Henri, Jennifer, & Josh – Group A

RUDIMENTARY FACIAL CLASSIFICATION WITH PRINCIPAL COMPONENT ANALYSIS AND SUPPORT VECTOR MACHINES



Outline

- Project goals
- 2. Data set
- 3. Principal Component Analysis and Support Vector Machines
- 4. Our method
- 5. Results of PCA, SVM, and classification
- 6. Areas of improvement
- 7. Conclusion

Project goals

Use principal component analysis and support vector machines to quickly and accurately classify a face



Result:
Laura Bush

(Not what we want)





- We used the dataset Labeled Faces in the Wild from University of Massachusetts^[2]
- 2. Total of 13293 images and 5752 people in the database.
- 3. We took 60 pictures from every person that had at least 60 pictures in the database (8 people)
- 4. Pictures needed to be in .jpg format and size 250x250
- 5. We deal with the grayscale and cropping of the pictures

Data set people



Ariel Sharon



Colin Powell



Donald Rumsfeld



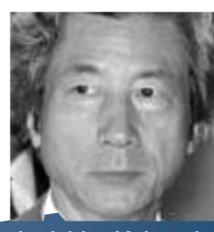
George W Bush



Gerhard Schroeder



Hugo Chavez



Junichiro Koizumi



Tony Blair

What is Principal Component Analysis (PCA)?

- PCA is a technique used for dimension reduction with minimal information loss^[1]
- Two of the biggest uses of PCA are clustering and dimension reduction
- In this project we use PCA to reduce the dimensions of the dataset to improve accuracy in facial classification

- We're given 480 250x250 images (crop them to 125x125)
- We convert each image into a 125² dimensional vector

$$x_1, x_2, \dots, x_{480} \in \mathbb{R}^{125^2}$$

Then we take the mean vector of all these images

$$\vec{\mu} = \frac{1}{480} \sum_{i=1}^{480} \vec{x}_i$$

Define the variance of any vector \vec{u} as following

$$var(\vec{u}) = \frac{1}{480} \sum_{i=1}^{480} |\vec{u}^*(\vec{x}_i - \vec{\mu})|^2$$

• We want to find the vector \vec{u} that maximizes this variance to find the direction that captures the most variation in the dataset

Notice that the expression can be written

$$\operatorname{var}(\vec{u}) = \vec{u}^* (YY^*) \vec{u} \text{ and } Y = \frac{1}{\sqrt{480}} [\vec{x}_1 - \vec{\mu}| \cdots | \vec{x}_{480} - \vec{\mu}]$$

$$C = \frac{1}{480} \sum_{i=1}^{480} (\vec{x}_i - \vec{\mu}) (\vec{x}_i - \vec{\mu})^*$$

• Where $C = YY^*$ is the sample covariance matrix and

$$var(\vec{u}) = \vec{u}^* C \vec{u}$$

 When we are trying to maximize the variance, this is the same as trying to maximize the previous expression

$$\max_{\|\vec{u}\|=1} \vec{u}^* C \vec{u} = \lambda_{\max}(C) = \sigma_1(Y^*)^2$$

• And the maximizing \vec{u} is the top left singular vector of Y, or the first eigenvector of C

- Define the principal components of a data set X to be the eigenvectors of the sample covariance matrix
- Then we have the following eigendecomposition with the eigenvalues in decreasing order

$$C = U\Lambda U^*$$

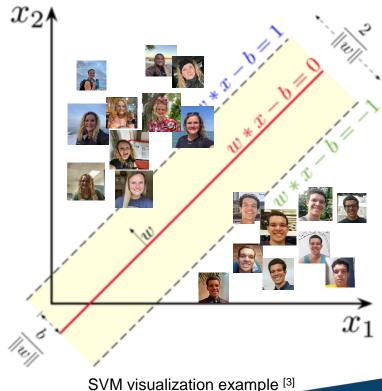
And the columns of *U* are the principal components

- In most datasets, only a few eigenvalues of C are large, so we can write each data point as a linear combination of only a few principal components with very little information lost
- This allows us to reduce the dimensions of the data significantly and avoid problems like collinearity when training our model

- First r principal components: multiply dataset by the 1st r columns of the matrix U to get the data in PCA form (reduced dimensions)
- Then use that reduced dimension data to train our SVM model
- Alternatively, you could use the function prcomp() in R and it does all that work for you

What is Support Vector Machines (SVM)?

- SVM is a supervised machine learning algorithm used for classification
- We use SVM to classify faces given images that have been transformed into lower dimensions with PCA
- For the math behind SVM, view Topic 10



Filter Data

```
filter_data <- function(path, number_pics, new_path=NULL, test=TRUE)
 if(test)
 all_paths<-list.dirs(path = path, full.names = TRUE, recursive = FALSE)
 # Create empty list, containing number of pics inside that path
 sizes<-c()
 # Loop through all the paths
 for (i in 1:length(all_paths)) {
   sizes<-c(sizes,length(list.files(all_paths[i])))</pre>
 list new paths <- c()
 for (i in 1:length(all_paths)){
   if(sizes[i]>=number_pics)
     list_new_paths<-c(list_new_paths, all_paths[i])
 # Create the new folders in the new location
 dir.create(new_path)
 file.copy(from=list_new_paths, to=new_path,
          overwrite = TRUE, recursive = TRUE,
          copy.mode = TRUE)
 return(list_new_paths)
```

Create Matrix of Images

```
create_matrix <- function(folders,nphotos=60,n=100)</pre>
 X <- c()
   for (i in 1:length(folders))
      setwd(paste0(reduced dir, "/",folders[i]))
      photos <- dir(path = paste0(reduced_dir, "/",folders[i]),</pre>
                    pattern = NULL, all.files = FALSE,
                    full.names = FALSE, recursive = FALSE,
                    ignore.case = FALSE, include.dirs = FALSE, no.. = FALSE)
      for (j in 1:nphotos)
        pic <= grayscale(load.image(photos[j]))</pre>
        pic <- resize(pic, n, n)
        pic <- matrix(pic,nrow=n)[-c(c(1:floor(0.25*n)),c(floor(0.75*n+1):n)),-c(c(1:floor(0.25*n)),c(floor(0.75*n+1):n))]
        vector <- matrix(pic, ncol=1)[,1]</pre>
        X <- cbind(X, vector)</pre>
  return(X)
```

- PCA
- Check for importance

```
faces_pca <- prcomp(t(matrix_images), center = FALSE, scale. = FALSE)
importance <- as.data.frame(summary(faces_pca)$importance)
imp val <- check importance(importance)</pre>
```

Create a response variable

Split data

```
data_vec <- 1:length(mydata[,1])
train_vec <- data_vec[data_vec%%60 <= 48 & data_vec%%60 != 0]
val_vec <- data_vec[data_vec%%60 <= 54 & data_vec%%60 > 48]
test_vec <- data_vec[data_vec%%60 <= 60 & data_vec%%60 > 54 | data_vec%%60 == 0]
train <- mydata[train_vec,]
val <- mydata[val_vec,]
test <- mydata[test_vec,]</pre>
```

Train model

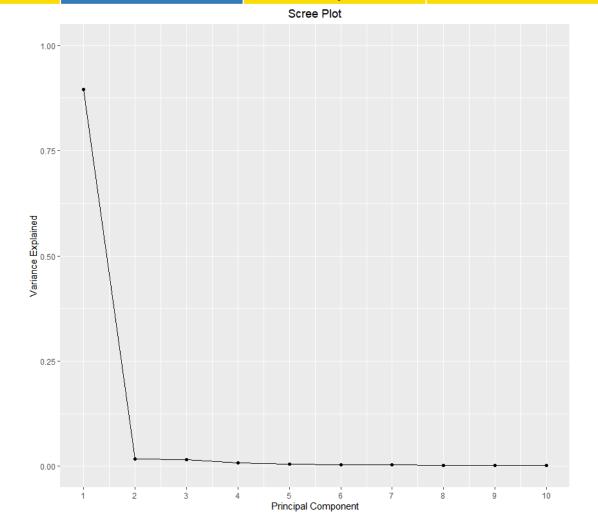
```
accuracy <- c()
max <- 0
b_gamma <- 0
gammas <- c(0.1,0.01,0.001,0.0001,0.00001,0.000001)
for (i in 1:length(gammas))
  classifier <- svm(y ~ ., data = train, gamma = gammas[i], kernel = "radial")</pre>
  prediction <- predict(classifier, newdata = val)</pre>
  m <- sum(prediction == val$y)/length(prediction)</pre>
  accuracy <- c(accuracy, m)
  if (m == max(accuracy))
    max <- m
  if (i == length(gammas))
    print(paste0("Best gamma is ", gammas[i]))
    b_gamma <- gammas[i]
```

Validate

```
classifier <- svm(y ~ ., data = train, gamma = b_gamma, kernel = "radial")
prediction <- predict(classifier, newdata = val)</pre>
```

Results of PCA

- There is a massive dropoff in explained variance as we increase the number of principal components
- Only 104 of the 480 PCs are needed to explain over 99% of the variance in the data!



Results of SVM

- At first our model struggled to classify accurately more than 40-50% of the time
- If we crop the pictures to the center, most of the images would be just the face and have less background to throw our model off









Results of SVM

- We split our 480 images into 80% training, 10% validation, and 10% testing
- Created multiple SVM models with different kernel parameters (with radial kernel type)
- Chose the model with the highest accuracy in the validation set as our model

Results of Model

Example of model prediction with PCA



Correct: George_W_Bush
Prediction: George_W_Bush



Correct: George_W_Bush
Prediction: Tony_Blair

Results of Model

 There was a massive difference in accuracy when using PCA compared to not using PCA

	Validation Accuracy	Testing Accuracy
With PCA	72.9%	70.8%
Without PCA	22.9%	14.6%

Results of validation classification

	folders <chr></chr>	gender <chr></chr>	Accuracy (%)	Predicted individual as	X2 <chr></chr>	X3 <chr></chr>	X4 <chr></chr>
11	Colin_Powell	M	66.67	Donald_Rumsfeld	Tony_Blair	NA	NA
13	Donald_Rumsfeld	M	83.33	Colin_Powell	NA	NA	NA
15	George_W_Bush	M	66.67	Donald_Rumsfeld	Hugo_Chavez	NA	NA
16	Gerhard_Schroeder	M	33.33	Donald_Rumsfeld	George_W_Bush	Hugo_Chavez	Hugo_Chavez
23	Hugo_Chavez	M	66.67	George_W_Bush	Tony_Blair	NA	NA
62	Tony_Blair	M	66.67	George_W_Bush	George_W_Bush	NA	NA

- Total Accuracy of the algorithm for validation 72.92%
- The name that was predicted the most was George B with a count of 4
- The highest number of inaccurate predictions per person was: 4

Results of testing classification

	folders <chr></chr>	gender <chr></chr>	Accuracy (%)	Predicted individual as	X2 <chr></chr>	X3 <chr></chr>	X4 <chr></chr>
6	Ariel_Sharon	M	83.33	Colin_Powell	NA	NA	NA
11	Colin_Powell	M	33.33	Ariel_Sharon	George_W_Bush	Gerhard_Schroeder	Junichiro_Koizumi
15	George_W_Bush	M	83.33	Tony_Blair	NA	NA	NA
16	Gerhard_Schroeder	M	50.00	Donald_Rumsfeld	Hugo_Chavez	Tony_Blair	NA
23	Hugo_Chavez	M	66.67	Junichiro_Koizumi	Tony_Blair	NA	NA
39	Junichiro_Koizumi	M	83.33	Ariel_Sharon	NA	NA	NA
62	Tony_Blair	M	66.67	Colin_Powell	Gerhard_Schroeder	NA	NA

- Total Accuracy of the algorithm for testing 70.83%
- The name that was predicted the most was Tony Blair with a count of 3
- The highest number of inaccurate predictions per person was: 4

Areas of Improvement



- NONE, WE ARE PERFECT 99
- Only had 8 people with over 60 images
 - More people with a higher quantity of images might increase accuracy
- Crop to faces better
 - Our cropping took the center of the image, some faces weren't centered
- To test other predicting/classification methods
 - logistic regression, K-nearest neighbors, or convolutional neural networks

Conclusion

- Converted 480 grayscale images from 8 separate people into column vectors and created a matrix of images
- Ran PCA on this matrix and reduced our dimensions of the model from 480 to 104
- Used SVM to create a model classifying an image
- Our model had 70.8% accuracy in classifying faces



Correct: George_W_Bush
Prediction: George_W_Bush



References

- 1. https://royalsocietypublishing.org/doi/10.1098/rsta.2015.0202
- 2. http://vis-www.cs.umass.edu/lfw/
- 3. https://en.wikipedia.org/wiki/File:SVM_margin.png
- 4. GitHub

Libraries

- rcpp
- imager

- tictoc
- ggplot2

- magick
- e1071

Questions?