# Bayesian Classifiers / unsupervised learning

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Applied Machine Learning (EDAN96)
Lecture 13
2023-12-11
Elin A.Topp
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Material based on Lecture Slides on Probabilistic Representation and Bayesian Learning, EDAF70, Spring 2018,
Lecture 11, EDAN95 Fall 2018
Goodfellow et al, "Deep Learning", and Russel/Norvig, "AI - A Modern Approach"

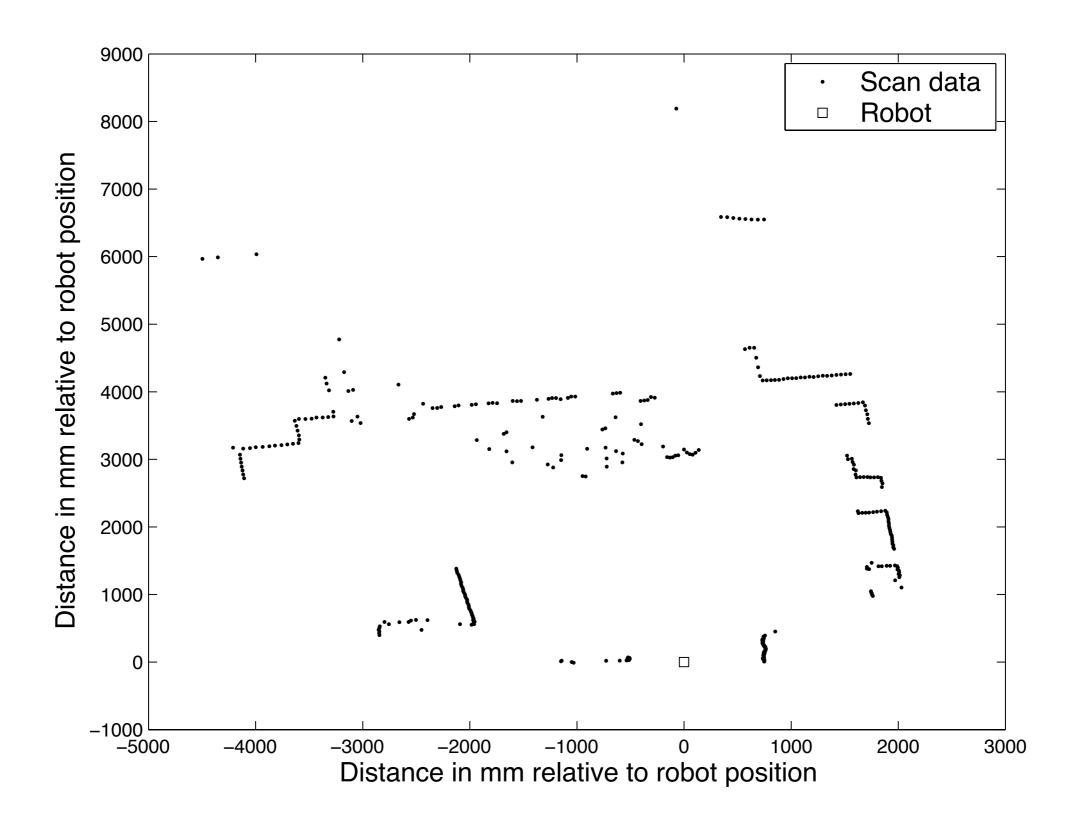
## Today's agenda

- Recap (if necessary) conditional / posterior probabilities, Bayes' rule, independence / conditional independence
- The Naive Bayesian Classifier and the Gaussian Naive Bayesian Classifier
- Learning a Bayesian Classifier
- The unsupervised version: EM (for GMMs or k-Means)

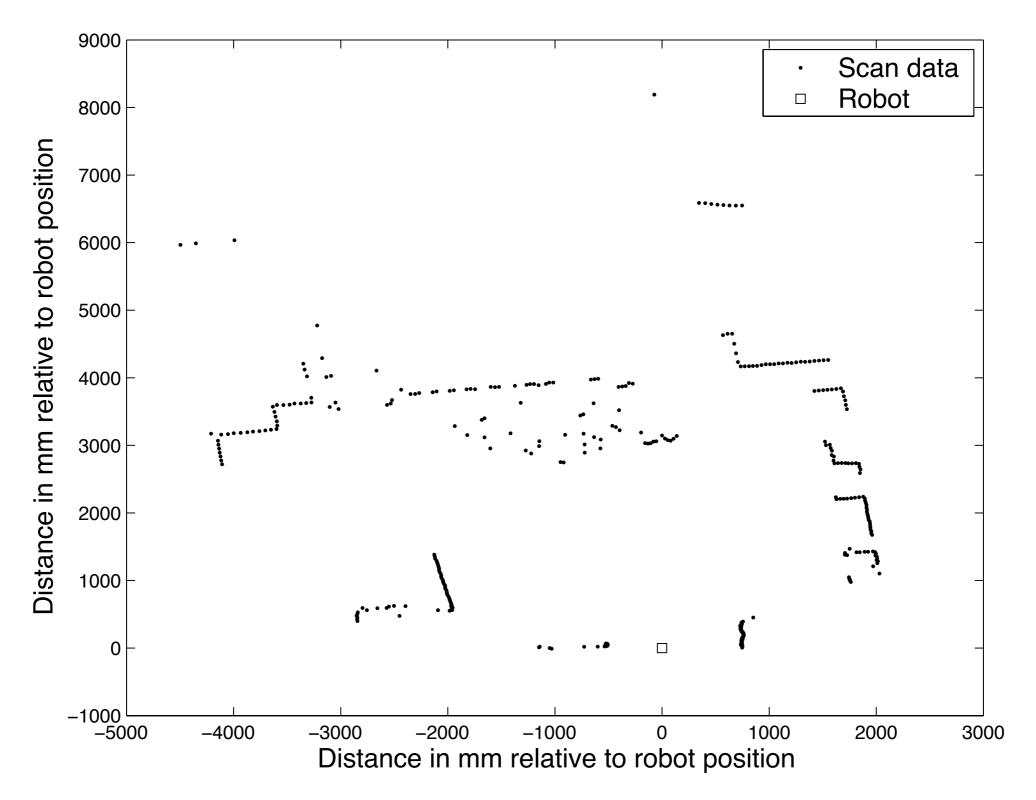
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#### A robot's view of the world...

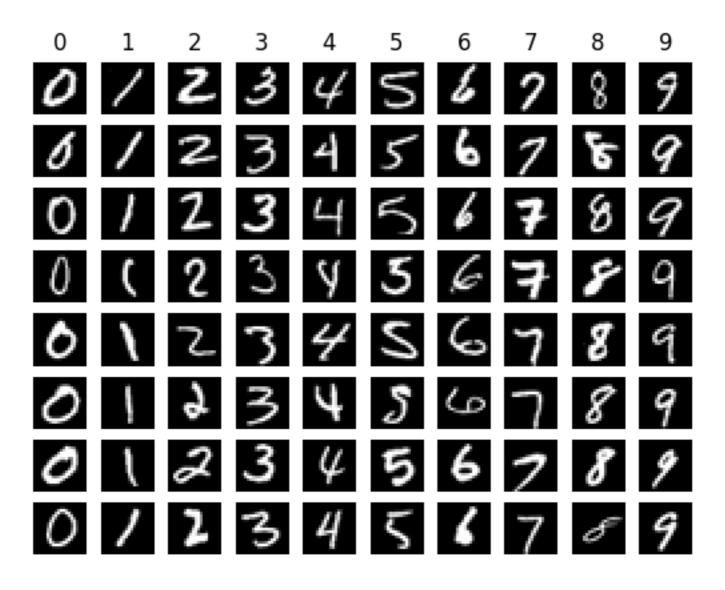


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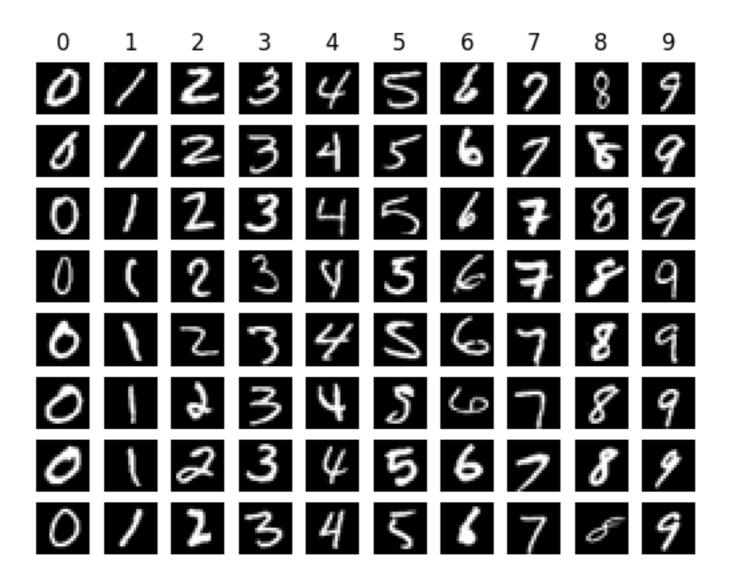


Which "leg-like" data point patterns were caused by a person's leg, which by furniture?

#### Or for MNIST-data:



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Which combinations of pixel values are most often seen for each of the numbers? Which number is it that best explains the pixel values of a specific sample?

We express propositions as random variables taking on certain values directly

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We look then for example at

$$p(X = x_i)$$
,  $i = 1,...$   $n$ , for all  $n$  values  $x_i$  of the Variable  $X$ 

E.g.:  $p(Class = c_1) = p(Class = c_2) = ... = p(Class = c_n) = 1/10$ 

with  $c_i$  = "the image is showing the number "i"

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For the distribution over the possible values of X we get then:

$$p(X) = \langle p(X = x_1), p(X = x_2), ..., p(X = x_n) \rangle$$

i.e., when joining distributions, we iterate over a subset of the values for X and Y in the computation:

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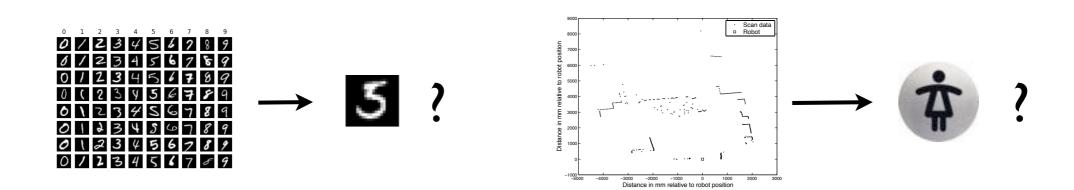
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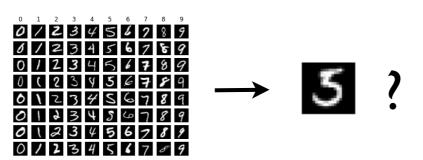
<sup>\*)</sup> There might be inconsistencies in the slides, anything that does not fit to this notation should be reported as faulty!

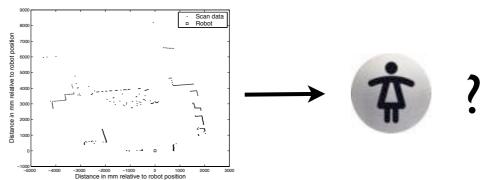
### Bayesian learning / classification

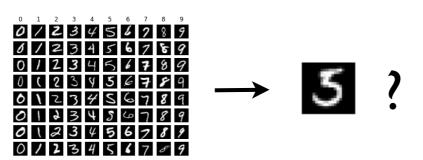
We want to classify / categorise / label new observations based on experience

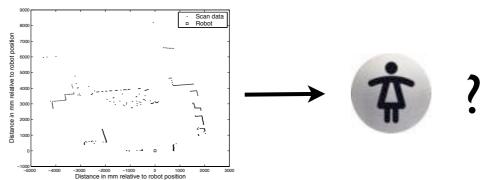
More general: We want to predict and explain based on (limited) experience, to find categories / labels for observations or even the model for "how things work" (transition models, sensor models) given a series of (explained) observations.

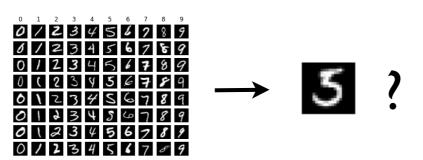


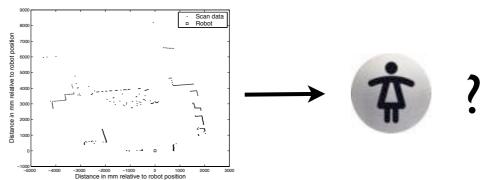


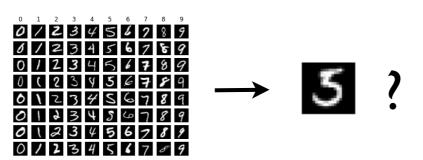


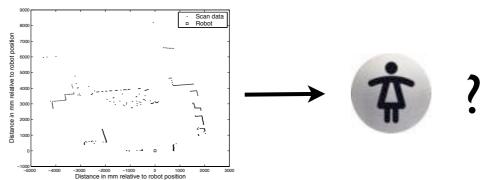


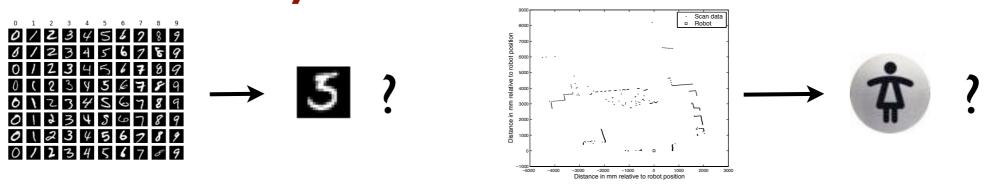




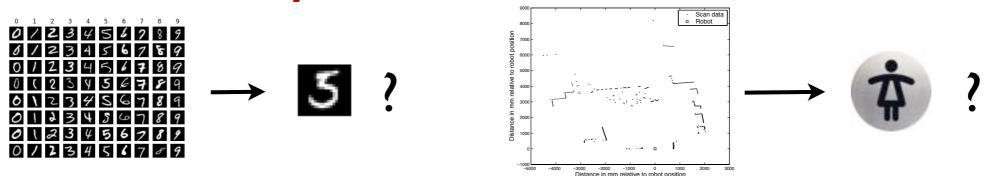








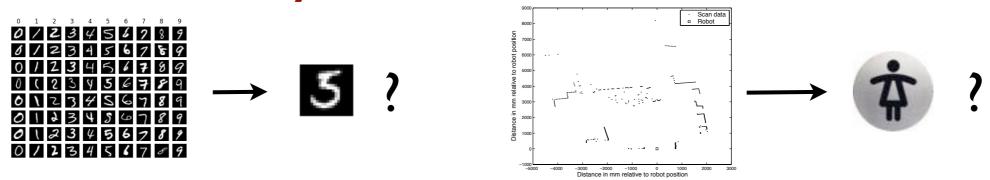
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Each image X has 64 pixels  $x_i$ , i = 0,...,63, that can take 17 values  $v_j = 0,...,16$ , these we can call features or attributes.



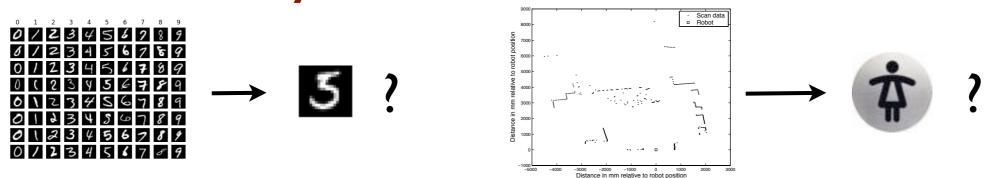
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or for the pattern problem:

Two classes of patterns, expressed through RV Person = {person,  $\neg person$ }, where each pattern has two features LegSize = {legSize,  $\neg legSize$ } and Curved = {curved,  $\neg curved$ }



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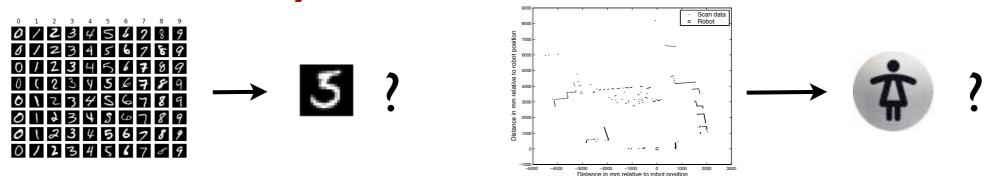
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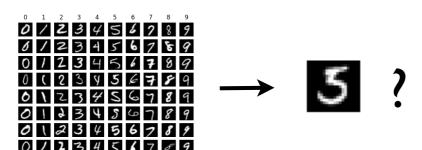
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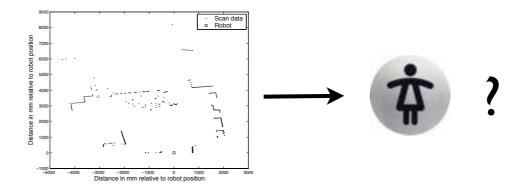
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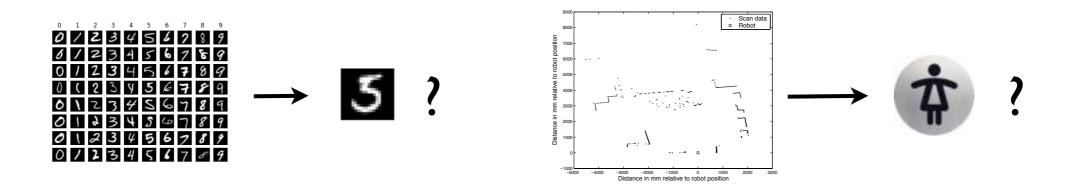
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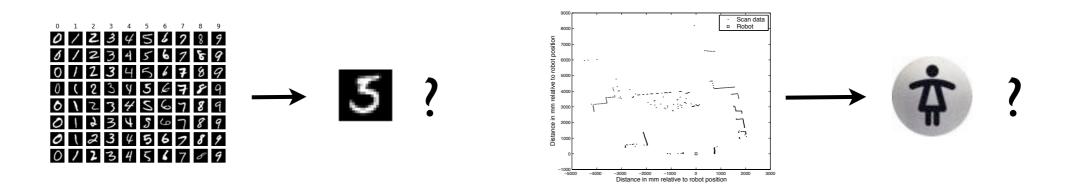
Now we make a new observation  $X_{new}$  of a combination of attributes (we see an image or pattern), and want to find the best hypothesis  $h^*$ , in this case directly corresponding to the class  $k^*$  or the value for *Person*.





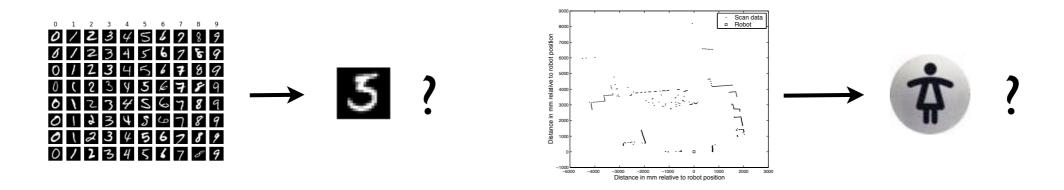


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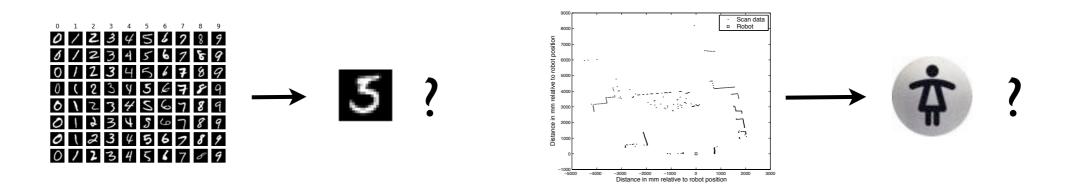
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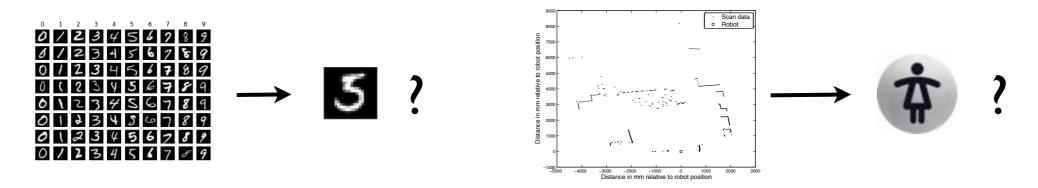
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What if one type of image is much more prevalent in the "world" (data set)? Would that be reflected?

# Maximum a Posteriori hypothesis



Use MAP, i.e., find the hypothesis / class that is best explained by the observation:

$$k^* = h_{MAP} = \underset{k}{\operatorname{argmax}} p(k | X_{new}) = \underset{k}{\operatorname{argmax}} \left[ p(k) p(X_{new} | k) \right]$$

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Can we do even better? This seems still bold, in particular if there is little data. What if our assumption of the underlying distribution is not correct?

Will get back ... first, there is another problem: how do we compute  $p(X_{new} | k)$ ?

Remember:  $X_{new}$  is a combination of attribute values  $x_i$ , i.e., we want

$$p(X_{new} | k) = p(x_0 = v_0, x_1 = v_1, \dots x_n = v_n | k)$$

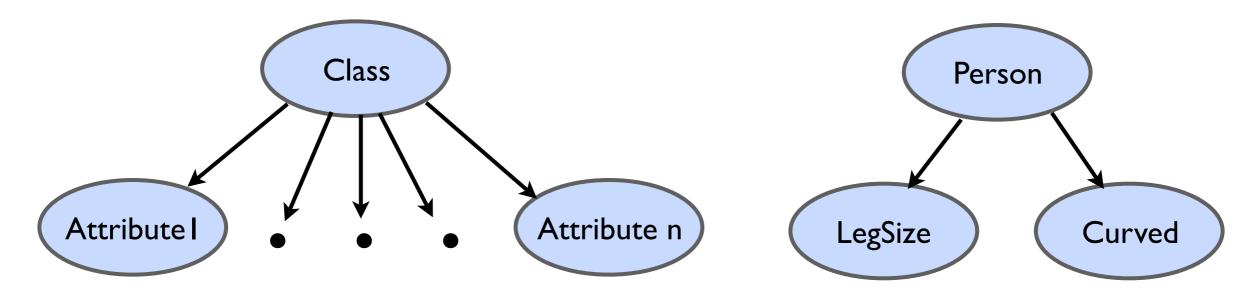
which becomes very complex with only few features, if we cannot assume independence.

## Conditional independence in the MAP-approach

```
p(Class | Attribute I = aI \land Attribute 2 = a2)
= \alpha p(aI \land a2 | Class) p(Class)
= \alpha p(aI | Class)p(a2 | Class) p(Class)
```

This gives a naive Bayes model (think Class = Cause, Attribute = Effect):

$$p(Class, Attribute_{1,...,} Attribute_{n}) = p(Class) \prod_{i} p(Attribute_{i} | Class)$$



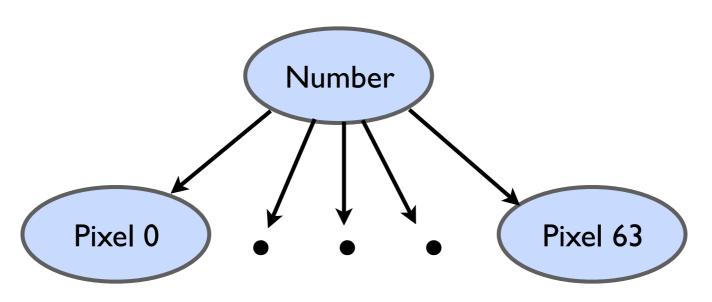
The total number of parameters is linear in n, and

$$p(Class = c^*) = \underset{c}{\operatorname{argmax}}[p(Class = c, Attribute_0, \dots, Attribute_{n-1})]$$

$$= \underset{c}{\operatorname{argmax}}[p(Class = c) \prod_{i} p(Attribute_i = av_j | Class = c)]$$

where  $av_i$  is the value that is observed for  $Attribute_i$ 

## A (super naïve) NBC for the digits data



$$CPT_{0jk} = p(Pixel_0 = v_j | Number = k)$$

$$CPT_{63jk} = p(Pixel_{63} = v_j | Number = k)$$

For a given (unknown) image  $X_{new}$  with  $pixel_{new_i}$  representing the value of  $Pixel_i$  in  $X_{new}$  we want to classify we get:

$$Number(X_{new}) = \arg\max_{k} [p(Number = k, pixel_{new_0}, \dots, pixel_{new_{n-1}})]$$

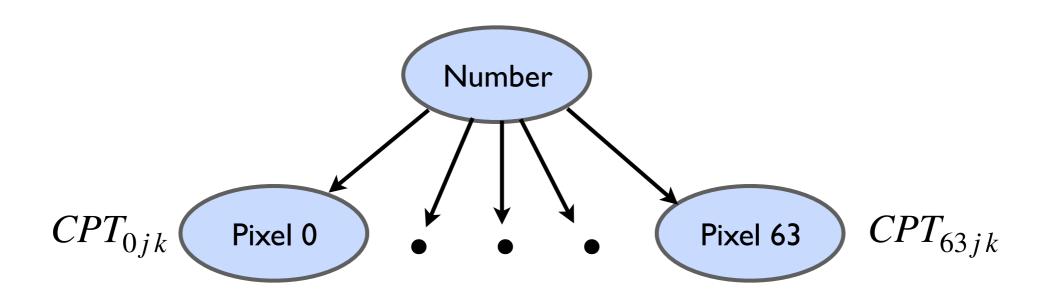
$$= \arg\max_{k} [p(Number = k) \prod_{i} p(x_{new_i} | Number = k)]$$

note: all possible attribute values are here the same for each attribute, hence  $v_i$ , not  $av_i$ 

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# Learning = "assigning" (?)



Simply set 
$$CPT_{ijk} = p(pixel_i = v_j | Number = k) = \frac{|X_{iv_jk}|}{|X_k|}$$

with

 $X_{iv_ik} = \{examples \ X \ belonging \ to \ class/number \ k, \ where \ pixel \ i \ in \ X \ has \ value \ v_j\}$ 

# What if...?

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Then 
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This is "super naïve", but why is it actually bad?

Instead of 
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 use  $CPT_{ijk} = \frac{|X_{iv_jk}| + mp}{|X_k| + m}$ 

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Technically, the m-estimate in itself can be learned from / adapted to the data in the process.

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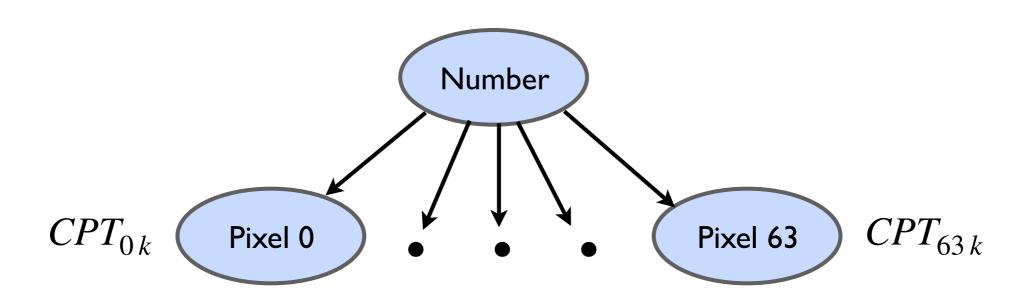
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- "Blur" the probabilities by using a suitable distribution (often, not always, a Gaussian Normal Distribution can help)

#### Gaussian Mixture Model

- Assume that the *n* attributes in the example set form the axes of an *n*-dimensional feature space, i.e., each example is a "point" in that space.
- The examples belonging to a class will then somehow "gather" around some centre "point"
- The degree of "belonging" can be expressed as a continuous PDF very often a Gaussian Normal distribution is suitable, which gives then a Gaussian Mixture Model (the multidimensional bell "curves" will most likely overlap, hence, there is a mixture of several distributions that explain a given data point the sample to be classified).
- see Goodfellow (3) / Murphy (2) for other "standard" distributions

## Gaussian Naive Bayesian Classifier



$$CPT_{ik} = (\mu_{ik}, \sigma_{ik}), \quad p(pixel_i = x | Number = k) = \frac{1}{\sqrt{2\pi\sigma_{ik}^2}} e^{-\frac{1}{2\sigma_{ik}^2}(x - \mu_{ik})^2}$$

with  $\mu_{ik} = mean(x_i)$  and  $\sigma_{ik}^2 = var(x_i)$   $\forall X \in \{examples \ where \ Number = k\}$  and  $x_i$  the value of  $Pixel \ i$  in a given image X

Classification is then handled for unseen sample  $X_{new}$ :

$$Number(X_{new}) = \underset{k}{\operatorname{argmax}} [p(Number = k, x_{new_0}, \dots, x_{new_{n-1}})]$$

$$= \underset{k}{\operatorname{argmax}} [p(Number = k) \prod_{i} p(x_{new_i}) | Number = k)]$$

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# EM-algorithm for GMMs

This is the EM-algorithm as given by Murphy, "Machine Learning - A Probabilistic Approach", p 353. assume data set X with examples  $\overrightarrow{x_i}$ , i=1,...,N and K classes you want to cluster X into.

#### EM-for-GMM( X, K)

- I. Initialize  $\theta_k^0 = (\pi_k^0, \overrightarrow{\mu_k}^0, Cov_k^0)$ , where
  - $\pi_k$  is the class prior for class k (e.g., assume uniform distribution here initially)

 $\overrightarrow{\mu_k}$  are the means for the attribute values per attribute a in class k (initially a random subset of X)

 $Cov_k$  is the covariance for the attribute values in class k (can be simplified to variance  $\sigma_{ak}^2$  for each attribute a if a G-NBC is assumed as the model)

2. Iterate over E and M steps as follows:

E-step:

compute 
$$r_{ik}^t = \frac{\pi_k^{t-1} p(\vec{x}_i \mid \theta_k^{t-1})}{\sum_{k'} \pi_{k'} p(\vec{x}_i \mid \theta_{k'}^{t-1})}$$
 where  $p(\vec{x}_i \mid \theta_k^{t-1}) = \prod_a \frac{1}{\sqrt{2\pi\sigma_{ak}^2}} e^{-\frac{1}{2\sigma_{ak}^2}(x - \mu_{ak}^{t-1})^2}$ , assuming that

the covariance can be substituted with  $\sigma_{ak}^2$  for attribute a and class k. Index i runs over the samples, index a runs over the attributes, and k over clusters ("classes")

M-step:

compute 
$$r_k^t = \sum_i r_{ik}^t$$
 and  $\pi_k^t = \frac{r_k^t}{N}$ , then update the means and variances:

$$\overrightarrow{\mu_k}^t = \frac{\sum_i r_{ik}^t \overrightarrow{x_i}}{r_k^t} \quad and \quad Cov_k^t = \frac{\sum_i r_{ik}^t \overrightarrow{x_i} \overrightarrow{x_i}^T}{r_k^t} - \overrightarrow{\mu_k^t} \overrightarrow{\mu_k^t}^T \text{ (from which the new } \sigma_{ak}^2 \text{ can be extracted)}$$

3. Stop, when the  $\overrightarrow{\mu_k}$  and  $Cov_k$  are not changing significantly anymore.

#### EM-algorithm for k-Means

This is the EM-algorithm as given by Murphy, "Machine Learning - A Probabilistic Approach", p 356. assume data set  $\mathbf{X}$  with examples  $\overrightarrow{x_i}$ , i=1,...,N and K classes you want to cluster  $\mathbf{X}$  into.

k-Means(X, K)

- I. Initialize  $\overrightarrow{\mu_k}^0$ , assume fixed class priors  $\pi_k$
- 2. Iterate over E and M steps as follows:

E-step:

Assign each data point to its closest cluster centre:  $z_i = \underset{k}{\operatorname{argmin}} \|\vec{x}_i - \overrightarrow{\mu}_k\|_2^2 = L_2(\vec{x}_i - \overrightarrow{\mu}_k)^2$ 

M-step:

Update each cluster centre by computing the means of all points assigned to it:

$$\overrightarrow{\mu}_k = \frac{1}{N_k} \sum_{i: z_i = k} \overrightarrow{x}_i$$

Until converged

## Today's summary

- (Refreshed memory on conditional probabilities, Bayes' rule, independence, conditional independence)
- Introduced Naive Bayesian Classifiers
- Introduced GNBCs (very briefly)
- Rushed through the EM-algorithm...

- Reading:
  - Goodfellow, ch 3, Murphy, ch 2
  - Lecture slides lecture 11, 2018
  - Mitchell, chapter 6