Project 4_G16

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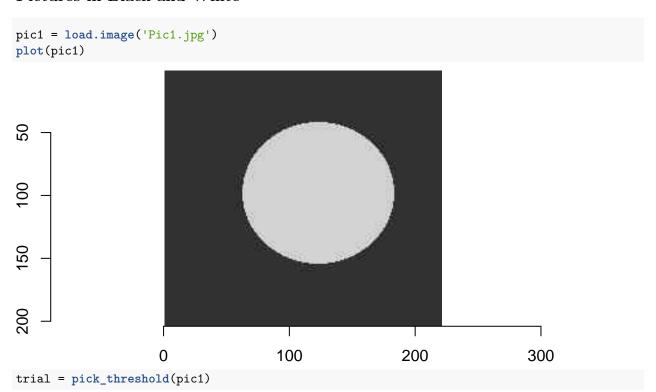
Part 1

```
pick_threshold = function(picture){
  histogram = hist(picture)
  freq = histogram$counts

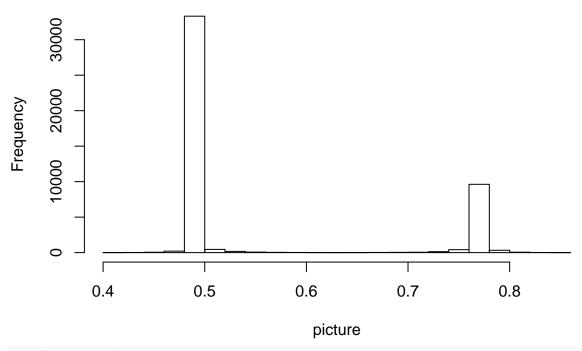
derivatives = diff(sign(diff(freq)))==2
  midpoint = which(derivatives)+1

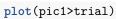
thresh = histogram$breaks[midpoint]
  return(mean(thresh))
}
```

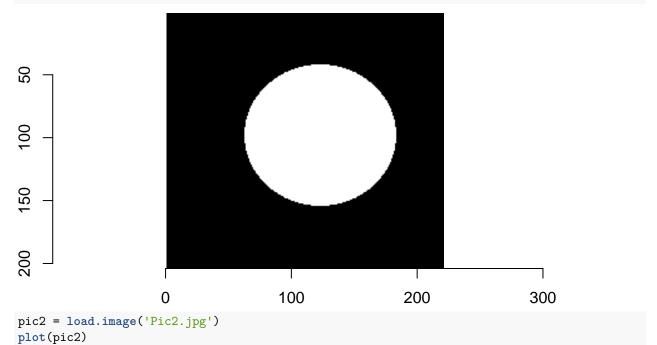
Pictures in Black-and-White

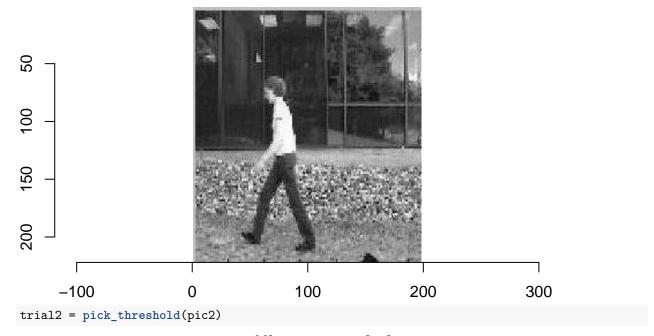


Histogram of picture

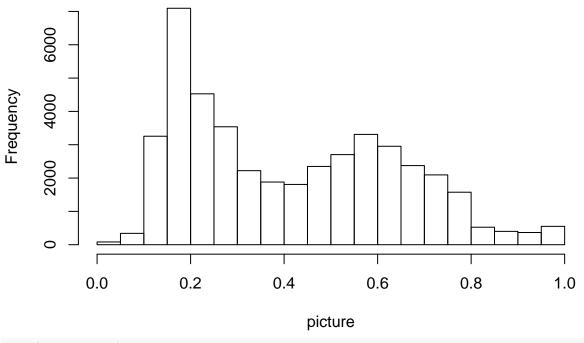


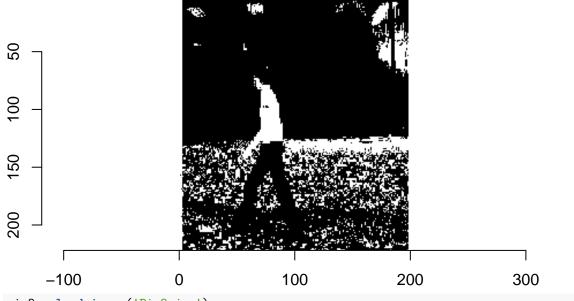




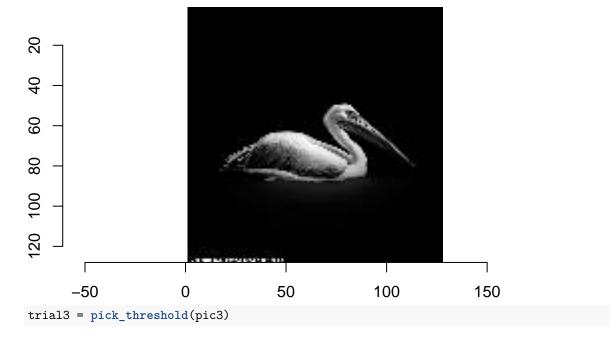


Histogram of picture

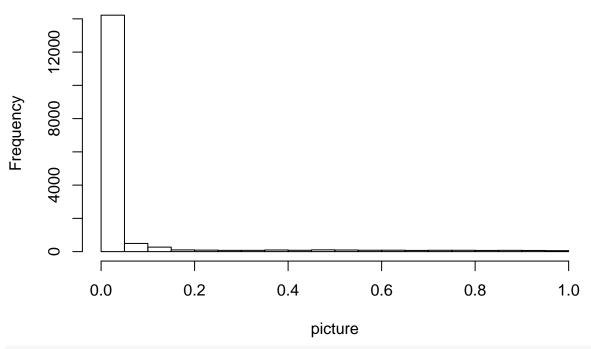


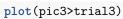


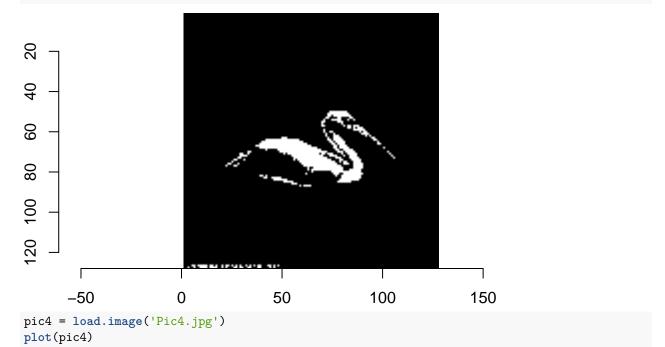
pic3 = load.image('Pic3.jpg')
plot(pic3)

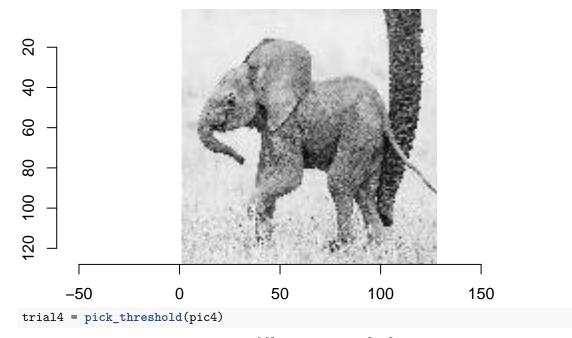


Histogram of picture

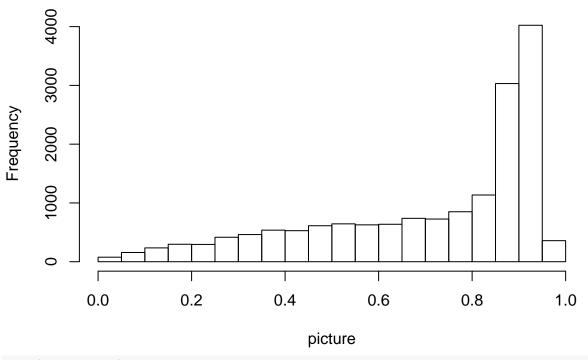




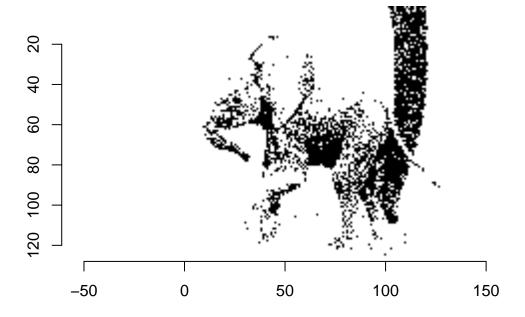




Histogram of picture



plot(pic4>trial4)



Part 2

```
image_segmentation = function(picture){
 pic = resize(picture, 128, 128)
 xrange = c(10, 120)
  yrange = c(40, 128)
  tempforeground = imsub(pic,x %inr% xrange, y %inr% yrange)
  tempbackground = imsub(pic, !(x %inr% xrange), !(y %inr% yrange))
  #compute the mean for each subimage
  p_bar_f = mean(c(tempforeground))
 p_bar_b = mean(c(tempbackground))
  #make the a and b lists
 picture_matrix = matrix(pic, 128,128)
 a_{vector} = c()
  b_vector = c()
 for (p_i in c(pic)){
   a_i = -\log(abs(p_i-p_bar_f)/(abs(p_i-p_bar_f)+abs(p_i-p_bar_b)))
   a_i = min(a_i, 1000000)
   b_i = -\log(abs(p_i-p_bar_b)/(abs(p_i-p_bar_f)+abs(p_i-p_bar_b)))
   b_i = min(b_i, 1000000)
   a_vector= c(a_vector,a_i)
   b_vector= c(b_vector,b_i)
```

```
# get the column pairs
x = c(1:(dim(picture_matrix)[1]*dim(picture_matrix)[2]))
# adjacency matrix
adjacent_matrix = matrix(x, dim(picture_matrix)[1], dim(picture_matrix)[2], byrow=TRUE)
f1=head(adjacent matrix, -1)
t1 = tail(adjacent_matrix, 127)
# transposed adjacency matrix
transpose_adjacent = t(adjacent_matrix)
f2 = head(transpose_adjacent, -1)
t2 = tail(transpose_adjacent, 127)
# top portion of dataframe of to and from
to = c(c(t2), c(t1))
from = c(c(f2), c(f1))
top_df = data.frame(from, to)
# Initialize before loop
pic_list = c(picture_matrix)
c_{vec} = c()
K = .01
sigma = 1
# Find capacities
for (i in 1:nrow(top_df)){
  row = top_df[i,]
  p_i = pic_list[row$from]
  p_j = pic_list[row$to]
  numer = -((p_i-p_j)^2)
  denom = (sigma^2)
  cij = K*exp(numer/denom)
  c_{vec} = c(c_{vec}, cij)
}
# add capacity to top portion of dataframe
top_df$capacity = c_vec
\# Create and stack bottom dataframe
bottom_df = data.frame(top_df$to,top_df$from,top_df$capacity)
names(bottom_df) = c("from", "to", "capacity")
# Initialize
lower_df = rbind(top_df,bottom_df)
```

```
# add source to nodes and nodes to sink
nodes=seq(1,128^2)
s=rep(1,128^2)
t=rep(128<sup>2</sup>,128<sup>2</sup>)
nodes_from=c(s,nodes)
nodes_to=c(nodes,t)
weights=c(a_vector,b_vector)
upper_df=data.frame(nodes_from,nodes_to,weights)
names(upper_df) <- c("from", "to","capacity")</pre>
final_df<-rbind(upper_df, lower_df)</pre>
graph=graph_from_data_frame(final_df, directed = TRUE, vertices = NULL)
maxflow=max_flow(graph, 1, 128^2)
new_img = rep(0,128^2)
new_img[maxflow$partition1]=0
new_img[maxflow$partition2]=1
new_img = matrix(new_img,128,128)
new_img = array(new_img, c(128,128,1,1))
new_img = as.cimg(new_img)
color_scale <- scales::gradient_n_pal(c("red","blue"),c(1,0))</pre>
par(mfrow=c(1,2))
plot(pic,main = 'Input')
plot(new_img,main='Segmented', colourscale=color_scale,rescale=FALSE)
```

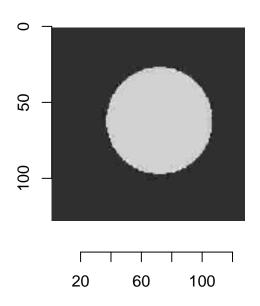
Image Segmentation

Image 1

```
image_segmentation(pic1)
```

Input

Segmented



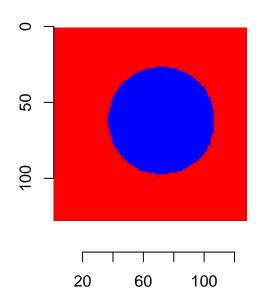
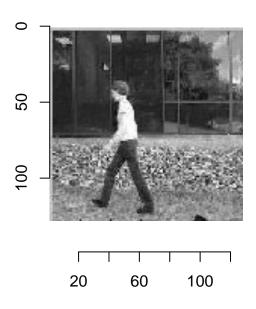


Image 2

image_segmentation(pic2)

Input

Segmented



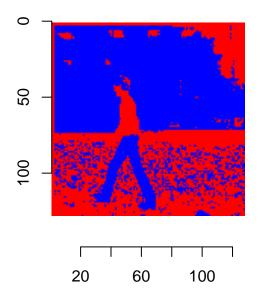
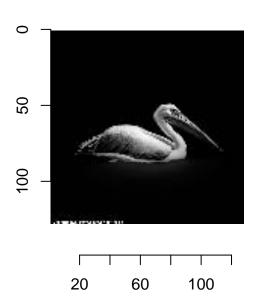


Image 3

image_segmentation(pic3)



Segmented



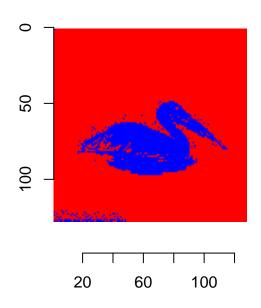


Image 4

image_segmentation(pic4)

Input

Segmented

