

# CPSC 340: Machine Learning and Data Mining

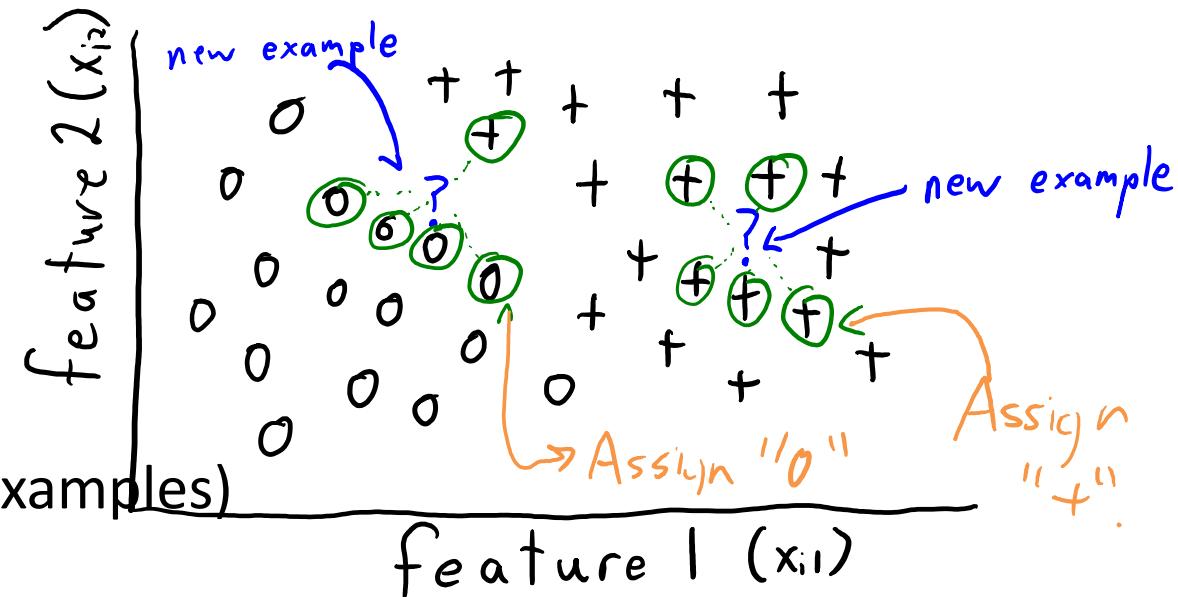
Ensemble Methods  
Spring 2022 (2021W2)

# Admin

- Course webpage:
  - <https://github.com/UBC-CS/cpsc340-2021w1>
- Assignment 2 is out
  - Due Friday of next week. It's long – start early
  - Keep an eye on Piazza and/or commits on the site for updates/fixes
- Midterm
  - **Thursday Feb 17 (6 - 7:30 pm)**
  - Will be online
  - You can take it from anywhere

# Last Time: K-Nearest Neighbours (KNN)

- K-nearest neighbours algorithm for classifying  $\tilde{x}_i$ :
  - Find ‘k’ values of  $x_i$  that are most similar to  $\tilde{x}_i$ .
  - Use mode of corresponding  $y_i$ .
- Lazy learning:
  - To “train” you just store X and y.
- Non-parametric:
  - Size of model grows with ‘n’ (number of examples)
    - Good short article on parametric vs. “non”
  - Nearly-optimal test error with infinite data.
- But high prediction cost and may need large ‘n’ if ‘d’ is large.



# Defining “Distance” with “Norms”

- A common way to define the “distance” between examples:
  - Take the “norm” of the difference between feature vectors.

$$\|x_i - \tilde{x}_i\|_2 = \sqrt{\sum_{j=1}^d (x_{ij} - \tilde{x}_{ij})^2}$$

*train example*      *test example*      "L<sub>2</sub>-norm"

- Norms are a way to measure the “length” of a vector.
  - The most common norm is the “L2-norm” (or “Euclidean norm”):

$$\|r\|_2 = \sqrt{\sum_{j=1}^d r_j^2}$$

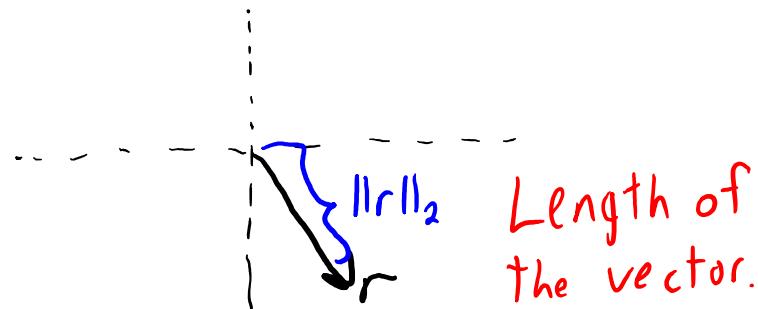
- Here, the “norm” of the difference is the standard Euclidean distance.

# L<sub>2</sub>-norm, L<sub>1</sub>-norm, and L <sub>$\infty$</sub> -Norms.

- The three most common norms: L<sub>2</sub>-norm, L<sub>1</sub>-norm, and L <sub>$\infty$</sub> -norm.
  - Definitions of these norms with two-dimensions:

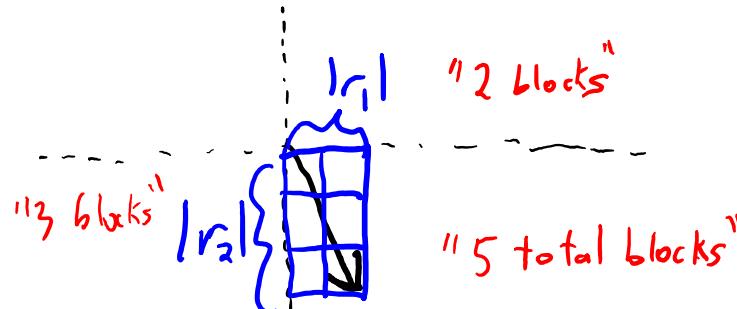
L<sub>2</sub> or "Euclidean" norm.

$$\|r\|_2 = \sqrt{r_1^2 + r_2^2}$$



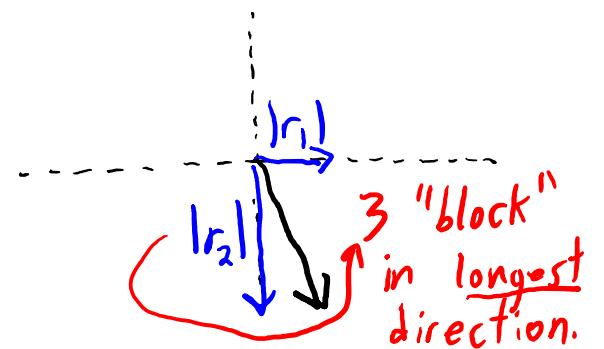
L<sub>1</sub> or "Manhattan" norm:

$$\|r\|_1 = |r_1| + |r_2|$$



L <sub>$\infty$</sub>  or "max" norm:

$$\|r\|_\infty = \max\{|r_1|, |r_2|\}$$



- Definitions of these norms in d-dimensions.

$$L_2: \|r\|_2 = \sqrt{\sum_{j=1}^d r_j^2}$$

$$L_1: \|r\|_1 = \sum_{j=1}^d |r_j|$$

$$L_\infty: \max_j \{|r_j|\}$$

# Norm and Norm<sup>p</sup> Notation (MEMORIZE)

- Notation:

- We often leave out the “2” for the L2-norm: We use  $\|r\|$  for  $\|r\|_2$
- We use superscripts for raising norms to powers: We use  $\|r\|^2$  for  $(\|r\|)^2$
- You should understand why all of the following quantities are equal:

$$\|r\|^2 = \|r\|_2^2 = (\|r\|_2)^2 = \left( \sqrt{\sum_{j=1}^d r_j^2} \right)^2 = \sum_{j=1}^d r_j^2 = \sum_{j=1}^d r_j r_j = r^T r$$

$$= \langle r, r \rangle$$

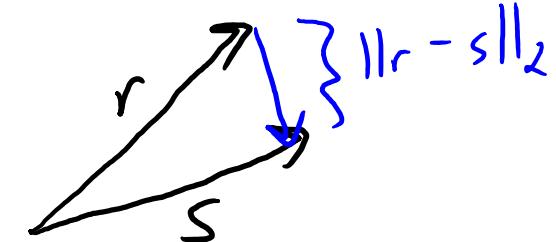
(we'll use  
these later)

# Norms as Measures of Distance

- By taking norm of difference, we get a “distance” between vectors:

$$\|r - s\|_2 = \sqrt{(r_1 - s_1)^2 + (r_2 - s_2)^2}$$

$= \|r - s\|$  "Euclidean distance"



$$\|r - s\|_1 = |r_1 - s_1| + |r_2 - s_2|$$

"Number of blocks you need to walk to get from  $r$  to  $s$ ."

$$\|r - s\|_\infty = \max \{ |r_1 - s_1|, |r_2 - s_2| \}$$

"Most number of blocks in any direction you would have to walk."

- Place different “weights” on large differences:
  - $L_1$ : differences are equally notable.
  - $L_2$ : bigger differences are more important (because of squaring).
  - $L_\infty$ : only biggest difference is important.

bonus!

# KNN Distance Functions

- Most common KNN distance functions:  $\text{norm}(x_i - x_j)$ .

- L1-, L2-, and L $\infty$ -norm.

- Weighted norms (if some features are more important):

$$\sum_{j=1}^d v_j |x_j|$$

$\uparrow$  "weight" of feature  $j$

- “Mahalanobis” distance (takes into account correlations).

- See bonus slide for what functions define a “norm”.

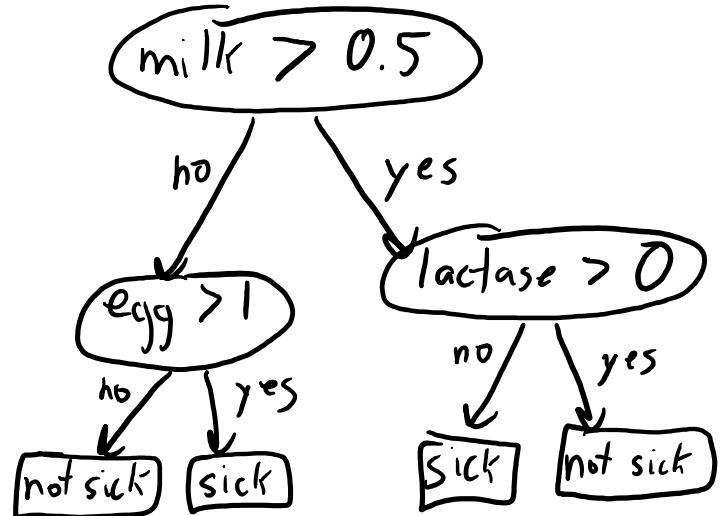
- But we can consider other distance/similarity functions:

- Jaccard similarity (if  $x_i$  are sets).

- Edit distance (if  $x_i$  are strings).

- Metric learning (*learn* the best distance function).

# Decision Trees vs. Naïve Bayes vs. KNN



$$\begin{aligned} p(\text{sick} \mid \text{milk}, \text{egg}, \text{lactase}) \\ \approx p(\text{milk} \mid \text{sick}) p(\text{egg} \mid \text{sick}) p(\text{lactase} \mid \text{sick}) p(\text{sick}) \end{aligned}$$

$(\text{milk} = 0.6, \text{egg} = 2, \text{lactase} = 0, ?)$  is close to  
 $(\text{milk} = 0.7, \text{egg} = 2, \text{lactase} = 0, \text{sick})$  so predict sick.

# Application: Optical Character Recognition

- To scan documents, we want to turn images into characters:
  - “Optical character recognition” (OCR).

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# Application: Optical Character Recognition

- To scan documents, we want to turn images into characters:
  - “Optical character recognition” (OCR).



- Turning this into a supervised learning problem (with 28 by 28 images):

$$X = \begin{bmatrix} (1,1) & (2,1) & (3,1) & \dots & (28,1) & (1,2) & (2,2) & \dots & (14,14) & \dots & (28,28) \end{bmatrix}$$
$$Y = \begin{bmatrix} 3 \\ 6 \\ 0 \\ 9 \end{bmatrix}$$

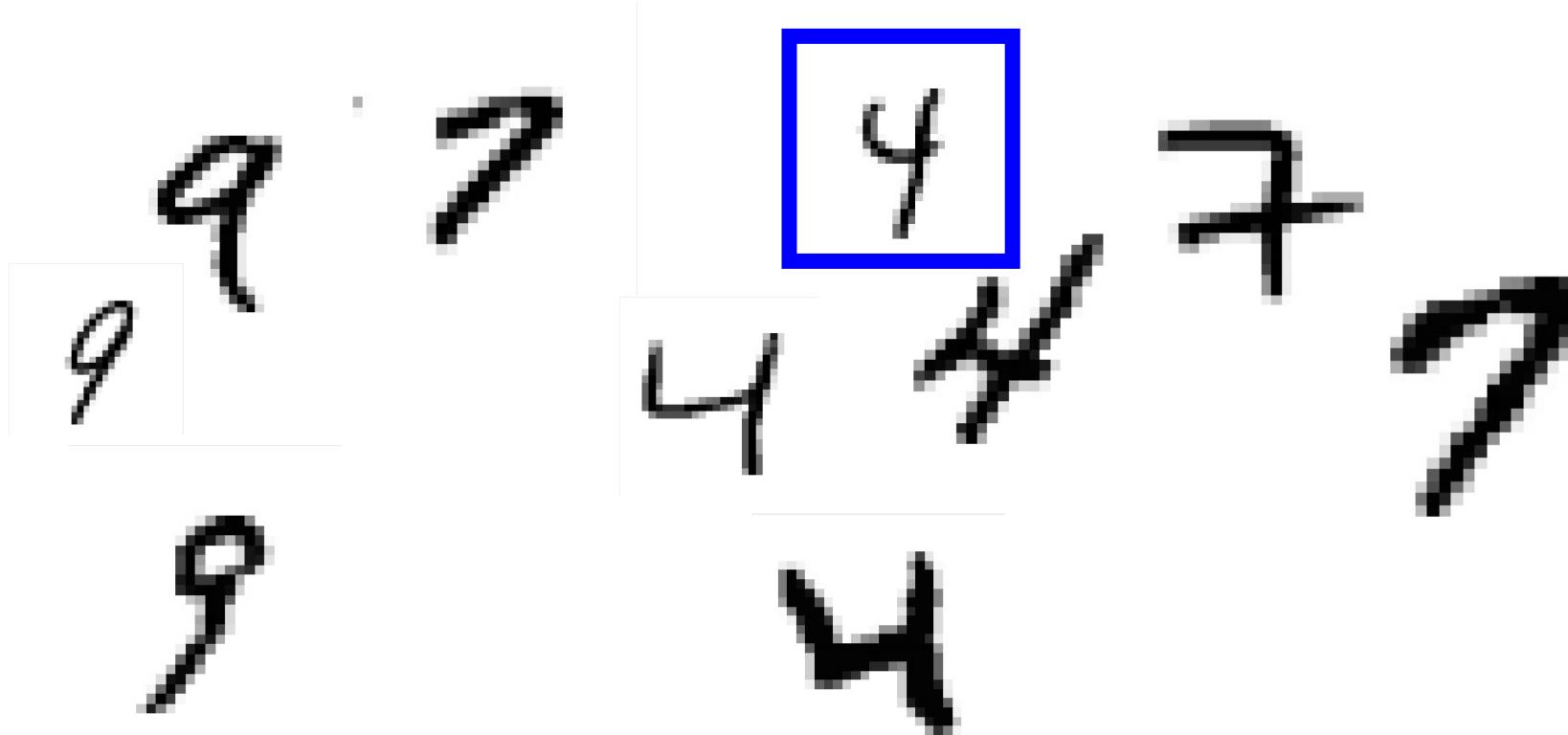
The matrix  $X$  represents a 28x28 image of the digit '3' as a feature vector. The matrix  $Y$  represents the target variable, which is the character '3'.

Each feature is grayscale intensity of one of the 784 pixels

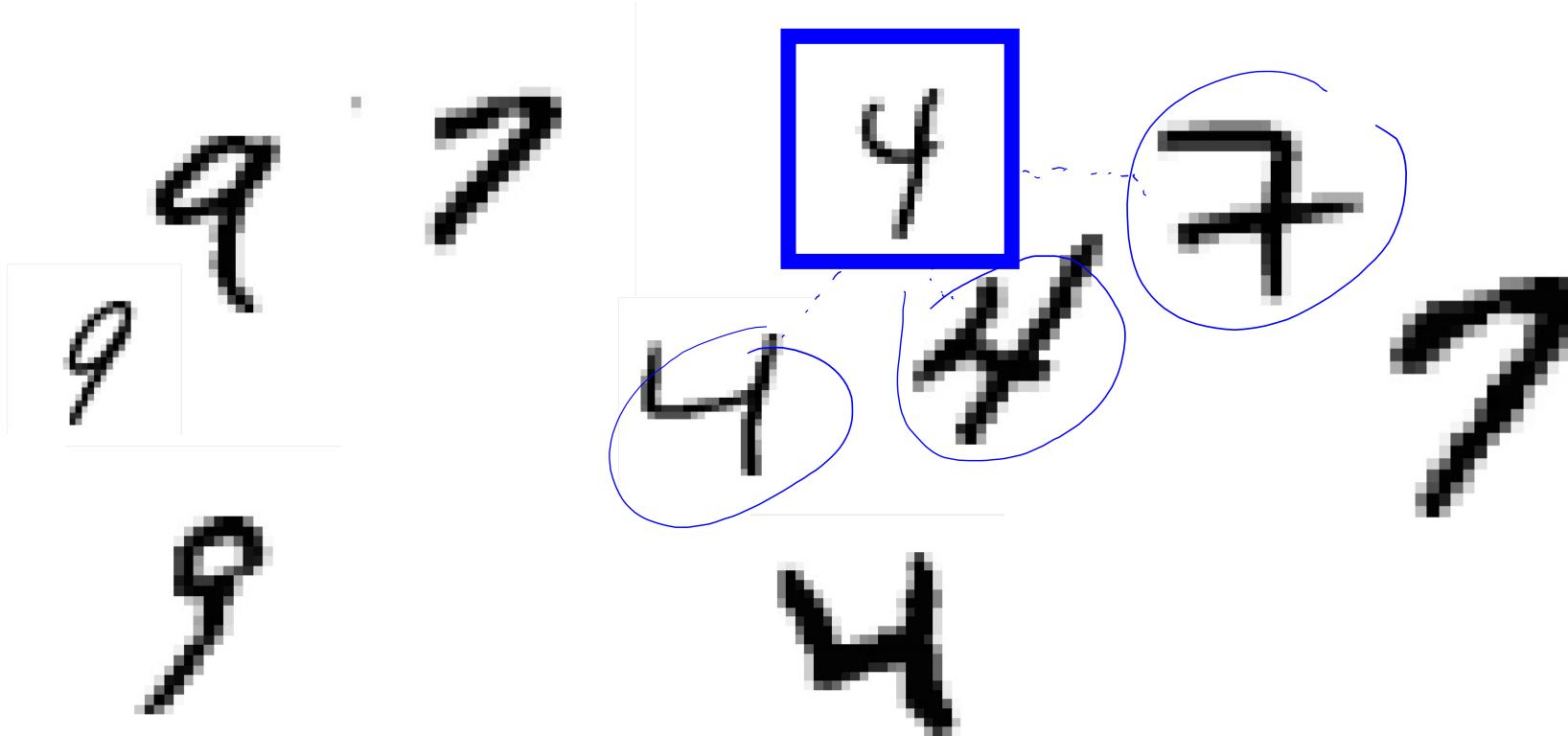
# KNN for Optical Character Recognition



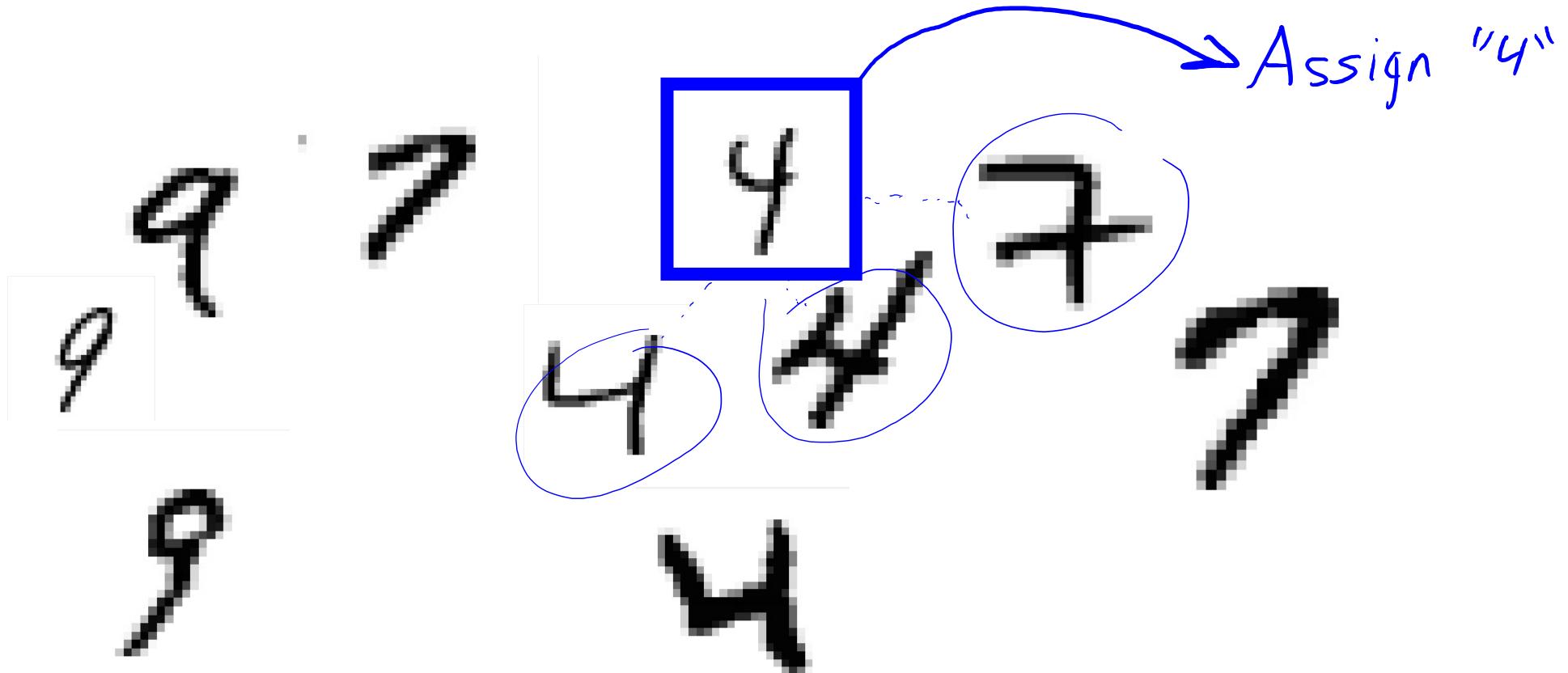
# KNN for Optical Character Recognition



# KNN for Optical Character Recognition



# KNN for Optical Character Recognition



# Human vs. Machine Perception

- There is **huge difference** between what we see and what KNN sees:

What we see:



What the computer “sees”:

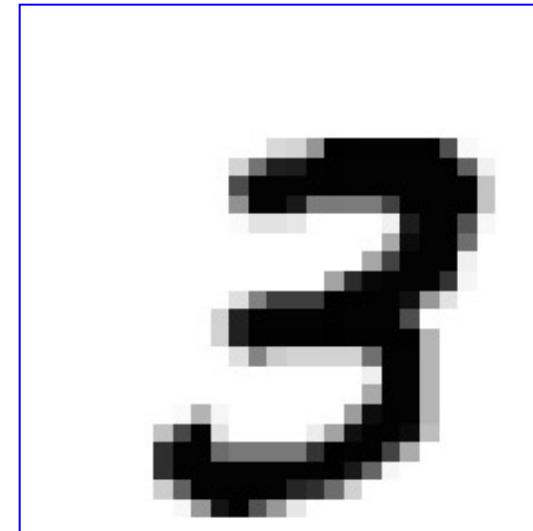
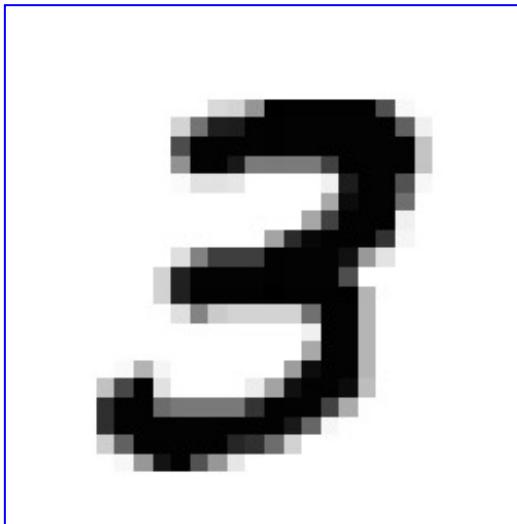


Actually, it's worse:



# What the Computer Sees

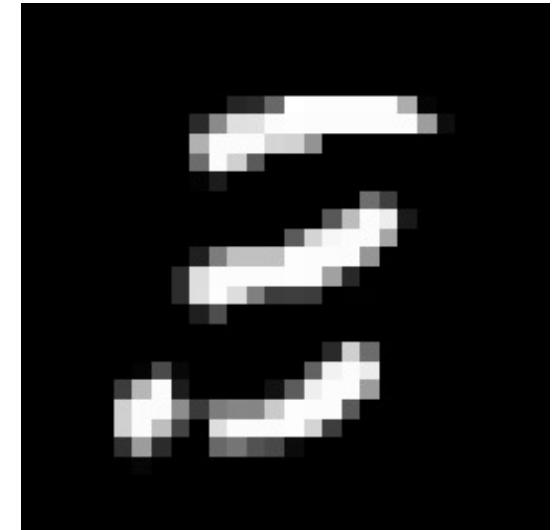
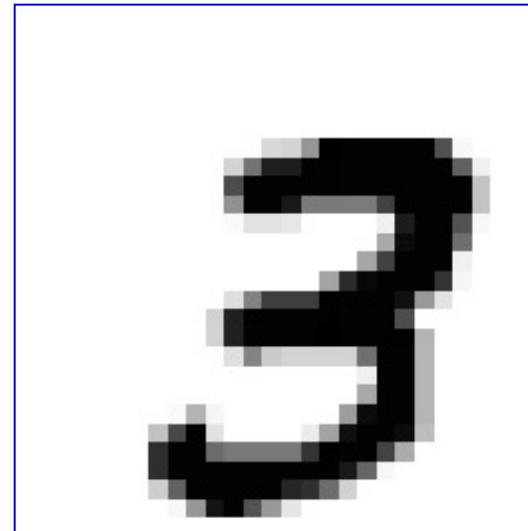
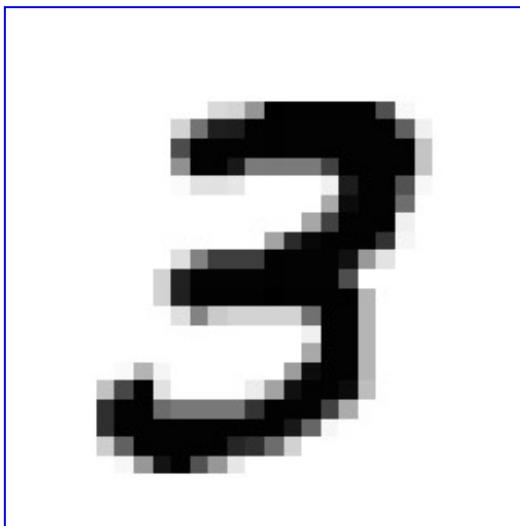
- Are these two images “similar”?



# What the Computer Sees

- Are these two images “similar”?

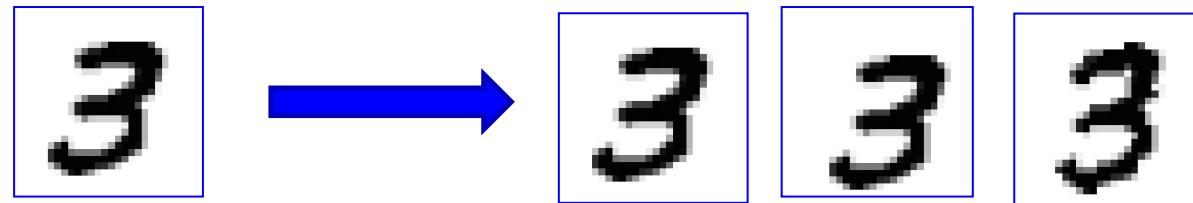
Difference:



- KNN does not know that labels should be translation invariant.

# Encouraging Invariance

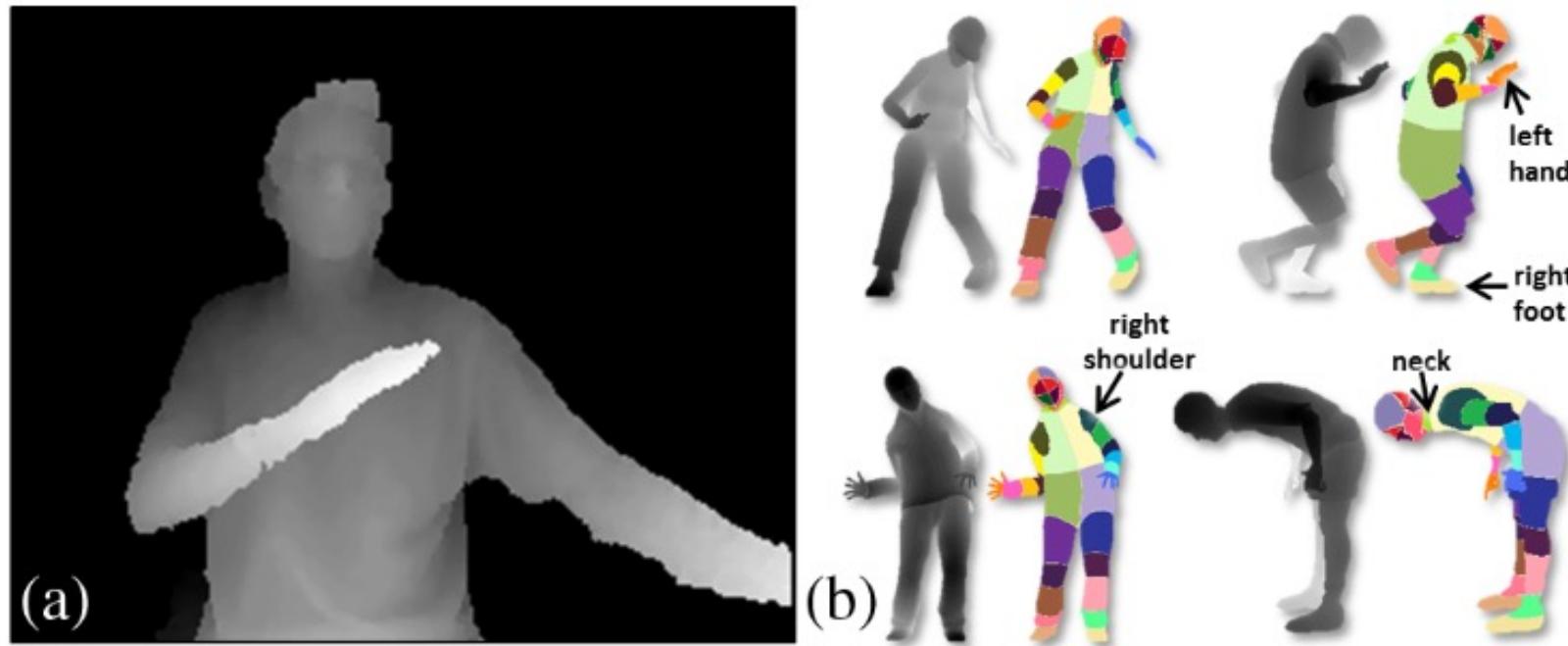
- May want classifier to be invariant to certain feature transforms.
  - Images: translations, small rotations, changes in size, mild warping,...
- The **hard/slow way** is to modify your distance function:
  - Find neighbours that require the “smallest” transformation of image.
- The **easy/fast way** is to just **add transformed data** during training:
  - Add translated/rotate/resized/warped versions of training images.



- “Data augmentation”: crucial part of many successful vision systems.
- Also really important for sound (translate, change volume, and so on).

# Application: Body-Part Recognition

- Microsoft Kinect:
  - Real-time recognition of 31 body parts from laser depth data.



- How could we write a program to do this?

bonus!

# Some Ingredients of Kinect

1. Collect **hundreds of thousands of labeled images** (motion capture).
  - Variety of pose, age, shape, clothing, and crop.
2. Build a **simulator that fills space of images** by making even more images.



3. Extract **features of each location**, that are cheap enough for real-time calculation (depth differences between pixel and pixels nearby.)
4. Treat **classifying body part of a pixel as a supervised learning** problem.
5. Run **classifier in parallel on all pixels** using graphical processing unit (GPU).

# Supervised Learning Step

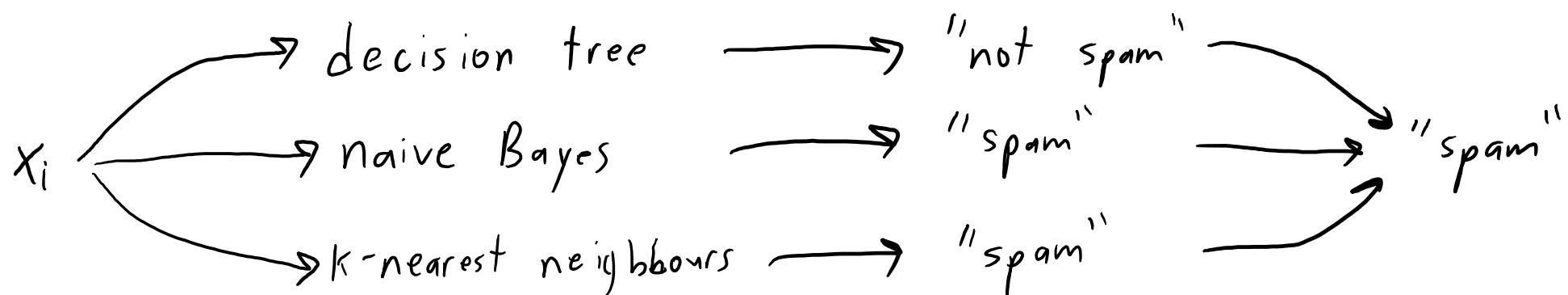
- ALL steps are important, but we'll focus on the learning step.
- Do we have any classifiers that are accurate and run in real time?
  - Decision trees and naïve Bayes are fast, but often not very accurate.
  - KNN is often accurate, but not very fast.
- Deployed system uses an ensemble method called random forests.

# Ensemble Methods

- Ensemble methods are classifiers that combine other classifiers.
- They have the best names:
  - Averaging.
  - Blending.
  - Boosting.
  - Bootstrapping.
  - Bagging.
  - Cascading.
  - Random Forests.
  - Stacking.
  - Voting.
- Ensemble methods often have higher accuracy than input classifiers.

# Ensemble Method Example: Voting

- Ensemble methods use predictions of a set of models.
  - For example, we could use:
    - Decision trees make one prediction.
    - Naïve Bayes makes another prediction.
    - KNN makes another prediction.
- One of the simplest ensemble methods is voting:
  - Take the mode of the predictions across the classifiers.



bonus!

# Why can Voting Work?

- Consider 3 binary classifiers, each **independently correct** with probability 0.80:
- With voting, **ensemble prediction is correct if we have “at least 2 right”**:
  - $P(\text{all 3 right}) = 0.8^3 = 0.512.$
  - $P(2 \text{ rights}, 1 \text{ wrong}) = 3 * 0.8^2(1-0.8) = 0.384.$
  - $P(1 \text{ right}, 2 \text{ wrongs}) = 3 * (1-0.8)^20.8 = 0.096.$
  - $P(\text{all 3 wrong}) = (1-0.8)^3 = 0.008.$
  - So **ensemble is right with probability 0.896** (which is  $0.512+0.384$ ).
- Notes:
  - For voting to work, **errors of classifiers need to be at least somewhat independent**.
  - You also want the probability of being right to be  $> 0.5$ , otherwise it can do much worse.
  - Probabilities also shouldn’t be too different (otherwise, it might be better to take most accurate).

# Why can Voting Work?

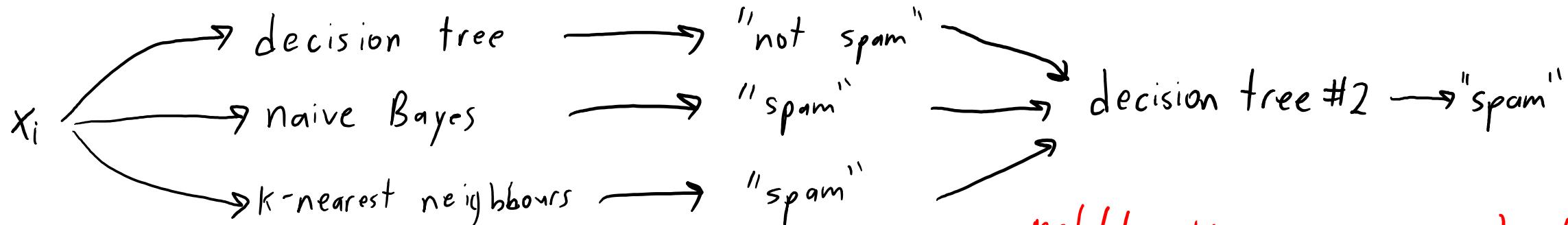
- Why can voting lead to better results?
- Consider classifiers that overfit (like deep decision trees):
  - If they all overfit in exactly the same way, voting does nothing.
- But if they make **independent errors**:
  - Probability that “vote” is wrong can be lower than for each classifier.
  - Less attention to specific overfitting of each classifier.

# Why can Voting Work?

- Consider a set of classifiers that make these predictions:
  - Classifier 1: “spam”.
  - Classifier 2: “spam”.
  - Classifier 3: “spam”.
  - Classifier 4: “not spam”.
  - Classifier 5: “spam”.
  - Classifier 6: “not spam”.
  - Classifier 7: “spam”.
  - Classifier 8: “spam”.
  - Classifier 9: “spam”.
  - Classifier 10: “spam”.
- If these independently get 80% accuracy, mode will be close to 100%.
  - In practice errors won’t be completely independent (due to noise in labels).

# Digression: Stacking

- Another variation on voting is **stacking**
  - Fit another classifier that uses the predictions as features.



- Can tune second classifier using validation data.
  - Sometimes called "**blending**".
- Stacking often performs **better than individual models**.
  - Typically used by Kaggle winners.
  - E.g., Netflix \$1M user-rating competition winner was stacked classifier.

$$X = \begin{bmatrix} \text{model 1} & \text{model 2} & \text{model 3} \\ \text{not spam} & \text{spam} & \text{spam} \\ \text{spam} & \text{spam} & \text{spam} \\ \text{not spam} & \text{not spam} & \text{spam} \\ \vdots & \vdots & \vdots \end{bmatrix}$$
$$y = \begin{bmatrix} \text{true label} \\ \text{spam} \\ \text{spam} \\ \text{not spam} \\ \vdots \end{bmatrix}$$

# Random Forests

- Random forests take vote from a set of deep decision trees.
  - Tend to be one of the best “out of the box” classifiers.
    - Often close to the best performance of any method on the first run.
  - And predictions are very fast.
- Do deep decision trees make independent errors?
  - No: with the same training data you’ll get the same decision tree.
- Two key ingredients in random forests:
  - Bootstrapping.
  - Random trees.

# Bootstrap Sampling

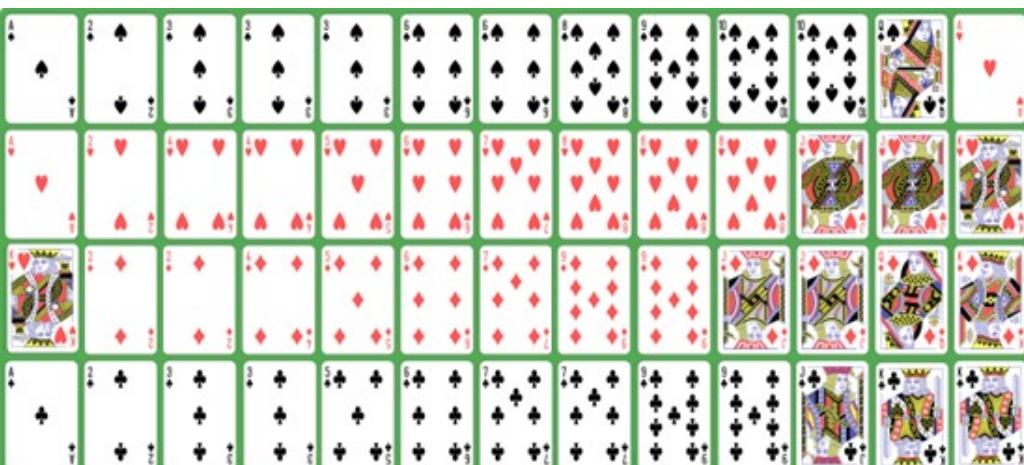
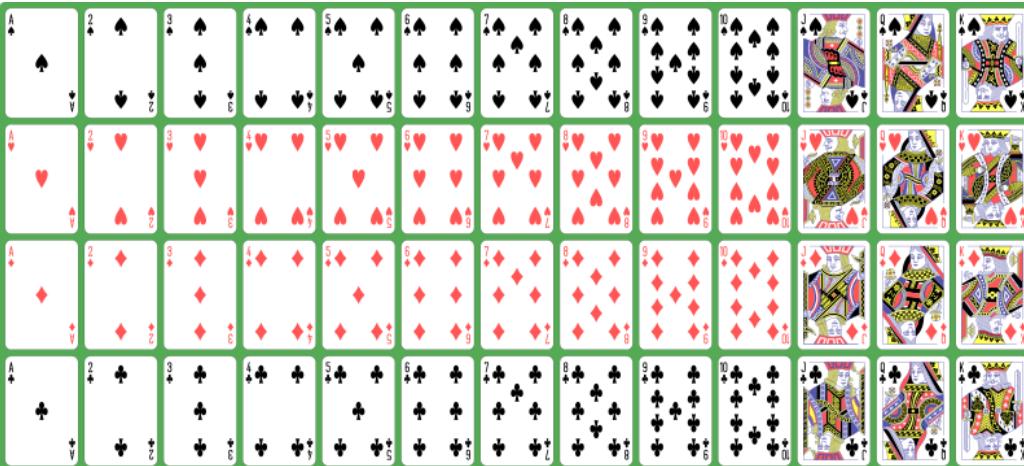
- Start with a standard deck of 52 cards:
  1. Sample a random card:  
(put it back and re-shuffle)  

  2. Sample a random card:  
(put it back and re-shuffle)  

  3. Sample a random card:  
(put it back and re-shuffle)  

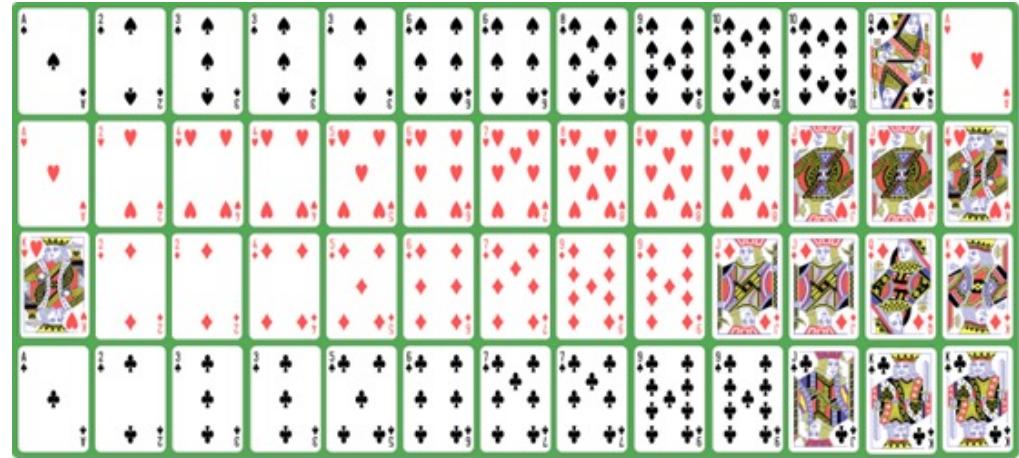
  - ...
  52. Sample a random card:  
(which may be a repeat)  

- Makes a new deck of the 52 samples:



# Bootstrap Sampling

- New 52-card deck is called a “bootstrap sample”:



- Some cards will be missing, and some cards will be duplicated.
  - So calculations on the bootstrap sample will give different results than original data.
- However, the bootstrap sample roughly maintains trends:
  - Roughly 25% of the cards will be diamonds.
  - Roughly 3/13 of the cards will be “face” cards.
  - There will be roughly four “10” cards.
- Common use: compute a statistic based on several bootstrap samples.
  - Gives you an idea of how the statistic varies as you vary the data.

# Random Forest Ingredient 1: Bootstrap

- **Bootstrap sample** of a list of ‘n’ examples:

- A new set of size ‘n’ chosen independently with replacement.

```
for i in 1:n  
    j = rand(1:n) # pick a random number from {1,2,...,n}  
    X_bootstrap[i,:] = X[j,:] # use the random sample
```

- Gives new dataset of ‘n’ examples, with some duplicated and some missing.
    - For large ‘n’, approximately 63% of original examples are included.

- **Bagging**: using bootstrap samples for ensemble learning.

- Generate several **bootstrap samples of the examples** ( $x_i, y_i$ ).
  - Fit a **classifier** to each bootstrap sample.
  - At test time, take **vote based on the predictions**.

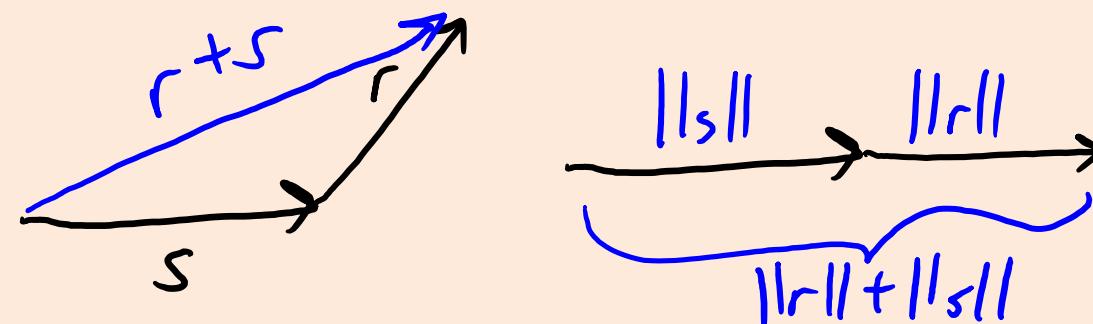
# Summary

- Encouraging invariance with **data augmentation**:
  - Add transformed data to be insensitive to the transformation.
- **Ensemble methods** take multiplier classifiers as inputs.
- **Voting ensemble method**:
  - Improves predictions of multiple classifiers if errors are independent.
- **Bagging**:
  - Ensemble method where we apply same classifier to “bootstrap samples”.
- Next time:
  - Unsupervised learning.

bonus!

# 3 Defining Properties of Norms

- A “norm” is any function satisfying the following 3 properties:
  1. Only ‘0’ has a ‘length’ of zero.
  2. Multiplying ‘r’ by constant ‘ $\alpha$ ’ multiplies length by  $|\alpha|$ 
    - “If  $r$  will twice as long if you multiply by 2”:  $\| \alpha r \| = |\alpha| \cdot \| r \|$ .
    - Implication is that norms cannot be negative.
  3. Length of ‘ $r+s$ ’ is not more than length of ‘ $r$ ’ plus length of ‘ $s$ ’:
    - “You can’t get there faster by a detour”.
    - “Triangle inequality”:  $\| r + s \| \leq \| r \| + \| s \|$ .



bonus!

# Squared/Euclidean-Norm Notation

We're using the following conventions:

The subscript after the norm is used to denote the p-norm, as in these examples:

$$\|x\|_2 = \sqrt{\sum_{j=1}^d w_j^2}.$$

$$\|x\|_1 = \sum_{j=1}^d |w_j|.$$

If the subscript is omitted, we mean the 2-norm:

$$\|x\| = \|x\|_2.$$

If we want to talk about the *squared* value of the norm we use a superscript of "2":

$$\|x\|_2^2 = \sum_{j=1}^d w_j^2.$$

$$\|x\|_1^2 = \left( \sum_{j=1}^d |w_j| \right)^2.$$

If we omit the subscript and have a superscript of "2", we're taking about the squared L2-norm:

$$\|x\|^2 = \sum_{j=1}^d w_j^2.$$

bonus!

# L<sub>p</sub>-norms

- The L<sub>1</sub>-, L<sub>2</sub>-, and L<sub>∞</sub>-norms are special cases of L<sub>p</sub>-norms:

$$\|x\|_p = (|x_1|^p + |x_2|^p + \cdots + |x_n|^p)^{1/p}$$

- This gives a norm for any (real-valued) p ≥ 1.
  - The L<sub>∞</sub>-norm is the limit as 'p' goes to ∞.
- For p < 1, not a norm because triangle inequality not satisfied.

# Why does Bootstrapping select approximately 63%?

bonus!

- Probability of an arbitrary  $x_i$  being selected in a bootstrap sample:

$p(\text{selected at least once in } n \text{ trials})$

$$= 1 - p(\text{not selected in any of } n \text{ trials})$$

$$= 1 - (p(\text{not selected in one trial}))^n$$

$$= 1 - (1 - 1/n)^n$$

$$\approx 1 - 1/e$$

$$\approx 0.63$$

(trials are independent)

(prob =  $\frac{n-1}{n}$  for choosing  
any of the  $n-1$  other samples)

( $(1 - 1/n)^n \rightarrow e^{-1}$  as  $n \rightarrow \infty$ )

bonus!

# Why Averaging Works

- Consider ‘k’ independent classifiers, whose errors have a variance of  $\sigma^2$ .
- If the errors are IID, the variance of the vote is  $\sigma^2/k$ .
  - So the more classifiers that vote, the more you decrease error variance.  
(And the more the training error approximates the test error.)
- Generalization to case where classifiers are not independent is:

$$c \sigma^2 + \frac{(1-c)}{k} \sigma^2$$

- Where ‘c’ is the correlation.
- So the less correlation you have the closer you get to independent case.
- Randomization in random forests decreases correlation between trees.
  - See also “[Sensitivity of Independence Assumptions](#)”.

# How these concepts often show up in practice

bonus!

- Here is a recent e-mail related to many ideas we've recently covered:
  - “However, the performance did not improve while the model goes deeper and with augmentation. The best result I got on validation set was 80% with LeNet-5 and NO augmentation (LeNet-5 with augmentation I got 79.15%), and later 16 and 50 layer structures both got 70%~75% accuracy.

In addition, there was a software that can use mathematical equations to extract numerical information for me, so I trained the same dataset with nearly 100 features on random forest with 500 trees. The accuracy was 90% on validation set.

I really don't understand that how could deep learning perform worse as the number of hidden layers increases, in addition to that I have changed from VGG to ResNet, which are theoretically trained differently. Moreover, why deep learning algorithm cannot surpass machine learning algorithm?”
- Above there is data augmentation, validation error, effect of the fundamental trade-off, the no free lunch theorem, and the effectiveness of random forests.