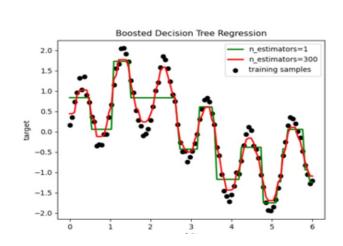
Predicting a continuous-valued attribute associated with an object.

Applications: Drug response, Stock

prices.

Algorithms: SVR, nearest neighbors,

random forest, and more...



Clustering

Automatic grouping of similar objects into sets.

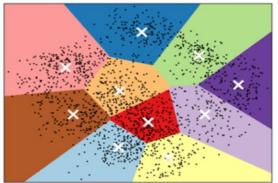
Applications: Customer segmentation,

Grouping experiment outcomes

Algorithms: k-Means, spectral cluster-

ing, mean-shift, and more...







Popular Machine Learning Applications

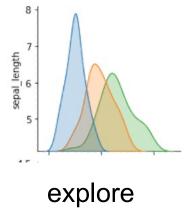
| Anomaly detection | Data mining social media (like Facebook, Twitter, LinkedIn) | Predict mortgage loan defaults |
|--|--|--|
| Chatbots | Detecting objects in scenes | Natural language translation (English to Spanish, French to Japanese, etc.) |
| Classifying emails as spam or not spam | Detecting patterns in data | Recommender systems ("people who bought this product also bought") |
| Classifying news articles as sports, financial, politics, etc. | Diagnostic medicine | Self-Driving cars (more generally, autonomous vehicles) |
| Computer vision and image classification | Facial recognition | Sentiment analysis (like classifying movie reviews as positive, negative or neutral) |
| Credit-card fraud detection | Handwriting recognition | Spam filtering |
| Customer churn prediction | Insurance fraud detection | Time series predictions like stock-price forecasting and weather forecasting |
| Data compression | Intrusion detection in computer networks | Voice recognition |
| Data exploration | Marketing: Divide customers into clusters | |
| | | |

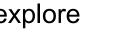


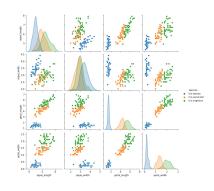
Machine Learning Recipe

5.1,3.8,1.6,0.2,Iris-setosa 4.6,3.2,1.4,0.2,Iris-setosa 5.3,3.7,1.5,0.2,Iris-setosa 5.0,3.3,1.4,0.2,Iris-setosa 7.0,3.2,4.7,1.4,Iris-versicolor 6.4,3.2,4.5,1.5,Iris-versicolor 6.9,3.1,4.9,1.5,Iris-versicolor 5.5,2.3,4.0,1.3, Iris-versicolor

> load data



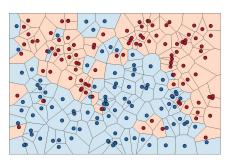




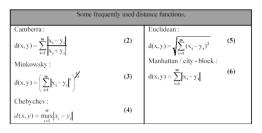
transform



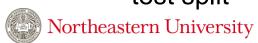
training & test split



model building

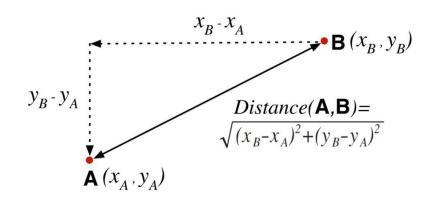


model tuning



Euclidean Distance

| Attribute | Person A | Person B |
|---|----------|----------|
| Age | 23 | 40 |
| Years at current address | 2 | 10 |
| Residential status (1=Owner, 2=Renter, 3=Other) | 2 | 1 |



$$d(A, B) = \sqrt{(23 - 40)^2 + (2 - 10)^2 + (2 - 1)^2}$$

\$\approx 18.8\$

Distance metrics

Some frequently used distance functions.

Camberra:

$$d(x, y) = \sum_{i=1}^{m} \frac{x_i - y_i}{x_i + y_i}$$

$$d(x, y) = \sum_{i=1}^{m} \frac{|x_i - y_i|}{|x_i + y_i|}$$
Minkowsky:
$$d(x, y) = \left(\sum_{i=1}^{m} |x_i - y_i|^r\right)^{r}$$
Chebychev:
$$d(x, y) = \max_{i=1}^{m} |x_i - y_i|$$

$$d(x, y) = \max_{i=1}^{m} |x_i - y_i|$$

(4)

Euclidean:

(2)
$$d(x,y) = \sqrt{\sum_{i=1}^{m} (x_i - y_i)^2}$$
 Manhattan / city - block :
$$d(x,y) = \sum_{i=1}^{m} |x_i - y_i|$$
 (6)

$$d(x, y) = \sum_{i=1}^{m} |x_i - y_i|$$
 (6)

Euclidean Distance on Iris Dataset

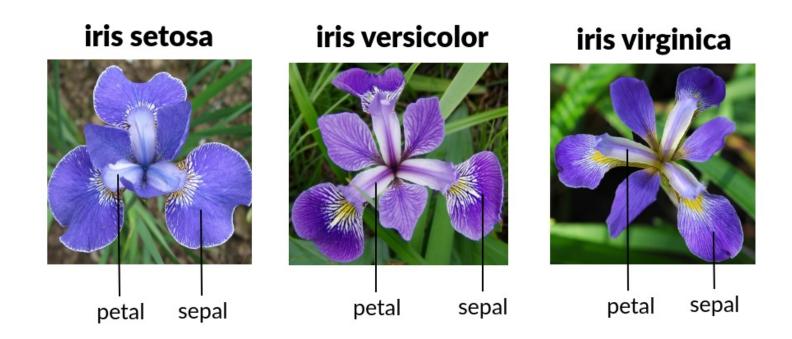
$$diff = \sqrt{(\Delta Slen)^2 + (\Delta Swid)^2 + (\Delta Plen)^2 + (\Delta Pwid)^2}$$

Slen = Sepal Length

Swid = Sepal Width

Plen = Petal Length

Pwid = Petal Width



A more general distance measure

$$diff = w_1 |\Delta A_1|^r + w_2 |\Delta A_2|^r + ... + w_n |\Delta A_n|^r$$

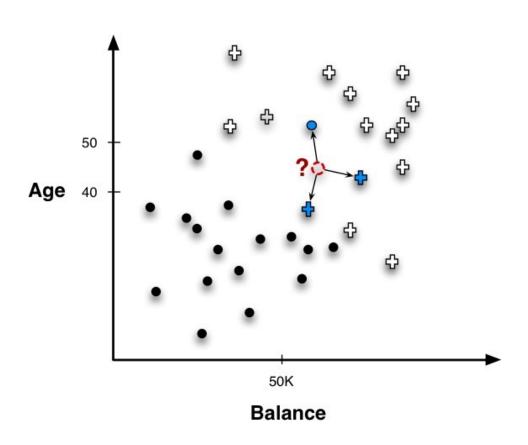
where

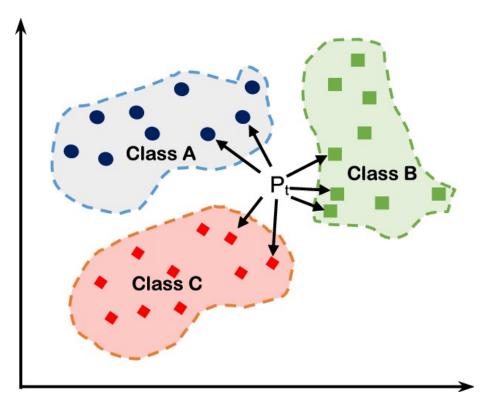
 ΔA_i is the difference with respect to feature i,

 w_k is a feature weighting factor (0.0 to 1.0), and

r is an exponent (> 0.0)

Multiple classes

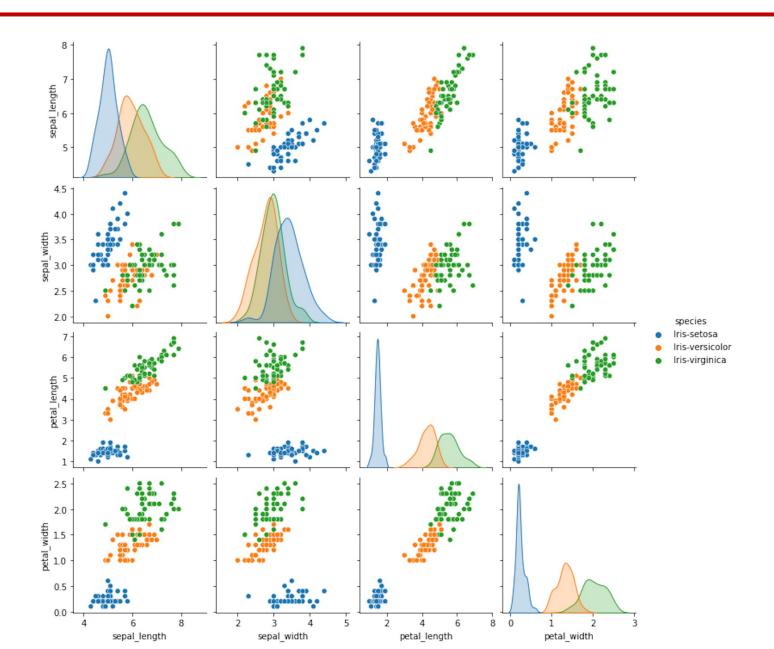




Source: https://laptrinhx.com/k-nearest-neighbors-unlocked-454254569/

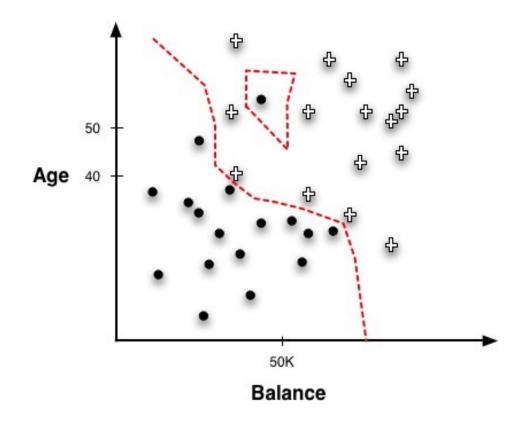


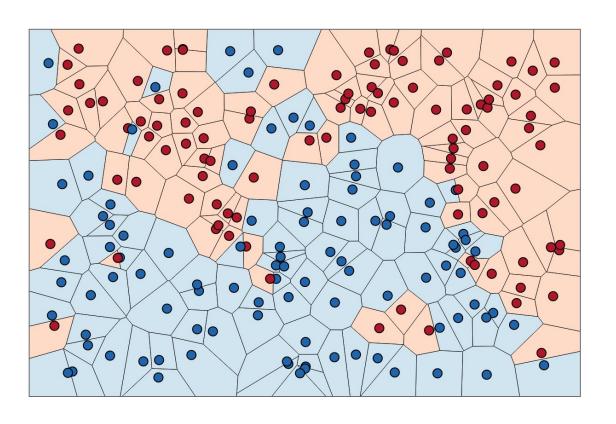
Why 1-NN makes mistakes on the iris dataset





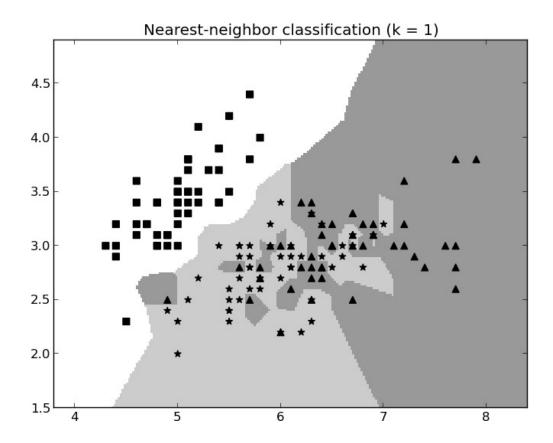
Overfitting

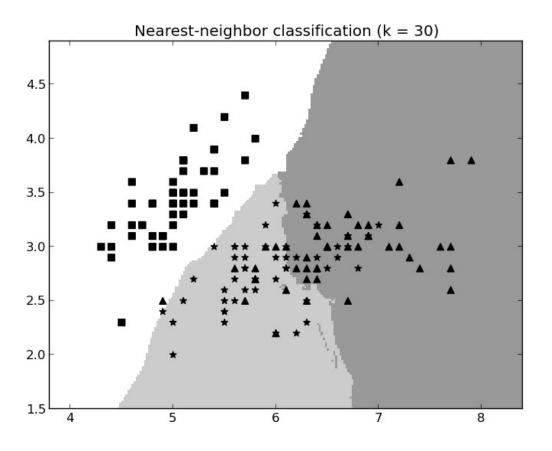






Justifying the k-value





Intelligibility

"The movie Billy Elliot was recommended based on your interest in Amadeus, The Constant Gardener and Little Miss Sunshine"

Amazon: Customers with similar searches purchased....

Amazon: Related to Items You've Viewed.....

We declined your mortgage application because you remind us of the Smiths and the Mitchells, who both defaulted."



Efficiency Issues

Training: very fast (simply store the instances)

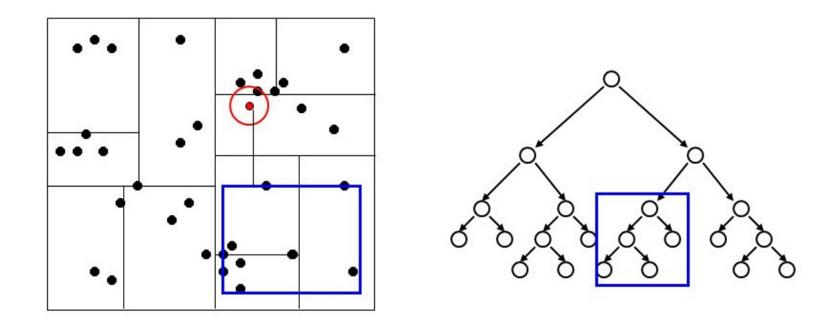
Classification: Finding k nearest neighbors requires computing distance to each instance, and sorting \rightarrow **O**(n log n) for each instance.

Reduced to $O(kn) \sim O(n)$ if k is small using selection by partial sorting.

```
function select(list[1..n], k)
  for i from 1 to k
    minIndex = i
    minValue = list[i]
    for j from i+1 to n do
        if list[j] < minValue then
            minIndex = j
            minValue = list[j]
        swap list[i] and list[minIndex]
  return list[k]</pre>
```



KD-Trees



Using the distance bounds and the bounds of the data below each node, we can prune parts of the tree that could NOT include the nearest neighbor.



Heterogeneous Attributes

| Attribute | Person A | Person B |
|---|----------|----------|
| Sex | Male | Female |
| Age | 23 | 40 |
| Years at current address | 2 | 10 |
| Residential status (1=Owner, 2=Renter, 3=Other) | 2 | 1 |
| Income | 50,000 | 90,000 |

Handling categorical attributes:

- 1. Ignore?
- 2. Convert to numbers?
- 3. Cosine similarity?

The k-NN algorithm

Algorithm:

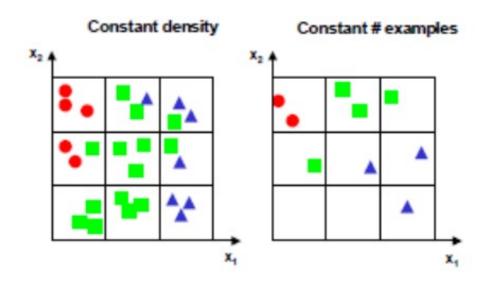
Suppose you have a training data of size D with d samples, and t is a new test sample.

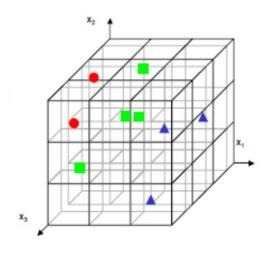
- 1. Select a value of K, which is the value of the nearest neighbors to be used in the computation of the algorithm.
- 2. For i=0 to d; find the distance of the test point from every point in the training data set.
- 3. Make a set S of the K smallest distances.
- 4. Return the majority label from set S.

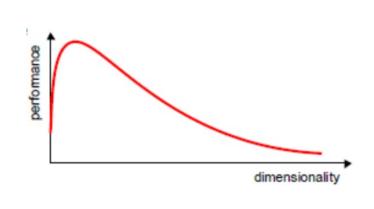


Curse of Dimensionality

Since all attributes contribute to the distance calculations, instance similarity can be confused and misled by the presence of too many irrelevant attributes.









Precision and Recall / Sensitivity and Specificity

10 photos, 7 dogs



Perfect Classification:

| | | True/Actual | | |
|-----------|--------------|--------------|----------|--|
| | | Positive (😭) | Negative | |
| Pred | Positive (😫) | 7 | 0 | |
| Predicted | Negative | 0 | 3 | |

Source: https://towardsdatascience.com/multi-class-metrics-made-simple-part-i-precision-and-recall-9250280bddc2



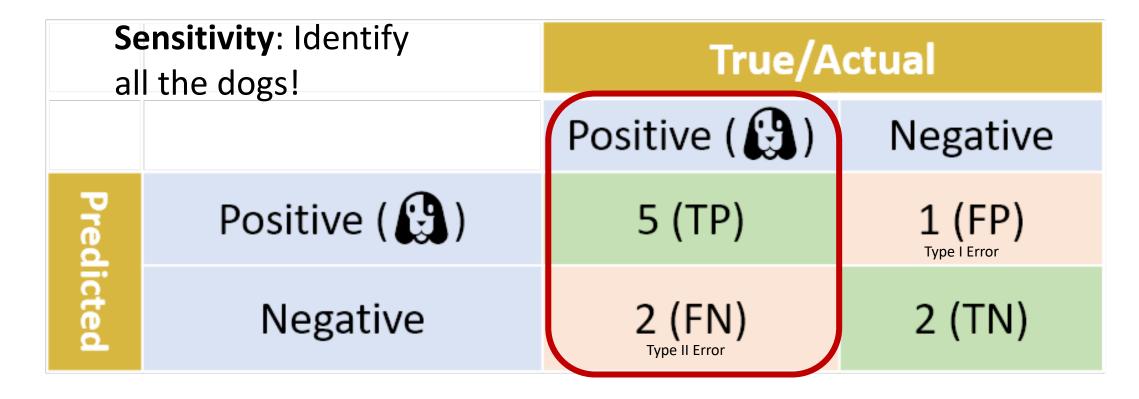
Imperfect Classification

| | | True/Actual | | |
|-----------|--------------|----------------------|---------------------|--|
| | | Positive (😭) | Negative | |
| Pred | Positive (😭) | 5 (TP) | 1 (FP) Type I Error | |
| Predicted | Negative | 2 (FN) Type II Error | 2 (TN) | |

Accuracy: What proportion of photos are correctly classified? Accuracy = (TP + TN) / (TP + TN + FP + FN) = 7 / 10 = 70.0%



Imperfect Classification: Sensitivity



Sensitivity: What proportion of **dogs** (*positives*) are identified? **Sensitivity** = TP / (TP + FN) = TP / T = 5 / (5 + 2) = 71.4% a.k.a. **Recall**



Imperfect Classification: Specificity

| | ecificity: Don't assify cats as dogs! | True/Actual | | |
|-----------|---------------------------------------|----------------------|---------------------|--|
| | • | Positive (😭) | Negative | |
| Pred | Positive (😭) | 5 (TP) | 1 (FP) Type I Error | |
| Predicted | Negative | 2 (FN) Type II Error | 2 (TN) | |

What proportion of the cats (negatives) are identified?

Specificity =
$$TN / (TN + FP) = 2 / (1 + 2) = 66.7\%$$

Specificity = 1 – False Positive Rate



Imperfect Classification

| | recision: Identify ogs | True/Actual | |
|-----------|------------------------|----------------------|---------------------|
| | | Positive (😝) | Negative |
| Pred | Positive (😭) | 5 (TP) | 1 (FP) Type I Error |
| Predicted | Negative | 2 (FN) Type II Error | 2 (TN) |

Precision: What proportion of *predicted dogs (positives)* are truly dogs?

$$Precision = TP / (TP + FP) = TP / P = 5 / (5 + 1) = 83.3\%$$



Multiple classes

| | | True/Actual | | | |
|-----------|------------------|-------------|---|-----------|------------------|
| | | Cat (🐷 |) | Fish (��) | Hen (4) |
| Pr | Cat (🐯) | 4 | | 6 | 3 |
| Predicted | Fish (��) | 1 | | 2 | 0 |
| ed | Hen (﴿) | 1 | | 2 | 6 |

| pr | ecision | recall | f1-score |
|------|---------|--------|----------|
| Cat | 0.308 | 0.667 | 0.421 |
| Fish | 0.667 | 0.200 | 0.308 |
| Hen | 0.667 | 0.667 | 0.667 |

Cat: Precision = 4 / 13 = 0.308, Recall = 4 / 6 = 0.667

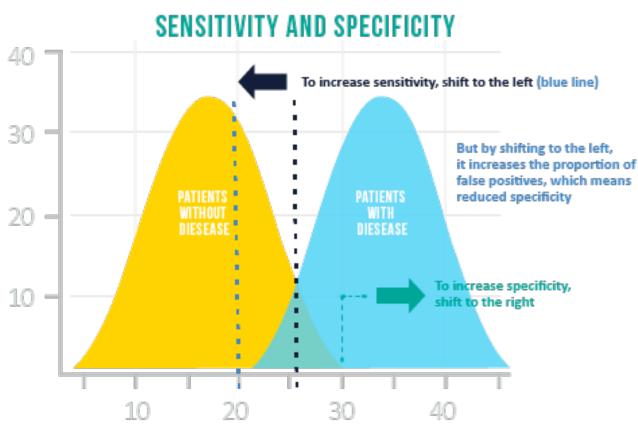
Fish: Precision = 2/3 = 0.667, Recall = 2/10 = 0.200

Hen: Precision = 6/9 = 0.667, Recall = 6/9 = 0.667



Medical Testing

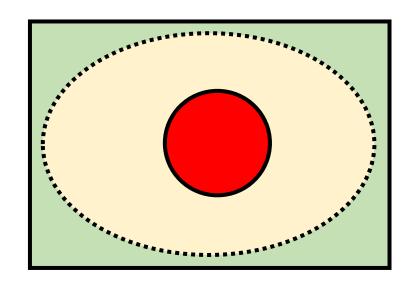


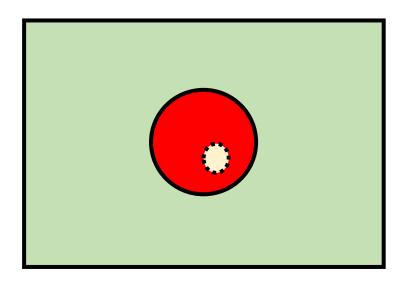


How does this impact patient care and public policy?



Recall/Precision and Sensitivity/Specificity





Perfect recall / sensitivity (TP/P):

Everyone truly sick is detected.

Poor precision (TP/(TP+FP)):

Few of the positives are really sick.

Poor specificity (TN/N):

Many healthy people are test positive.

Poor recall / sensitivity:

Many sick people aren't detected.

Perfect precision:

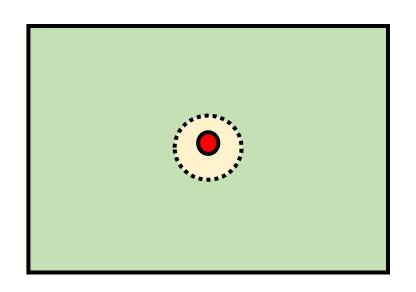
Every positive is truly sick.

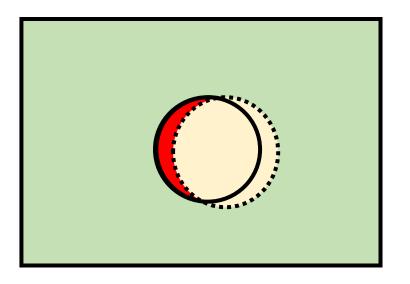
Perfect specificity:

All healthy people test negative.



Recall/Precision and Sensitivity/Specificity





Perfect recall / sensitivity (TP/P):

Everyone truly sick is detected.

Poor precision (TP/(TP+FP)):

Few of the positives are really sick.

High specificity (TN/N):

Most healthy people are test negative.

Good recall / sensitivity:

Most sick people are detected positive.

Good precision:

Most positives are really sick.

Good specificity:

Most healthy people test negative



Precision or Recall? It depends.

Covid-19 Testing:

- → High recall / sensitivity → We detect most patients with Covid (screening)

 High precision → A high proportion of positives are truly sick
- → High specificity → A high proportion of negatives are indeed healthy (diagnosis)

Movie Recommendations:

High recall / sensitivity → We recommend most movies a user would like

→ High precision → Most recommendations are indeed liked

High specificity → Most movies not recommended would not be liked

Output

Description:

Ou

Search Engines / Information Retrieval

High recall / sensitivity → Find all relevant information

★ High precision → Most hits are relevant
High specificity → Most ignored information is not relevant



F1 - Score

Classifier A: precision > recall

Classifier B: recall > precision

Which is better?

The F1-Score is a way of combining precision and recall into a single overall score:

F1-score = $2 \times (precision \times recall)/(precision + recall)$



Combining F1-Scores

| | | True/Actual | | |
|-----------|----------|-------------|-----------|------------------|
| | | Cat (🐯) | Fish (��) | Hen (4) |
| Pr | Cat (👹) | 4 | 6 | 3 |
| Predicted | Fish (¶) | 1 | 2 | 0 |
| ed | Hen (🐴) | 1 | 2 | 6 |

| Class | Precision | Recall | F1-score |
|-------|-----------|--------|----------|
| Cat | 30.8% | 66.7% | 42.1% |
| Fish | 66.7% | 20.0% | 30.8% |
| Hen | 66.7% | 66.7% | 66.7% |

Macro-F1 = (42.1% + 30.8% + 66.7%) / 3 = 46.5%

Weighted-F1 = $(6 \times 42.1\% + 10 \times 30.8\% + 9 \times 66.7\%) / 25 = 46.4\%$



Iris Classification Report

| | Setosa | Virginica | Versicolor |
|------------|--------|-----------|------------|
| Setosa | 50 | 0 | 0 |
| Virginica | 0 | 47 | 3 |
| Versicolor | 0 | 3 | 47 |

| Classification | Report: | | | |
|----------------|-----------|--------|----------|---------|
| | precision | recall | f1-score | support |
| setosa | 1.000 | 1.000 | 1.000 | 50 |
| versicolor | 0.940 | 0.940 | 0.940 | 50 |
| virginica | 0.940 | 0.940 | 0.940 | 50 |
| accuracy | | | 0.960 | 150 |
| macro avg | 0.960 | 0.960 | 0.960 | 150 |
| weighted avg | 0.960 | 0.960 | 0.960 | 150 |



Limitations of F1-Score

Main problem: The F1-Score gives equal weight to Precision and Recall

A more general F score, F_{β} , that uses a positive real factor β , where β is chosen such that recall is considered β times as important as precision, is:

$$F_{eta} = (1 + eta^2) \cdot rac{ ext{precision} \cdot ext{recall}}{(eta^2 \cdot ext{precision}) + ext{recall}}.$$



General Distance Measure

$$diff = w_1 |\Delta A_1|^r + w_2 |\Delta A_2|^r + ... + w_n |\Delta A_n|^r$$

where

 ΔA_i is the difference with respect to feature i,

 w_k is a feature weighting factor (0.0 to 1.0), and

r is an exponent (> 0.0)

Can we evolve an optimal distance metric?

 $diff = w_1 |\Delta A_1|^r + w_2 |\Delta A_2|^r + \dots + w_n |\Delta A_n|^r$

where

 ΔA_i is the difference with respect to feature i,

 w_k is a feature weighting factor (0.0 to 1.0), and

r is an exponent (> 0.0)

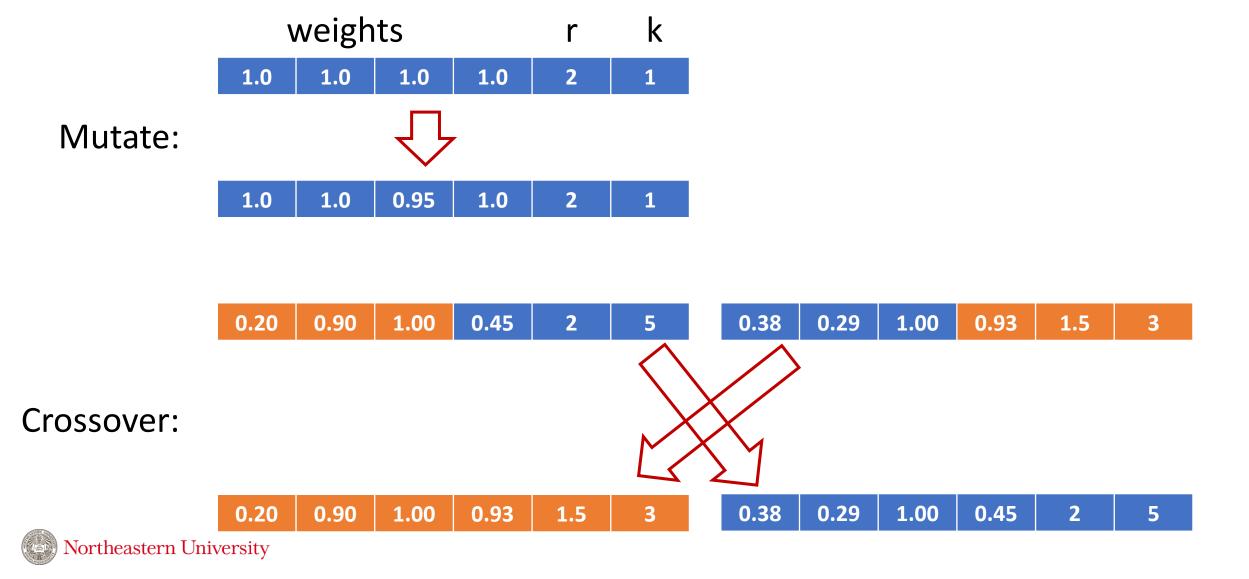
A <u>solution</u> consists of a specification of the distance metric parameters: $w_1...w_n$, r, k (# of nearest neighbors). So n+2 parameters altogether.

We evaluate the solution by testing it against a dataset and finding:

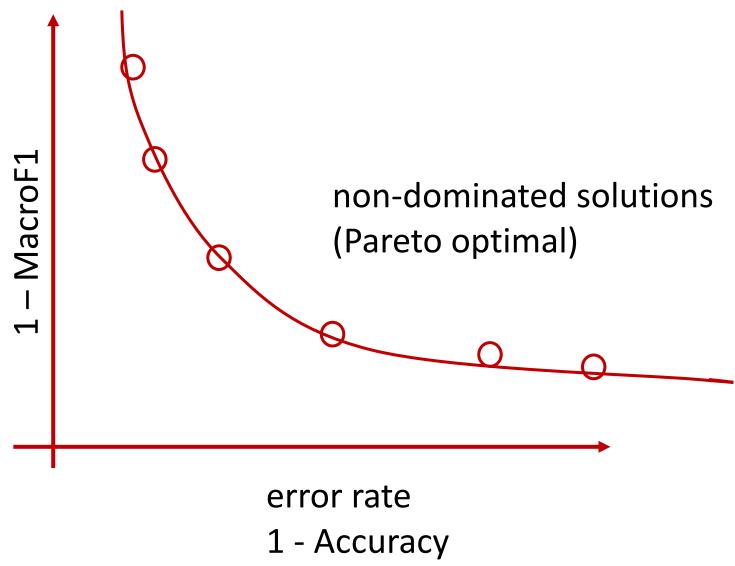
- 1 classification accuracy (error rate)
- 1 macro-F1 / weighted-F1
- # of weights > 0
- 1 macro-precision / weighted-precision?
- 1 macro-recall / weighted-recall?



Tweaking / modifying a solution

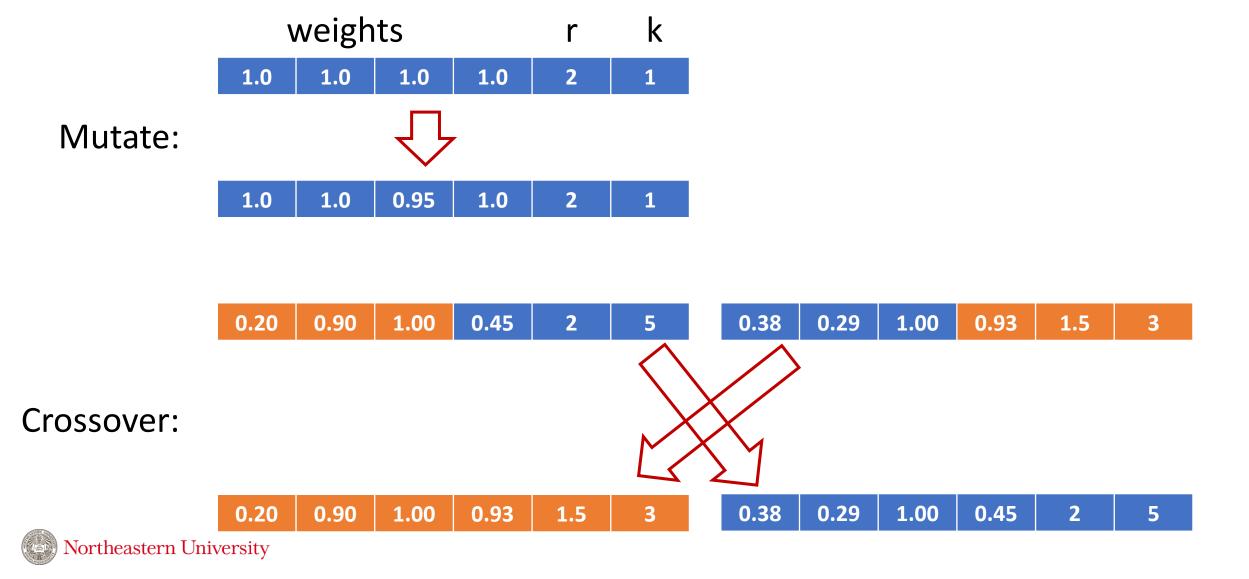


Tradeoffs





Tweaking / modifying a solution

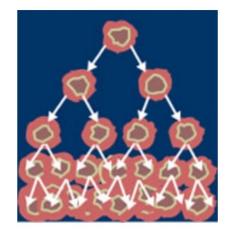


Predicting recurrence of breast cancer



Breast Cancer Wisconsin (Prognostic) Data Set Download: Data Folder, Data Set Description

Abstract: Prognostic Wisconsin Breast Cancer Database

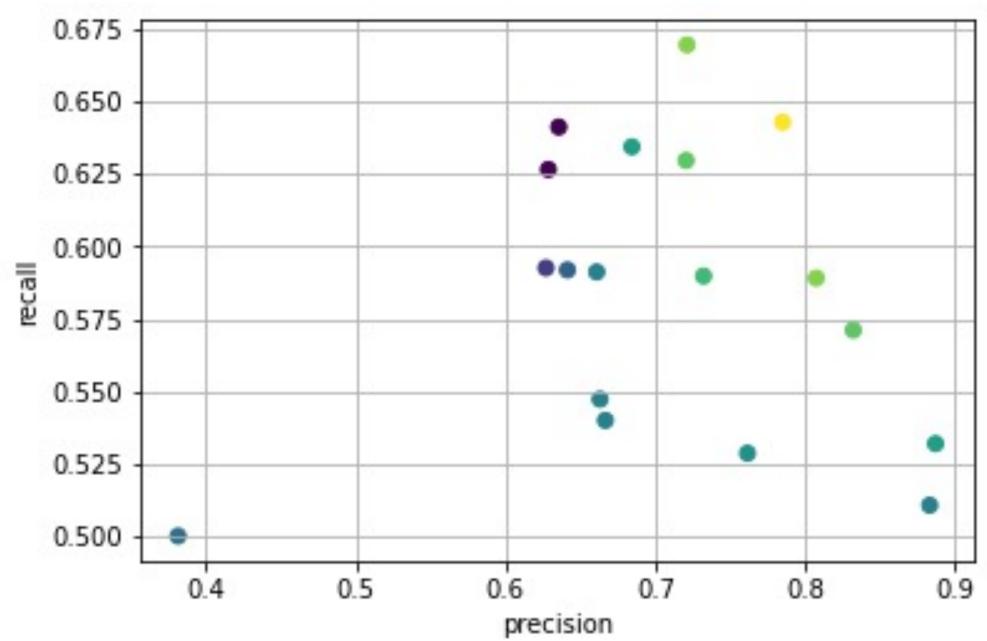


| Data Set Characteristics: | Multivariate | Number of Instances: | 198 | Area: | Life |
|----------------------------|----------------------------|-----------------------|-----|---------------------|------------|
| Attribute Characteristics: | Real | Number of Attributes: | 34 | Date Donated | 1995-12-01 |
| Associated Tasks: | Classification, Regression | Missing Values? | Yes | Number of Web Hits: | 224238 |



```
predicted
              R
        N
actual
   N [[151
              0]
              3]]
       [ 44
                     precision
                                    recall
                                            f1-score
                                                         support
                  N
                       0.77436
                                   1.00000
                                              0.87283
                                                              151
                  R
                       1.00000
                                  0.06383
                                              0.12000
                                                               47
                                              0.77778
                                                              198
          accuracy
                       0.88718
                                  0.53191
                                              0.49642
                                                             198
         macro avg
     weighted avg
                       0.82792
                                  0.77778
                                              0.69413
                                                              198
```

| discarded | accuracy | recall | precision | weights | R | Κ |
|-----------|----------|---------|-----------|---|--------|----|
| 32 | 0.76263 | 0.50000 | 0.38131 | 000000000000000000000000000000000000000 | 2.0085 | 3 |
| 31 | 0.74747 | 0.59264 | 0.62700 | 000000000000000000000000000000000000000 | 2.1085 | 4 |
| 31 | 0.73232 | 0.62667 | 0.62858 | 000000000000000000000000000000000000000 | 2.1085 | 1 |
| 30 | 0.73232 | 0.64133 | 0.63554 | 000000001000000000000000000000000000000 | 1.0264 | 2 |
| 31 | 0.75758 | 0.59194 | 0.64133 | 00000000000000000000001000000000 | 2.0085 | 5 |
| 31 | 0.76768 | 0.59124 | 0.66087 | 00000000001000000000000000000000 | 2.6851 | 5 |
| 31 | 0.76768 | 0.54727 | 0.66310 | 0000000000000010000000000000000 | 2.0085 | 8 |
| 31 | 0.76768 | 0.53995 | 0.66667 | 000000000000000100000000000000000 | 2.0085 | 10 |
| 30 | 0.77778 | 0.63449 | 0.68437 | 000000001000000000000000000000000000000 | 1.0537 | 3 |
| 30 | 0.79293 | 0.62977 | 0.72055 | 000000010000000000000000000000000000000 | 1.1662 | 5 |
| 29 | 0.79798 | 0.66972 | 0.72121 | 0000000010000000000000001000010 | 1.7124 | 3 |
| 30 | 0.78788 | 0.58983 | 0.73224 | 000000001000000000000000000000000000000 | 1.4039 | 7 |
| 31 | 0.77273 | 0.52860 | 0.76160 | 00000000001000000000000000000000 | 2.0223 | 22 |
| 29 | 0.81313 | 0.64302 | 0.78511 | 000001010000000000000000000000000000000 | 1.0179 | 9 |
| 30 | 0.79798 | 0.58912 | 0.80749 | 000000010100000000000000000000000 | 2.9913 | 7 |
| 30 | 0.79293 | 0.57116 | 0.83224 | 000000001000000000000000000000000000000 | 2.7197 | 8 |
| 31 | 0.76768 | 0.51064 | 0.88325 | 0000000000000000010000000000000 | 2.2906 | 17 |
| 30 | 0.77778 | 0.53191 | 0.88718 | 0010000000000000000010000000000 | 2.1623 | 19 |





predicted N R N [[138 13] お R [27 20]]

R=1.7, K=3, Features = 3

| [27 20]] | precision | recall | f1-score | support |
|---------------------------------------|--------------------|--------------------|-------------------------------|-------------------|
| N R | 0.83636 0.60606 | 0.91391 0.42553 | 0.87342 0.50000 | 151 47 |
| accuracy macro avg weighted avg | 0.72121 0.78170 | 0.66972 0.79798 | 0.79798 0.68671 0.78478 | 198 198 198 |

0.675 [[123 [25 28] 0.650 2211 0.625 0.600 0.575 0.550 0.525 0.500 -0.5 0.6 0.7 0.8 0.9 precision

Recall: What fraction of the recurring cancers are we detecting (predicted to recur)? (20 / 47 = 0.42553)

Precision: What fraction of those cancers predicted to recur actually recurred? (20 / 33 = 0.60606)

| actual | N R | predic N [[151 [44 | eted R 0] 3]] | | | | |
|--------|---------------------------|------------------------------|----------------------------|--------------------|--------------------|-------------------------------|-------------------|
| Ø | | | | precision | recall | f1-score | support |
| | | | N R | 0.77436 1.00000 | 1.00000 0.06383 | 0.87283 0.12000 | 151 47 |
| | | | curacy ro avg ed avg | 0.88718 0.82792 | 0.53191 0.77778 | 0.77778 0.49642 0.69413 | 198 198 198 |
| | R=2.3, K=17, Features = 1 | | | | | | |

