

AUTOMATIC IMAGE SEGMENTATION BY HISTOGRAM ANALYSIS

RECONNAISSANCE DES FORMES

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CHAPTER 1

Introduction

1.1 CONTEXT

This report is the result of a university assignment. It aims to prove that the student understood the motivation, goals and means of studying pattern recognition. Therefore, this is a way of summarizing a series of observations and experiments done on images containing shapes.

1.2 MOTIVATION

L'objectif du TP est de segmenter une image en niveaux de gris par le seuillage automatique des niveaux de gris par des macros R que vous allez concevoir. In order to be able to work with shapes, we must first acquire them. Most of the time, they are not given to us in their simplest form, but found in images with all kinds of backgrounds. They must be extracted as precisely as possible, making sure we differentiate them from the background as accurately as we can. Where there's a need for great precision and accuracy, there's also a need for research, so in this report we'll be looking over some methods we can use to identify objects (shapes).

1.3 GOALS

Having multiple grayscale images, we want to be able to do a *binary segmentation* on them in order to differentiate between the objects and the image's background. Given the fact that we only have gray values at our disposal, we must look into ways of using them in order to state that a certain pixel belongs to either an object or to the background.

CHAPTER 2

BINARISATION USING BAYES' THEOREM

2.1 LOADING DATA

The image we'll try to apply binarisation to is the following:



FIGURE 2.1 Image to experiment on

Which is composed of the following 2 images:

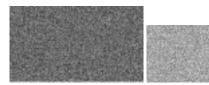


FIGURE 2.2
Separate images

The images will be be loaded as gray images:

```
# Chargement d'une image en niveaux de gris
rdfReadGreyImage <- function (nom) {
    image <- readImage (nom)
    if (length (dim (image)) == 2) {
        image
    } else {
        channel (image, 'red')
    }
}</pre>
```

LISTING 2.1 Loading data

2.2 FIXED THRESHOLD

Taking a look at the histogram, we can have a "manual" attempt at binarising the image: using a *fixed threshold*. We simply choose a value for the threshold and we assign, for each pixel, one of the predicted classes as it follows:

$$\hat{w_1} = \{ P \in I | I(P) < \hat{X} \}$$

$$\hat{w_2} = \{ P \in I | I(P) \ge \hat{X} \}$$

where \hat{X} is our threshold.

Histogram of as.vector(image)

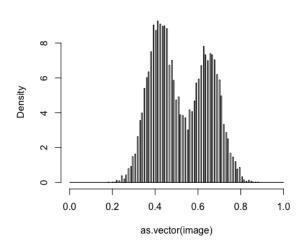


FIGURE 2.3
Histogram of grey values for

```
# for each threshold, calculate the binary image
# image has values between [0, 1] so our threshold has to be in the same range
# for each pixel, the expression (pixel_value - threshold) >= 0 will assign it
to one of two classes: w1 or w2

binaire50 <- (image - 0.5) >= 0

display (binaire50, "image binaire 0.35", method="raster", all=TRUE)
binaire55 <- (image - 0.55) >= 0

display (binaire55, "image binaire 0.35", method="raster", all=TRUE)
binaire60 <- (image - 0.6) >= 0

display (binaire60, "image binaire 0.35", method="raster", all=TRUE)
```

LISTING 2.2
Using fixed threshold

Let's take a look at the results, using threshold values of 0.5, 0.55 and 0.6.



FIGURE 2.4
Fixed threshold results

It is not hard to see that there is a lot of noise in our predictions and they are far from perfect. The threshold value of 0.6 might seem attractive, but it only manages to classify well most of the pixels from the image on the top. Further more, it heavily missclassifies the pixels from the image on the bottom.

2.3 A PRIORI CLASS PROBABILITIES

Our image 2.1 is formed of 2 separate images 2.2. We can try to calculate the following probabilities:

```
P(w_1), the probability of a pixel to be in w_1 class P(w_2), the probability of a pixel to be in w_1 class
```

Knowing that the dimension of the merged image is 100x100 and the 2 images are 100x43, 100x57, the probabilities above are easy to calculate.

```
calculateClassAPrioriProbabilities <- function() {
combined <- rdfReadGreyImage ("2classes_100_100_8bits_2016.png")
above <- rdfReadGreyImage ("2classes_100_100_8bits_omega1_2016.png")
below <- rdfReadGreyImage ("2classes_100_100_8bits_omega2_2016.png")

# pw1 = (number_of_pixels_from_the_first_image) / (number_of_total_pixels)
# pw1 = (number_of_pixels_from_the_second_image) / (number_of_total_pixels)
noTotalPixels = dim(combined)[1] * dim(combined)[2]
pw1 = (dim(above)[1] * dim(above)[2]) / noTotalPixels
pw2 = (dim(below)[1] * dim(below)[2]) / noTotalPixels
c(pw1, pw2)</pre>
```

LISTING 2.3
A priori class probabilities

In this case, the probabilities are $P(w_1) = 0.57$ and $P(w_2) = 0.43$.

2.4 CONDITIONAL PROBABILITIES

Having $P(w_1)$ and $P(w_2)$, we can take a step further and calculate the following conditional probabilities:

 $P(X|w_1)$, the probability of a pixel having the gray value of X if it's in the w_1 class $P(X|w_2)$, the probability of a pixel having the gray value of X if it's in the w_2 class

These are, according to Wikipedia 2020, equal to:

$$P(X|w) = \frac{P(w|X) * P(X)}{P(w)}$$

We already have P(w) and P(X) is simply the number of pixels with the gray value of X divided by the total number of pixels. P(w|X) is the number of pixels with the value X in the image corresponding to X divided by the total number of pixels with a value of X.

```
calculateConditionalProbability <- function (X) {</pre>
           probs <- calculateClassAPrioriProbabilities()</pre>
           nom <- "2classes_100_100_8bits_2016.png"</pre>
3
           image <- rdfReadGreyImage (nom)</pre>
           nbins <- 256
           h <- hist (as.vector (image), freq=FALSE, breaks = seq (0, 1, 1 / nbins))
           \# we can notice that h$counts[X + 1] gives us the number of pixels that have
       the gray value of X
           # the "+ 1" above is because in R arrays start from 1
           \# so, P(X \mid I) = the probability of a pixel from the merged image to have
      the gray value of X
          px <- h$counts[X + 1] / sum(h$counts)</pre>
10
           \# P(X \mid w) = the probability of a pixel from the merged image to have the
      gray value of X
           # IF that pixel is from the w class
12
           # Using bayes theorem, we get:
13
           \# P(X|W) = (P(W|X) * P(X)) / P(W)
14
           \# and P(w \mid X) = how many pixels of value X are in the image corresponding
15
      to w / the total number of pixels with a value of X
16
           above <- rdfReadGreyImage ("2classes_100_100_8bits_omega1_2016.png")</pre>
17
          below <- rdfReadGreyImage ("2classes_100_100_8bits_omega2_2016.png")
18
          h1 <- hist (as.vector (above), freq=FALSE, breaks = seq (0, 1, 1 / nbins))
19
20
          h2 <- hist (as.vector (below), freq=FALSE, breaks = seq (0, 1, 1 / nbins))
          p_w1_{if_x} \leftarrow h1$counts[X + 1] / h$counts[X + 1]
23
          p_x_and_w1 \leftarrow p_w1_if_x * px
          p_x_{if_w1} \leftarrow p_x_{and_w1} / probs[1]
24
          p_w2_{if_x} \leftarrow h2$counts[X + 1] / h$counts[X + 1]
25
          p_x_and_w2 \leftarrow p_w2_if_x * px
          p_x_{if}w2 \leftarrow p_x_{and}w2 / probs[2]
27
          c(px, p_x_if_w1, p_x_if_w2)
28
```

LISTING 2.4
Calculating conditional probabilities

Running the above function for X = 141, we get the following probabilities:

$$P(X|I) = 0.011800000$$

 $P(X|w_1) = 0.008947368$
 $P(X|w_2) = 0.015581395$

where P(X|I) refers to the probability of a pixel to have the gray value of X in the whole (merged) image. Of course, $P(X|w_1) * P(w_1) + P(X|w_2) * P(w_2) = P(X|I)$.

2.5 AUTOMATIC THRESHOLDING USING BAYES

Having a threshold \hat{X} , we can define an assignment error as being the following:

$$P(\epsilon|\hat{X}) = \underset{x}{\operatorname{argmin}} \left(\sum_{X \in \hat{w}_2} P(X|w_1) * P(w_1) + \sum_{X \in \hat{w}_1} P(X|w_2) * P(w_2) \right)$$

Our goal is to find the \hat{X} that minimizes $P(\epsilon|\hat{X})$. For this, we can try each value for \hat{X} and see which is the best.

CHAPTER 3

CONCLUSION

Using only the gray values or the texture values as attributes did not result in a great segmentation. As we've seen in the experiments, the error rate, when using each attribute individually, can be pretty high in certain cases, even though for some textures the methods are really precise.

However, where one of them did bad, the other did well. By combining the two, we have a more precise way of segmenting the image. In conclusion, we may state that the combination of the two attributes is a worthy contender for binary segmentation.

BIBLIOGRAPHY

Wikipedia (2020). *Bayes' Theorem*. Wikimedia Foundation. URL: https://en.wikipedia.org/wiki/Bayes%27_theorem (visited on 27th Feb. 2020).