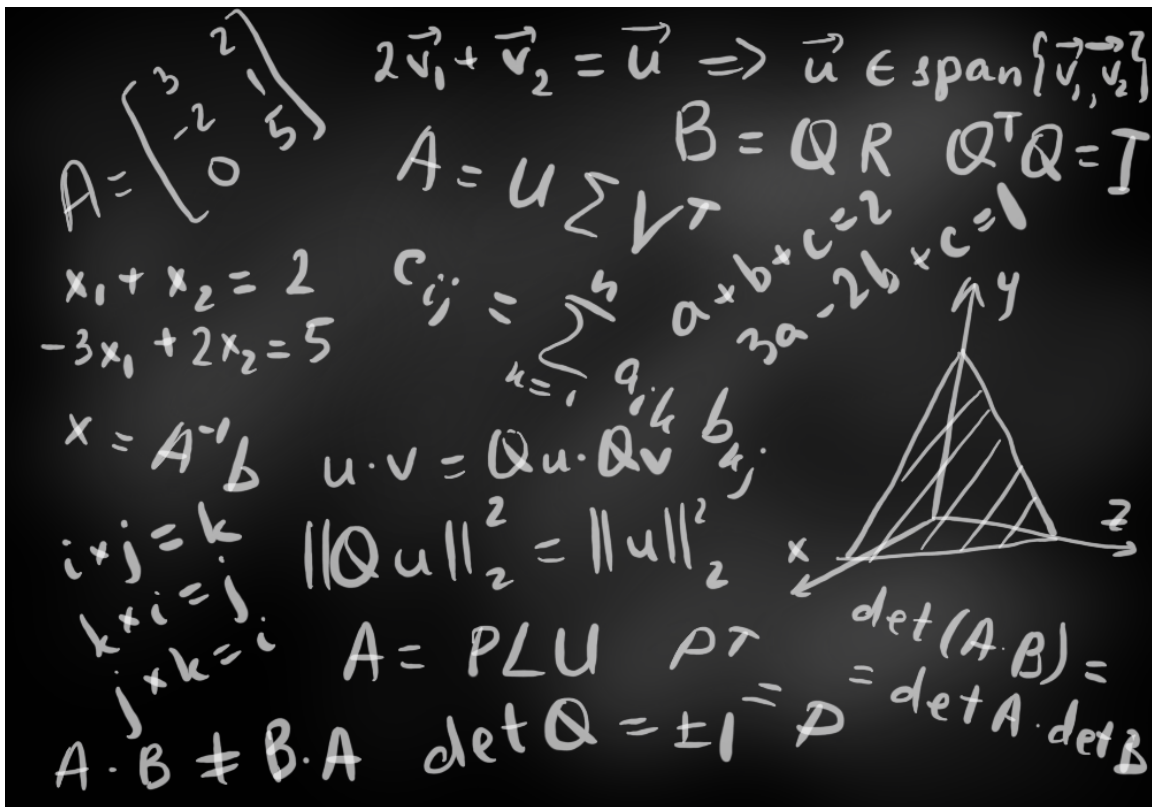
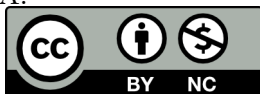


# APPLIED PROJECTS FOR AN INTRODUCTORY LINEAR ALGEBRA CLASS

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## About this book

This book is based on the Applied Matrix Theory curriculum which the author taught at Kansas State University. Applied Matrix Theory is an introductory linear algebra undergraduate class geared primarily towards engineering students. Unlike a proof-based class that is typically offered to mathematics majors, the focus here is to introduce real life applications at the same time as teaching the standard topics of linear algebra such as matrix operations, linear systems, inverses, vector spaces, determinants, and eigenproblems. While teaching the class, the author of this book struggled with finding realistic applications of the theoretical material which can be reasonably completed during a class period. The search for real life applications has led the author to write this book which hopefully fills the gap between linear algebra theory and applications. Most of the projects in this book can be done with minimal knowledge of programming. It is author's hope that the projects will encourage the exploration of data using simple concepts from linear algebra. The projects are written in Matlab in order to harness the abundance of built in functions and computational power of Matlab. The author attempted to incorporate real data from various open sources into these projects. There are two labs which are dedicated to exploring operations with matrices and to solving linear systems in Matlab (the projects 1 and 4). There are also two labs which use more advanced material typically not covered in an introductory linear algebra class, such as singular value decomposition and normal equations (the projects 8 and 14).

The code in this book has been developed using the 2015a version of Matlab and tested on Matlab 2019a. Depending on the version of Matlab on your computer the syntax and the output might differ slightly. The projects in this book can be adapted to use with GNU Octave as well. The supplementary files necessary to complete some of the labs are available from the author's webpage at <http://www.math.ksu.edu/~azem>.

A variety of printed and Internet sources inspired this book. These sources are acknowledged in the Reference section as well as the sources of data for the projects. This book has been developed under the Open/Alternative Textbook Initiative grant from Kansas State University, the generous support of which is gratefully acknowledged here. The author is immensely grateful to her colleagues, Nathan Albin and Pietro Poggi-Corradini, for proofreading the book and for the corrections. The author is grateful to Jeffrey Sheldon for proofreading the book and suggesting style corrections.

## A brief introduction to Matlab

MATLAB is a numerical computing software application and a high level programming language developed by MathWorks, Inc. Initially, Matlab stood for “matrix laboratory” and was developed specifically to perform numerical linear algebra operations. Currently Matlab has a vast library of built in mathematical functions which allow users to perform complex computations without having to program numerical methods themselves. Because of this Matlab is the software of choice for many scientific and engineering applications.

In this book we will use Matlab to illustrate common linear algebra concepts and to perform computations on problems with real life applications. Before we go any further, let us review some basic commands and principles of using Matlab. For more information, please see Matlab Help or any Matlab manual of your choice.

### The Matlab desktop environment

Matlab’s desktop environment typically consists of several windows. By default, you are likely to see the command window with a prompt. There is also usually the “Current Folder” window which shows the directory on the computer you are working with and the “Workspace” window with any variables which have been initialized (if you did not define any variables yet this window will be empty). In some cases you can also see the “Command History” window which shows the commands which have been executed recently. The default configuration can be changed by clicking downward arrow in the upper right corner of each of these windows or by using the switch windows button located in the upper right corner of the main Matlab window. Matlab can be used by typing commands in a command prompt or by executing text files with commands. These files are called Matlab scripts and have the extension “.m”. Matlab comes with a built in editor for editing M-files. This editor can also be one of the windows you see in your Matlab environment when you start Matlab.

### The command line

The easiest way to get started with Matlab is by executing commands in the interactive command window. The command prompt can be seen in the main window of the Matlab environment:

```
>>
```

A Matlab command can be entered in this prompt and Matlab will produce an immediate result. For instance, entering

```
>> sin(pi/3)
```

will generate the output

```
ans =  
    0.8660
```

To suppress Matlab output you can put a semicolon at the end of your command. For instance, the command

```
>> sin(pi/3);
```

produces no output. However, observe that the last output which has not been assigned to any variable is stored in the Matlab variable **ans**. This variable has now changed its value to  $\sin \pi/3$  rounded up to the machine precision.

By default Matlab uses the short format for output on the screen which is given by the Matlab command `format short`. This command generates an output for real numbers with up to 4 decimal digits. The output format can be changed by executing the command

```
format long
```

The long format produces output for real numbers with up to 15 decimal digits.

If you want to know more about a particular command, you can always type

```
help <name of the command>
```

or

```
doc <name of the command>
```

This will open Matlab Help on the command you are interested in and also produce various examples of Matlab's use of this command. You can also open Matlab help directly by pressing the corresponding button in the Matlab environment. This button looks like a question mark and is generally located in the upper right corner of the Matlab "Home" menu.

Another useful tip to remember is that pressing "Up" arrow button on your keyboard while in the command line will bring up previous commands you have executed. You can scroll through the commands and select any of them to run again. This is especially useful if you have been typing lengthy expressions in the command line and need to correct misprints or simply need to run the command again.

Finally, sometimes it is necessary to interrupt Matlab computations. This might happen, for instance, if Matlab is stuck in an infinite loop or a computation simply takes too long to finish. You can interrupt the execution of Matlab code by pressing "Ctrl-C" while in the main Matlab window.

## Matlab scripts

While simple computations can be handled by typing commands directly into the command prompt, it is advisable to organize more complex computations in a form of Matlab scripts or M-files. You can create a new script file by going to the menu "Home" of the main Matlab window and clicking on the "New Script" button. This will open a new empty script file in the Matlab editor.

It is a good idea to keep your M-files well-documented by utilizing the comment option in Matlab. You can create a comment in a script file by starting the code line with the percentage symbol `%`. While interpreting the script, Matlab will ignore everything in the line after the `%` symbol.

You can also create code cells within the script. A new code cell can be started by using the double percentage symbol `%%`. You can also write a title of the cell after the `%%` sign. Matlab will treat everything after the `%%` sign as a comment. The creation of code cells allows you to break your code into smaller executable pieces and run those pieces individually. To do so you can use the "Run and Advance" button at the top of the Matlab editor window. Generally speaking, you want to split the code into code cells in some logical way. For instance, each code cell can be dedicated to performing a particular function in your code.

Finally, you can publish your code using the "Publish" section of the menu of the Matlab editor. By default, code is published in the HTML format. Publishing will create a document which involves a table of contents, the code itself, and all the plots and output which have been produced by your code.

## Variables, vectors, matrices

Variables are used to store and manipulate the values obtained during a Matlab session. Matlab uses the following standard types of variables: real which can be double and single precision (**double** or **single**), several integer types (**int8**, **int16**, **int32**, **uint8**, **uint16**, **uint32**, **uint64**), boolean (**logical**) and character (**char**).

By default all the numbers in Matlab are treated as the type **double** which uses 64 bits to store a real number.

Each variable needs to have an identifier or a name. There are some general considerations for variable names in Matlab:

- The name of a variable must start with a letter. After the first character the name may contain letters, numbers, and the underscore character.
- Variable names in Matlab are case-sensitive. This means that **Price** and **price** are treated as different variables.
- Certain words (such as reserved words or keywords) cannot be used as variable names.
- Names of built-in functions can, but generally should not, be used as variable names.

Matlab has several different ways to define matrices (two-dimensional arrays) and vectors (one-dimensional arrays). For instance, we can define arrays explicitly by typing in all the values:

```
>> vec=[1 2 3 4]
vec =
     1     2     3     4
>> mat=[1 2 3; 4 5 6]
mat =
     1     2     3
     4     5     6
```

In the example above a  $1 \times 4$  vector **vec** and a  $2 \times 3$  matrix **mat** are defined. Another way to define vectors and matrices is by using the colon operator “:”. For instance, the variables **vec** and **mat** from the example above can be also defined by commands

```
>> vec=1:4;
>> mat=[1:3;4:6];
```

We will explore other matrix operations further in this book. Matlab also has an ability to work with multi-dimensional arrays which will be useful for us later when manipulating color images.

Matlab has a set of operators to perform standard algebraic operations with variables:

Operator	Meaning
+, −	addition, subtraction, binary
−	negation, unitary
*, \	multiplication, division, binary
^	power
()	parentheses (change of the order of operations)

Keep in mind the priority in which these operators are executed: the parentheses **()** take the highest priority, followed by the power operator **^**, unitary negation **−**, multiplication and division **\***, **\**, and, finally, addition and subtraction **+**, **−**.

## Matlab functions

Matlab has a large library of built-in functions which makes it a convenient environment for many scientific and engineering computations. The full list of available functions can be found in Matlab Help and on the [mathworks.com](http://mathworks.com) webpage. You can also look into the “Help” menu of the Matlab main window (shown as a question mark in the upper right corner). The list of elementary functions can be viewed by executing the command

```
>> help elfun
```

from the command line.

Matlab also allows user to create their own user-defined functions which can be saved in separate M-files and can be later called from other scripts. The syntax of such functions in its simplest form looks like this:

```
function [OutputVariables] = FunctionName(InputVariables)
% Description of the function
    <Main body of the function>
end
```

## Control statements

There are several standard control statements in Matlab which change the order in which the commands of a script are executed or whether they are executed at all.

The control statements can be roughly divided into condition statements and loop statements. The first group of control statements we will look at is the conditional statements. The conditional statements allow one to perform different computations or procedures depending on whether the condition (which is given by a boolean expression) is true or false. In the simplest form, the conditional statement looks like this:

```
if <condition>
    <statements>
else
    <statements>
end
```

In the above construction, **else** statements can be skipped. Additionally, more options can be incorporated by using the **elseif** clause. A construction of this type has the form:

```
if <condition1>
    <statements1>
elseif <condition2>
    <statements2>
elseif <conditions3>
    <statements3>
...
else
    <statements>
end
```

The **switch** statement can also be used in place of **if...elseif** statements which use many **elseif** clauses. Conditions in each of the conditional statements represent logical (boolean) expressions which can be either true or false.

The second group of control statements consists of loop statements. A simple **for** loop looks like this



```

for LoopVariable = Range
    <statements>
end

```

Range here consists of all the values through which the `LoopVariable` will iterate. For instance, in the following code

```

for i=1:3
    fprintf('%d\n',i)
end

```

the loop variable `i` iterates through the integers 1, 2, and 3, and the code itself prints the integers from 1 to 3 in a column format.

Another loop statement in Matlab is the **while** loop. The general form of a **while** statement looks like this:

```

while <condition>
    <statements>
end

```

The `<statements>` part of the code is executed as long as the conditional statement given by `<condition>` remains true.

## Logical statements

Expressions which are either true or false are called boolean or logical expressions. The logical operators and relational operators in Matlab are given in the table below:

Operator	Meaning
> (<)	greater than (less than)
>= (<=)	greater or equal (less or equal)
==	logical equality
~=	logical inequality
	or
&&	and
~	not
<b>xor</b>	exclusive or

The difference between the logical equality `==` and the assignment operator `=` is that the assignment operator assigns the value on the right side of the operator to the variable on the left side of the operator. At the same time, the logical equality operator checks whether the two values on the right and the left sides of the operator are equal.

When we compare two expressions using relational operators the result is a boolean which can be true or false. For instance,

```

>> 3>2
ans =
    1

```

The variable `ans` here is of **logical** type.

## Input and output functions

The simplest way to create a user defined variables in Matlab is provided by the function `input`. The `input` function displays a prompt string on the screen and awaits for user input. For instance,

```
>> side=input('Enter the side of the square: ')
Enter the side of the square: 3
side =
    3
```

The code above asks to input a number (the side of the square) and stores it in the variable `side`. It is generally a good idea to give the user an explanation of what is expected from them. The string `'Enter the side of the square: '` above provides such an explanation.

Formatted output can be achieved in Matlab by using the `fprintf` function. For instance,

```
>> fprintf('The area of the square is equal to %d\n',side^2)
The area of the square is equal to 9
```

The expression `%d` here is a placeholder for the variable written after the string which, in this case, is the value `side2`. The symbol `\n` indicates that any output after this command should start with a new line.

More information about the functions `input` and `fprintf` can be obtained by calling Matlab Help with the `help input` or `help fprintf` commands.

## Plotting graphs

A standard way to display information in the form of a two-dimensional graph in Matlab is provided by the `plot` function. In the simplest form, the function `plot` looks like this:

```
>> plot(x_vector,y_vector,'linestyle')
```

Here, the variables `x_vector` and `y_vector` are the vectors which contain, correspondingly, the x- and y-coordinates of the points on the graph. The `'linestyle'` is a string variable which contains the parameters for determining the type of line used for the graph. For instance, `'k-o'` means that the graph will be drawn with black (`'k'`) solid (`'-'`) line and each data point will be shown with an empty circle (`'o'`).

It is possible to plot multiple graphs on the same figure at the same time:

```
>> plot(x1_vector,y1_vector,'linestyle1',x2_vector,y2_vector,'linestyle2', ...)
```

Alternatively, it is possible to add more graphs to the figure later by using the `hold on`, `hold off` commands. If you don't use the command `hold on` after the graph, calling the `plot` function again will plot your graph in the previous figure window, erasing everything which was on that figure before. If you would like a new plot to be shown in a new figure window without erasing any of the previous plots, you can create a new figure window by using the `figure` command. Finally, Matlab allows figures to be grouped together in the form of subfigures. This can be achieved by using the `subplot` command.

## Further reading

This introduction is designed to touch upon some of the basics of Matlab and is not meant to be exhaustive. Some of the concepts discussed here will be applied in the sections to follow. There are many other sources of additional information about Matlab, which include Matlab Help, the [mathworks.com](http://mathworks.com) website, and various Matlab manuals which can be purchased separately.

## Advice on good programming style

Usually the goal of a code developer is not only to write working code but also to make the code easily understandable to other people. Here are several fairly standard tips which the author found useful on how to write clean, easily readable, code:

1. Choose meaningful names for your variables and functions. This will make your code significantly easier to read. For instance, the meaning of the variables `Height` and `Weight` is much clearer than non descriptive `h` and `w`. If the variable name consists of several words, such as `carspeed` and `priceperitem`, then capitalizing each word (for instance, `CarSpeed` and `PricePerItem`) or using underscores (`car_speed` and `price_per_item`) can increase the readability of your code.
2. If your code is long, break it into smaller pieces. Give each piece only one function. In Matlab, you can create user-defined functions and code cells for this purpose. User defined functions can be created by using the `function` keyword (type `doc function` in the Matlab command line to learn more about it). Code cells can be created by entering the double percentage sign `%%` into your code. Each code cell can then be run individually by using the “Run and Advance” button in the Matlab editor.
3. Add comments to your code by using the single percentage sign `%`. Matlab will ignore the rest of the line after the percentage sign. Explain the purpose of the commands used, variables, and functions created in the comments. You will thank yourself for doing this when you look at your code six months later. Anybody else trying to understand your code will thank you too.
4. Don’t write dense code. Allow empty lines between the commands.
5. Use consistent indentation. This makes reading code with multiple cycles much easier. Compare the two examples in the Fig. 1. Which one is easier to read?

```
for i=1:10
    for j=1:10
        Sums(i,j)=i+j;
        Products(i,j)=i*j;
    end;
end;

for i=1:10
    for j=1:10
        Sums(i,j)=i+j;
        Products(i,j)=i*j;
    end;
end;
```

Figure 1: An example of indentation usage.

6. Don’t make your lines too long. Split long commands and computations using triple dots `...`
7. Clear code is usually better than clever code. Sometimes you can use a clever style of programming to compound 5-10 lines of code into one single line. However, this code is often much more difficult to understand.

You will discover many other tips and pointers on your own as you gain more experience with programming.

## Project overview

Each of the projects in this book is designed to test several concepts from linear algebra. The goals of each project and linear algebra concepts used in it are summarized below:

**Project 1:** To introduce operations with matrices in Matlab such as matrix addition, scalar multiplication, matrix multiplication, and matrix input functions in Matlab.

**Project 2:** To use the matrix operations introduced in the previous project to manipulate images. Lightening/darkening, cropping, and changing the contrast of the images are discussed.

**Project 3:** To use matrix multiplication to manipulate image colors similarly to photo filters for images.

**Project 4:** Introduction to solving linear systems in Matlab.

**Project 5:** To discuss some of the strategies for the ranking of sports teams with an example of the Big 12 conference in college football.

**Project 6:** To apply convolution of two matrices to blurring/sharpening of images.

**Project 7:** To create a simple recommender system based on the use of norms and inner products.

**Project 8:** To study basic interpolation techniques and the least squares method with application to climate data.

**Project 9:** To use orthogonal matrices for the rotation of 3D objects.

**Project 10:** To apply matrix mappings in order to generate fractals with the help of Chaos game.

**Project 11:** To apply eigenvalue problems and orthogonal projections in order to create a simple face recognition algorithm.

**Project 12:** To introduce Google's PageRank algorithm and look at the eigenproblems for the ranking of webpages.

**Project 13:** Application of eigenvalues and eigenvectors for the data clustering with the example of Facebook network.

**Project 14:** Application of singular value decomposition for image compression and noise reduction.

Sample questions to check students' understanding and checkpoints for introduced variables are highlighted in the text of the projects in yellow. The templates of the labs 9-14 are given in the appendix.

## Project 1: Basic operations with matrices in Matlab

**Goals:** To introduce matrix input functions and basic operations with matrices in Matlab. The matrix operations to be studied include matrix addition and subtraction, scalar product, matrix product and elementwise matrix product in Matlab, matrix concatenation, and selecting submatrices.

**To get started:** Create a new Matlab script file and save it as `lab01.m`.

**Matlab commands used:** `length`, `linspace`, `size`, `max`, `min`, `mean`, `sum`, `randi` (`rand`), `'`, `*`, `.*`, `eye`, `zeros`, `ones`, `diag`, `spdiags`.

**What you have to submit:** The file `lab01.m`, which you will create during the lab session.

### INTRODUCTION

It is assumed that the reader is familiar with basic matrix operations and their properties, so the definitions and facts here are given mostly for the sake of a review.

We can think of a matrix as a table of numbers:

$$\mathbf{A} = \begin{bmatrix} a_{11} & a_{12} & \dots & a_{1j} & \dots & a_{1n} \\ a_{21} & a_{22} & \dots & a_{2j} & \dots & a_{2n} \\ \dots & \dots & \ddots & \dots & \dots & \dots \\ a_{i1} & a_{i2} & \dots & a_{ij} & \dots & a_{in} \\ \dots & \dots & \dots & \dots & \ddots & \dots \\ a_{m1} & a_{m2} & \dots & a_{mj} & \dots & a_{mn} \end{bmatrix}$$

The matrix  $\mathbf{A}$  above contains  $m$  rows and  $n$  columns. We say that  $\mathbf{A}$  is an  $m \times n$  ( $m$ -by- $n$ ) matrix, and call  $m, n$  the dimensions of the matrix  $A$ . The matrix  $A$  above can also be compactly written as  $\mathbf{A} = (a_{ij})$ , where  $a_{ij}$  is called  $(i, j)$  element of the matrix  $\mathbf{A}$ .

Matlab (originally named “MATrix LABoratory”) has an extensive list of functions which work with matrices. The operations discussed in this project consist of matrix addition/subtraction, multiplication of a matrix by a constant (scalar multiplication), transposition of a matrix, selecting submatrices, and concatenating several smaller matrices into a larger one, matrix multiplication and componentwise matrix multiplication, and operations of finding maximal/minimal element in the matrix, and finding the sum/mean in the row/column of the matrix.

*Scalar multiplication* of a matrix  $\mathbf{A}$  by a constant  $c$  consists of multiplying every element of the matrix by this constant:

$$c\mathbf{A} = (ca_{ij}).$$

*Matrix addition and subtraction* are possible if the matrices have the same dimensions and is done componentwise:

$$\mathbf{A} \pm \mathbf{B} = (a_{ij} \pm b_{ij}).$$

If  $\mathbf{A}$  is an  $m \times n$  matrix then the transpose  $\mathbf{A}^T$  ( $\mathbf{A}'$  in Matlab) of the matrix  $\mathbf{A}$  is the  $n \times m$  matrix obtained by flipping  $A$  along its diagonal entries. Thus every column of  $\mathbf{A}$  becomes a row of  $\mathbf{A}^T$ , and every row of  $\mathbf{A}$  becomes a column of  $\mathbf{A}^T$ . For instance,

$$\begin{bmatrix} 1 & 2 & 3 \\ 4 & 5 & 6 \end{bmatrix}^T = \begin{bmatrix} 1 & 4 \\ 2 & 5 \\ 3 & 6 \end{bmatrix}.$$

If  $n$  is a positive integer then the  $n \times n$  identity matrix, denoted by  $\mathbf{I}_n$ , is a matrix which has 1 in every diagonal entry  $(i, i)$  for  $1 \leq i \leq n$ , and 0 in every other entry. There is one identity matrix for every dimension  $n$ . For instance,

$$\mathbf{I}_2 = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} \quad \mathbf{I}_3 = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$$

are the  $2 \times 2$  and  $3 \times 3$  identity matrices.

The identity matrices are special examples of diagonal matrices. A matrix  $\mathbf{A}$  is diagonal if  $A(i, j)$  is 0 whenever  $i \neq j$  (it is not required that  $a_{ii} \neq 0$  for any  $i$ ).

A matrix product  $\mathbf{C} = \mathbf{AB}$  of an  $m \times n$  matrix  $\mathbf{A}$  and an  $n \times p$  matrix  $\mathbf{B}$  exists if and only if  $n = l$ . If this condition is satisfied, then  $\mathbf{C}$  is an  $m \times p$  matrix with the elements

$$c_{ij} = \sum_{k=1}^n a_{ik}b_{kj}.$$

Observe that matrix multiplication is non-commutative,  $\mathbf{AB} \neq \mathbf{BA}$ .

For example if  $\mathbf{A} = \begin{bmatrix} 2 & 3 \\ 1 & 0 \\ -1 & -1 \end{bmatrix}$  and  $\mathbf{B} = \begin{bmatrix} 4 & 0 & 2 \\ 3 & -1 & 1 \end{bmatrix}$  then the product  $\mathbf{AB}$  is given by

$$\mathbf{AB} = \begin{bmatrix} (2 \cdot 4 + 3 \cdot 3) & (2 \cdot 0 + 3 \cdot -1) & (2 \cdot 2 + 3 \cdot 1) \\ (1 \cdot 4 + 0 \cdot 3) & (1 \cdot 0 + 0 \cdot -1) & (1 \cdot 2 + 0 \cdot 1) \\ (-1 \cdot 4 + -1 \cdot 3) & (-1 \cdot 0 + -1 \cdot -1) & (-1 \cdot 2 + -1 \cdot 1) \end{bmatrix} = \begin{bmatrix} 17 & -3 & 7 \\ 4 & 0 & 2 \\ -7 & 1 & -3 \end{bmatrix}.$$

If  $\mathbf{C} = \begin{bmatrix} 2 & 1 \\ 1 & 7 \end{bmatrix}$  and  $\mathbf{D} = \begin{bmatrix} 1 & 2 & 3 \\ 0 & 1 & 2 \\ 0 & 0 & 1 \end{bmatrix}$  then the product  $\mathbf{CD}$  is not defined because the number of columns of  $\mathbf{C}$  does not equal the number of rows of  $\mathbf{D}$ .

## TASKS

1. Open the lab01.m Matlab script you have created and type in the following commands:

```
%% basic operations with matrices
A=[1 2 -10 4; 3 4 5 -6; 3 3 -2 5];
B=[3 3 4 2];
```

This will create a new cell and define two new variables A and B. The variable A stores a  $3 \times 4$  matrix, while the variable B contains a  $1 \times 4$  row vector. Use the “Workspace” window of the Matlab environment to view the variables A and B.

Variables: A, B

2. Type in the following commands:

```
lengthA = length(A)
lengthB = length(B)
```

Variables: lengthA, lengthb

Q1: What does the command `length(A)` do?

3. Add the vector **B** as the fourth row of the matrix **A** and save the result as a new matrix **C** using the following code:

```
C=[A; B];
```

Variables: **C**

4. Create a new matrix **D** whose entries are located in the rows 2,3,4 and columns 3,4 of the matrix **C**:

```
D=C(2:4,3:4);
```

Variables: **D**

5. Create a transpose of the matrix **D** by using a quotation mark sign ' in Matlab and save the transposed matrix as the matrix **E**.

Variables: **E**

6. Check the size of the matrix **E** by using the Matlab **size** function and store the result of this operation as two variables **m**, **n**.

Variables: **m**, **n**

7. Matlab allows for the creation of equally spaced vectors which can be useful in many situations. For instance, run the following code:

```
EqualSpaced=0:pi/10:2*pi  
EqualSpaced1=linspace(0,2*pi,21)
```

Take a look at the output produced by Matlab. Observe that both of these commands created the vector with equally spaced entries ranging from 0 to  $2\pi$  and the distances between the elements equal to  $\pi/10$ .

Variables: **EqualSpaced**, **EqualSpaced1**

8. Create a new cell and find the maximal and minimal elements in each of the columns of the matrix **A**. Store the result respectively in the row vectors named **maxcolA** and **mincolA**. You can use the Matlab **max** and **min** functions:

```
%% Max, min, sum and mean  
maxcolA=max(A)  
mincolA=min(A)
```

Variables: **maxcolA** and **mincolA**

9. Now repeat the previous step but find the maximal and the minimal elements in each row of the matrix **A**. Observe that there are at least two ways to approach this: you can either use the **max** and **min** functions on the transpose of **A** or use these functions with an additional argument 2 to show that you are interested in the maximum and minimum along the second dimension (i.e. **max(A,[],2)**). Save the results as **maxrowA** and **minrowA**. Finally find the maximal and minimal elements in the whole matrix **A**, and save the results as **maxA** and **minA**.

Variables: **maxrowA**, **minrowA**, **maxA** and **minA**

10. Repeat the last two steps using the commands `mean` and `sum` instead of `max` and `min`. You should create six new variables, corresponding to the column/row means/sums, and the mean/sum of the elements of the entire matrix.

Variables: `meancolA`, `meanrowA`, `sumcolA`, `sumrowA`, `meanA`, `sumA`

Q2: What do the commands `mean` and `sum` do?

11. Create a new cell and use the command `randi([-4 4],5,3)` to create two matrices `F` and `G` with 5 rows and 3 columns and random integer entries from  $-4$  to  $4$ :

```
% Matrix addition/subtraction, and multiplication
F=randi([-4 4],5,3)
G=randi([-4 4],5,3)
```

Variables: `F`, `G`

12. Perform the operations `0.4*F`, `F+G`, `F-G`, `F.*G`, storing the results in `ScMultF`, `SumFG`, `DiffFG`, and `ElProdFG` respectively.

Variables: `ScMultF`, `SumFG`, `DiffFG`, `ElProdFG`

Q3: What does the last operation do?

13. Check the size of `F` and the size of `A`, storing the results in the variables `sizeF` and `sizeA`, respectively.

Variables: `sizeF`, `sizeA`

Q4: Can we matrix-multiply `F` and `A`? Explain.

14. Compute `H=F*A`.

Variables: `H`

Q5: What are the dimensions of `H`?

15. Generate the identity matrix `eye33` with 3 rows and 3 columns using the Matlab `eye` command.

Variables: `eye33`

16. Run the commands:

```
zeros53=zeros(5,3);
ones42=ones(4,2);
```

Variables: `zeros53`, `ones42`

Q6: What do the functions `zeros` and `ones` do?

17. Generate the diagonal  $3 \times 3$  matrix `S` with the diagonal elements  $\{1, 2, 7\}$  using the Matlab `diag` function.

Variables: `S`

18. Now let us do the opposite: extract the diagonal elements from the matrix and save them in a separate vector. The same function `diag` accomplishes that as well:

```
R=rand(6,6)
diagR=diag(R)
```



This creates the  $6 \times 6$  matrix **R** with random entries from the interval  $(0,1)$ , extracts the diagonal entries from it, and saves them in the  $1 \times 6$  vector **diagR**. Observe that the function **diag** has other interesting functions. To learn more about it type **help diag** in the command line.

Variables: **R,diagR**

19. Another function which allows to create diagonal matrices is **spdiags**. Technically, this function creates a sparse matrix which is a matrix with a large number of zero elements. These matrices are stored in a special form in Matlab with only non-zero elements are stored, and operations with them are also done in a special way. To convert sparse form of the matrix into the regular one, the command **full** can be used. Run the following code:

```
diag121=spdiags([-ones(10,1) 2*ones(10,1) -ones(10,1)],[-1 0 1],10,10);  
full(diag121)
```

Variables: **diag121**

Q7: What does this code do?

## Project 2: Matrix operations and image manipulation

**Goals:** To apply elementary matrix operations such as matrix addition, subtraction, and elementwise multiplication to image manipulation. We will look at the lightening/darkening of images, selecting subimages, changing contrast, rotating and flipping images.

**To get started:**

- Create a new matlab script and save it as `lab02.m`.
- Download the image file `einstein.jpg` and save it in your working directory <sup>1</sup>.

**Matlab commands used:** `imread`, `imshow`, `uint8`, `double`, `flip`, `size`, `'`, `zeros`, `rot90`, `floor`, `ceil`, `round`, `fix`.

**What you have to submit:** The file `lab02.m`, which you will create during the lab session.

### INTRODUCTION

An image in a computer memory can be stored as a matrix with each element of the matrix representing a pixel of the image and containing a number (or several numbers) which corresponds to the color of this pixel. If the image is a color image, then each pixel is characterized by three numbers corresponding to the intensities of Red, Green, and Blue (the so-called RGB color system). If the image is a grayscale image, then only one number for the intensity of gray is needed. The intensity of each color typically ranges from 0 (black) to 255 (white). This allows for the representation of over 16 million colors in the RGB system which is more than a human eye can distinguish. The images stored in this way are called bitmaps (there are also images which do not use pixels, such as vector graphics, but we will not talk about those here).

If you have ever used an image editor before, you may know that there are several fairly standard operations which you can perform with images. In this project we will replicate some of them using operations on matrices. By the end of the project you will understand how graphic editors process images to achieve different visual effects. As it turns out, many of these operations are accomplished by manipulating the values of the pixel colors.

### TASKS

1. To begin let us load the image into Matlab. This can be done by using the Matlab command `imread`. This command allows Matlab to read graphic files with different extensions. The output is an array with the dimensions equal to the dimensions of the image, and the values corresponding to the colors of the pixels. Here we will work with a grayscale image, so the elements in the matrix will be integers ranging from 0 to 255 in the Matlab integer format `uint8`. Type in the following code to load the file “`einstein.jpg`” into Matlab:

```
%% Load a grayscale jpg file and represent the data as a matrix
ImJPG=imread('einstein.jpg');
```

---

<sup>1</sup>The image is available at: [https://commons.wikimedia.org/wiki/File:Albert\\_Einstein\\_Head.jpg](https://commons.wikimedia.org/wiki/File:Albert_Einstein_Head.jpg) and has no known copyright restrictions. You can also use an alternative .jpg grayscale image of your choice. The answers and the variable values will differ from what is shown here in that case.

The array `ImJPG` is a two-dimensional array of the type `uint8` which contains values from 0 to 255 corresponding to the color of each individual pixel in the image, where 0 corresponds to black and 255 to white. You can visualize this array by going to the “Workspace” tab of the main Matlab window and double clicking on `ImJPG`.

Variables: `ImJPG`

2. Use the `size` function to check the dimensions of the obtained array `ImJPG`:

```
[m,n]=size(ImJPG)
```

Variables: `m`, `n`

Q1: What are the dimensions of the image?

3. Check the type of the array `ImJPG` by using the command `isinteger`:

```
isinteger(ImJPG)
```

The output of the `isinteger` command is boolean, meaning it will produce either 1 (**true**) or 0 (**false**).

4. Find the range of colors in the image by using `max` and `min` commands and save those elements as `maxImJPG` and `minImJPG`. Observe that execution of this command depends on the version of Matlab on your computer.

Variables: `maxImJPG`, `minImJPG`

5. Finally, display the image on the screen by using `imshow`:

```
imshow(ImJPG);
```

If you did everything correctly, you should see the image on the Fig. 2 displayed on your screen in a separate window.

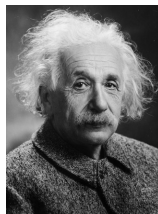


Figure 2: The original image

6. Now we are ready to work with this image. Let us look at some basic image operations we can do using matrices. It is possible to crop the image by selecting a submatrix of the matrix `ImJPG`. Selecting a submatrix can be done simply by naming the rows and columns you want to keep from the initial matrix. For instance, the following commands

```
%% Select and display a part of the image, paste into a different image;
ImJPG_center = ImJPG(100:m-100, 100:n-70);
figure;
imshow(ImJPG_center);
```

will create a new code cell, select the central part of the image leaving out 100 pixels from the top and bottom, 100 pixels on the left and 70 pixels on the right, and display the result in a new figure window.

Variables: **ImJPG\_center**

7. We can paste the selected part of the image into another image. To do this create a zero matrix using the command

```
ImJPG_border=uint8(zeros(m,n));
```

Then paste the preselected matrix **ImJPG\_center** into matrix **ImJPG\_border** and display the image:

```
ImJPG_border(100:m-100, 100:n-70)=ImJPG_center;  
figure;  
imshow(ImJPG_border);
```

Notice the conversion command **uint8**. It is necessary to use this command because by default the array will be of the type **double**, and **imshow** command does not work correctly with this type of array. To see this, try to remove **uint8** and observe what happens when you run the code above.

Variables: **ImJPG\_border**

8. The familiar operation of flipping the image vertically can be easily realized in Matlab by using the matrix command **flip**:

```
%% Rotate and flip the image horizontally/vertically  
ImJPG_vertflip=flip(ImJPG);
```

This command reverses the order of elements in each column of the matrix.

Variables: **ImJPG\_vertflip**

9. Transposing the matrix is achieved by adding an apostrophe ' at the end of the matrix name. Observe that transposing the image matrix is equivalent to rotating the image 90 degrees counterclockwise and flipping it horizontally. Try the following commands:

```
ImJPG_transpose=ImJPG';  
figure;  
imshow(ImJPG_transpose);
```

Variables: **ImJPG\_transpose**

10. Flipping the image horizontally is a little more involved and requires the use of a matrix transposition as well. For instance, this can be accomplished by using the command:

```
ImJPG_horflip=flip(ImJPG_transpose)';
```

Display the resulting matrix and observe what happened to the initial image.

Variables: **ImJPG\_horflip**

11. Apply the command **rot90** to the image using the following code:

```
ImJPG90=rot90(ImJPG)  
figure;  
imshow(ImJPG90)
```

Variables: `ImJPG90`

Q3: What does the command `rot90` do?

12. Try running the following commands:

```
%% Color inversion, lightening/darkening
ImJPG_inv = 255-ImJPG;
figure;
imshow(ImJPG_inv);
```

Display the resulting image in a new figure window. Observe also, that an array `ImJPG` is subtracted here from the constant 255 which mathematically does not make sense. However, Matlab treats the constant 255 as an array of the same size as `ImJPG` with all the elements equal to 255.

Variables: `ImJPG_inv`

Q4: Explain what happened to the image.

13. It is also easy to lighten or darken images using matrix addition. For instance, the following code will create a darker image:

```
ImJPG_dark=ImJPG-50;
```

You can darken the image even more by changing the constant to a number larger than 50. Observe that the command above can technically make some of the elements of the array to become negative. However, because the array type is `uint8`, those elements are automatically rounded to zero.

Variables: `ImJPG_dark`

Q5: How do you lighten the image?

14. Let us create Andy Warhol style art with the image provided. To do so we will arrange four copies of the image into a  $2 \times 2$  matrix. For the top left corner we will take the unaltered image. For the top right corner we will darken the image by 50 shades of gray. For the bottom left corner, lighten the image by 100 shades of gray, and finally, for the bottom right corner, lighten the image by 50 shades of gray. Then we will arrange the images together in one larger matrix using matrix concatenation. Finally, display the resulting block matrix as a single image. If you did everything correctly, you should see an image resembling the Fig. 3.

Variables: `ImJPG_Warhol`

15. Matlab has several commands which allow one to round any number to the nearest integer or a decimal fraction with a given number of digits after the decimal point. Those functions include: `floor` which rounds the number towards negative infinity (to the smaller value), `ceil` which rounds towards positive infinity (to the larger value), `round` which rounds towards the nearest decimal or integer, and `fix` which rounds towards zero.

A naive way to obtain black and white conversion of the image can be accomplished by making all the gray shades which are darker or equal to a medium gray (described by a value 128) to appear as a complete black, and all the shades of gray which are lighter than this medium gray to appear as white. This can be done, for instance, by using the code:

```
%% A naive conversion to black and white
ImJPG_bw=uint8(255*floor(ImJPG/128));
figure
imshow(ImJPG_bw);
```

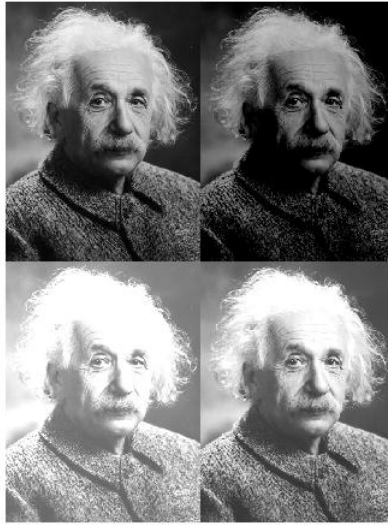


Figure 3: Andy Warhol style art with matrix concatenation

Observe that this conversion to black and white results in a loss of many details of the image. There are possibilities to create black and white conversions without losing so many details. Also notice the function `uint8` used to convert the result back to the integer format.

Variables: `ImJPG_bw`

16. Now let us write code which reduces the number of shades in the image from 256 to 8 by using `round` function. Save your array as `ImJPG8` and display it in a separate window.

Variables: `ImJPG8`

17. The contrast in an image can be increased by changing the range of the possible shades of gray in the image. Namely, we can increase the difference between the colors of the image, for instance, by making the dark shades of gray to appear even darker. One of the simplest ways to do this is to scalar multiply the matrix by some constant. Consider the code:

```
ImJPG_HighContrast=uint8(1.25*ImJPG);
```

Display the image and observe the result. The contrast here can be manipulated by increasing/decreasing the constant (we select 1.25 here). If we select a positive constant greater than 1, then the contrast is increased; if we select a constant less than 1, then it is decreased. Observe that this operation might lead to some of elements of the matrix to become outside of 0 – 255 range and, hence, might lead to data loss.

Variables: `ImJPG_HighContrast`

18. Finally, try another operation which is not linear but is still interesting. This operation is called gamma-correction. Run the following code and observe the result:

```
%% Gamma correction
ImJPG_Gamma05=uint8((double(ImJPG)).^0.95);
```

```
figure;  
imshow(ImJPG_Gamma05);  
ImJPG_Gamma15=uint8((double(ImJPG)).^1.05);  
figure;  
imshow(ImJPG_Gamma15);
```

Variables: ImJPG\_Gamma05, ImJPG\_Gamma15

## Project 3: Matrix multiplication, inversion, and photo filters

**Goals:** We will use matrix multiplication in order to manipulate image color and achieve different effects.

**To get started:**

- Create a new Matlab script and save it as `lab03.m`.
- Download the file `daylilies.jpg`<sup>2</sup> and save it in your working directory. You can use an alternative .jpg file of your choice. Your outputs will be different from shown here in that case.

**Matlab commands used:** `imread`, `imshow`, `size`, `figure`, `reshape`, `uint8`, `double`, `for... end`

**What you have to submit:** The file `lab03.m` which you will create during the lab session.

### INTRODUCTION

Have you ever wondered how photo filters work? In this project we will learn how to manipulate images in order to obtain different color effects. As we learned in the previous project, an image can be represented by a matrix with the dimensions corresponding to the dimensions of the image. Each of the elements of this matrix contains the number (or numbers) representing the color of the corresponding pixel. If the image is a color image, then the color of the pixel is represented by three numbers  $\{R, G, B\}$  (Red, Green and Blue) with each number ranging from 0 to 255 (RGB system). We will look at several different ways to manipulate the image colors to obtain effects similar to those of Instagram and other graphics software.

### TASKS

1. Start by loading the image into Matlab and displaying it on the screen by using the familiar commands `imshow` and `imread`. Save the resulting array as `ImJPG`. You should see something similar to the image on the Fig. 4.

Variables: `ImJPG`

2. Check the dimensions of the array `ImJPG` using the command

```
[m,n,1]=size(ImJPG)
```

Observe that Matlab will return three numbers. Since the image is colored, the resulting array is three-dimensional. Among other things, Matlab allows operations on multidimensional arrays. The first two dimensions of the array `ImJPG` correspond to the horizontal and vertical dimensions (number of pixels) in the image, and the third dimension stores three numbers corresponding to the values of Red, Green, and Blue for each pixel. These three colors mixed together produce all other colors in the image. The values for Red, Green and Blue can range from 0 to 255, allowing us to create  $256^3 = 16,777,216$  colors. In this project, we will be manipulating these three numbers to achieve various visual effects (similar to the way photo filters work).

Variables: `m,n,1`

---

<sup>2</sup>The copyright to this photo belongs to the author.





Figure 4: The original image showing orange and yellow daylilies

3. First of all, let us look into amount of Red, Green, and Blue color in the image. To do this, let us extract individual layers or color channels from the image array. To obtain the red channel of color in the image, use the following command:

```
redChannel = ImJPG(:, :, 1);
```

Similarly, extract green and blue color channels and save them in the arrays **greenChannel** and **blueChannel**. Display each array in a separate figure using **figure** and **imshow** commands. Observe that individual color channels will be shown by Matlab as grayscale images. That is due to the fact that we have taken “layers” off of the matrix **ImJPG** and each of these layers individually looks like a two-dimensional array with numbers ranging from 0 to 255 which is exactly the form in which grayscale images are represented in Matlab. If you did everything correctly, you should see something similar to the Fig. 5. The whiter the pixel appears on a particular color channel, the larger amount of that color is contained in that pixel. On the other hand, the darker the area is, the less corresponding color is in that part of the image. The file **daylilies.jpg** appears to contain mostly green and red (surprise, surprise!).

Variables: **redChannel**, **greenChannel**, **blueChannel**



Figure 5: Different color channels

4. Let us convert the image into a grayscale image by using the following filter matrix:

$$GrayMatrix = \begin{bmatrix} 1/3 & 1/3 & 1/3 \\ 1/3 & 1/3 & 1/3 \\ 1/3 & 1/3 & 1/3 \end{bmatrix}$$

Multiplication by this matrix allows to compute an average of red, green, and blue in each pixel. Use the following code:

```

GrayMatrix=[1/3 1/3 1/3; 1/3 1/3 1/3; 1/3 1/3 1/3];
for i=1:m
    for j=1:n
        PixelColor=reshape(double(ImJPG(i,j,:)),3,1);
        ImJPG_Gray(i,j,:)=uint8(GrayMatrix*PixelColor);
    end;
end;
figure;
imshow(ImJPG_Gray)

```

This code produces an array `ImJPG_Gray` in which the individual values of Red, Green, and Blue have been replaced by the averages of these three colors. Observe the use of the commands `reshape`, `uint8`, `double`. For every pixel of the image, first, the color attributes (RGB) are extracted from the matrix `ImJPG`, then these color attributes are treated as a vector with three components  $[R, G, B]$ , and finally, the new color attributes are obtained by multiplying the vector  $[R, G, B]$  by the filter matrix *GrayMatrix* on the left. The result is saved as color attributes of the corresponding pixel in the array `ImJPG_Gray`. All the pixels of the array `ImJPG_Gray` have equal number of Red, Green, and Blue; this produces different shades of gray color. Observe that there are different ways to create a grayscale conversion, and this is just one of them.

Variables: `ImJPG_Gray`

Q1: Explain the purpose of `uint8` and `double` commands in the code above.

5. Modify the code above to produce a sepia conversion of the image. Instead of `GrayMatrix` use the following filter matrix, and reproduce the code above with this matrix:

$$SepiaMatrix = \begin{bmatrix} 0.393 & 0.769 & 0.189 \\ 0.349 & 0.686 & 0.168 \\ 0.272 & 0.534 & 0.131 \end{bmatrix}$$

Save the result in the `ImJPG_Sepia` array and display this array using `imshow` command.

Variables: `ImJPG_Sepia`

6. Next, consider the filter matrix

$$RedMatrix = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix}$$

Modify the code again, using the matrix above as a filter matrix. Save the result as a new array `ImJPG_Red`. Display the image.

Variables: `ImJPG_Red`

Q2: What does the transformation above do?

7. Let us permute the colors in the image. To do this repeat the steps above with the matrix:

$$PermuteMatrix = \begin{bmatrix} 0 & 0 & 1 \\ 0 & 1 & 0 \\ 1 & 0 & 0 \end{bmatrix}$$

Save the result as a new array `ImJPG_Permute`. If you did everything correctly, you should see the image shown on the Fig. 6. Observe that the flowers turned purplish blue. The

matrix above is an example of a hue rotation filter. You can produce other hue rotation effects with this more general transformation:

$$\text{HueRotateMatrix} = \begin{bmatrix} 0.213 & 0.715 & 0.072 \\ 0.213 & 0.715 & 0.072 \\ 0.213 & 0.715 & 0.072 \end{bmatrix} + \cos(\theta) \begin{bmatrix} 0.787 & -0.715 & -0.072 \\ -0.213 & 0.285 & -0.072 \\ -0.213 & -0.715 & 0.928 \end{bmatrix} + \sin(\theta) \begin{bmatrix} 0.213 & -0.715 & 0.928 \\ 0.143 & 0.140 & -0.283 \\ -0.787 & 0.715 & 0.072 \end{bmatrix}.$$

Here  $\theta$  is the angle of rotation. You can try experimenting with various values of the angle  $\theta$  to get different color effects.

Variables: **ImJPG\_Permute**



Figure 6: Colors permuted

8. Now let us delete one of the colors in the image, say, green. First, produce the filter matrix which deletes Green in the image and keeps the values of Red and Blue intact. Use it to filter the image. Save the result as **ImJPG\_DeleteGreen**.

Variables: **ImJPG\_DeleteGreen**

9. It is possible to invert the colors of the image by using the following code:

```
ImJPG_Invert=255-ImJPG;
figure;
imshow(ImJPG_Invert);
```

Observe that here again Matlab automatically substitutes a matrix of appropriate dimension (in this case  $m \times n \times l$ ) with all the elements equal to 255 instead of the constant 255.

Variables: **ImJPG\_invert**

10. It is also possible to enhance/mute individual colors in the image. For instance, consider the color transformation with the matrix:

$$\text{SaturateMatrix} = \begin{bmatrix} 1.2 & 0 & 0 \\ 0 & 0.75 & 0 \\ 0 & 0 & 2 \end{bmatrix}$$

Save the result as **ImJPG\_Saturate**.

Variables: **ImJPG\_saturate**

Q3: Explain what the transformation above does to the image.

11. Consider the color adjusting filter generated by the matrix:

$$UserMatrix = \begin{bmatrix} 0.7 & 0.15 & 0.15 \\ 0.15 & 0.7 & 0.15 \\ 0.15 & 0.15 & 0.7 \end{bmatrix}$$

Apply this filter to the array `ImJPG` and save the result as `ImJPG_User`.

Variables: `ImJPG_User`

Q4: Explain what the transformation above does to the image.

12. Consider the matrix in the previous step. Is the transformation defined by this matrix invertible? Which means, can you get the original image colors back once you applied the filter? Or is some of the information lost? Use the `inv` command to create the matrix for the inverse filter. Apply this matrix to the arrays `ImJPG_User` and `ImJPG`. Display the resulting images.

Q5: What does the inverse transformation do with the colors?

13. Repeat the steps above to see if the sepia transformation is invertible.

Q6: What do you see? Why do you think this happened?

14. Finally, try the non-linear color transformation such as Gamma-conversion:

```
ImJPG_Gamma1=uint8(double(ImJPG).^0.9)+30;  
ImJPG_Gamma2=uint8(double(ImJPG).^1.1)-50;
```

Display the resulting images given by the matrices `ImJPG_Gamma1`, `ImJPG_Gamma2`.

Variables: `ImJPG_Gamma1`, `ImJPG_Gamma2`

## Project 4: Solving linear systems in Matlab

**Goals:** To explore the built-in functions in Matlab used to solve systems of linear equations  $\mathbf{Ax} = \mathbf{b}$ . To compare different options in terms of accuracy and efficiency.

**To get started:** Create a new script file and save it as `lab04.m`.

**Matlab commands used:** `\`, `inv`, `rref`, `pinv`, `null`, `tic...` `toc`

**What you have to submit:** The file `lab04.m`, which you will create during the lab session.

### INTRODUCTION

All the methods of solution of systems of linear equations can be divided into two large groups: direct and iterative methods. The direct methods aim to obtain an “exact” (up to numerical round off error) solution of the system in a finite number of steps. The iterative methods aim to create a convergent sequence of approximations to the exact solution of a linear system. Iterative methods are usually used when the matrix of a system has a known specific structure. Matlab has many built-in functions designed to solve systems of linear equations by both direct and iterative methods. In this project we will discuss only direct methods such as Gaussian elimination.

Gaussian elimination in Matlab is based on  $LU$ - or  $PLU$ -decomposition of the matrix of the system  $\mathbf{A}$ . We can reduce the matrix  $\mathbf{A}$  to an upper triangular form using forward elimination and then solve the resulting triangular system by backward substitution. In the process of doing this, we factor the matrix  $\mathbf{A}$  into a product of two matrices: a lower triangular matrix  $\mathbf{L}$  and an upper triangular matrix  $\mathbf{U}$ . Observe that this process cannot be always carried out without interchanges of rows, and, hence, we need an additional permutation matrix  $\mathbf{P}$  in our matrix decomposition. Observe, that the matrix  $\mathbf{P}$  is necessary even in the cases when  $LU$ -decomposition can be carried out due to the matters of numerical stability (this is called pivoting). Gaussian elimination is the preferred method of solving the system numerically if you do not have any additional information about the structure of the matrix  $\mathbf{A}$ .

If the matrix  $\mathbf{A}$  is square, we can define the inverse matrix of  $\mathbf{A}$  (denoted as  $\mathbf{A}^{-1}$ ). The matrix  $\mathbf{A}^{-1}$  is also a square matrix of the same size as  $\mathbf{A}$ . The inverse matrix exists as long as  $\mathbf{A}$  is not singular, and satisfies the following equality:

$$\mathbf{A} \cdot \mathbf{A}^{-1} = \mathbf{A}^{-1} \cdot \mathbf{A} = \mathbf{I},$$

where  $\mathbf{I}$  is the identity matrix of the same size as  $\mathbf{A}$  and  $\mathbf{A}^{-1}$ . If the matrix  $\mathbf{A}$  is invertible, then we can multiply both sides of the system  $\mathbf{Ax} = \mathbf{b}$  by the inverse matrix  $\mathbf{A}^{-1}$  on the left, and obtain the following formula for the solution of the system:  $\mathbf{x} = \mathbf{A}^{-1}\mathbf{b}$ . However, as it often happens in numerical linear algebra, this theoretical formula is inconvenient for numerical computation of the solution. The reasons for this are two-fold. Firstly, the computation of inverse matrix is significantly more costly than simply solving the system. Secondly, the accuracy of the resulting solution will be worse than if  $LU$ -decomposition is used.

Finally, yet another possibility to solve a linear system is to use the reduced row echelon form of the augmented matrix of the system. Theoretically, this is equivalent to using Gauss-Jordan elimination. This method of solution will be explored in this project as well. Analysis of the corresponding algorithm shows that the resulting solution is likely to have a larger error compared to Gaussian elimination and is likely to take more time to compute.

## TASKS

1. Create a new cell and type in the following commands to generate the matrix **A** and the column vector **b**:

```
%% Matlab backslash solver
clear;
A=magic(5);
b=[10; 26; 42; 59; 38]
```

Variables: **A, b**

2. If nothing is known about the structure of the matrix of the system, the best approach to solution is to use the Matlab backslash “\” solver:

```
x=A\b
```

In most cases, this method will produce the numerical solution of the system  $\mathbf{Ax}=\mathbf{b}$  with the highest possible accuracy and in the shortest computational time.

Variables: **x**

3. Whenever we are talking about the accuracy of the solution, ideally we want to find the difference between the approximate numerical solution  $x_{num}$  and the exact solution  $x_{ex}$  of the system. However, this is not feasible in most situations since the exact solution  $x_{ex}$  is generally unknown (otherwise, there is no point in using numerical methods at all). The next best thing is to find the residual  $\mathbf{r}=\mathbf{A}\mathbf{x}-\mathbf{b}$ . If the residual is small then we can assume, with certain limitations, that our approximate solution is close enough to the exact solution (this is not always true, however). Let us check how accurate the solution is which we found in the previous step. Type the following line into your Matlab script:

```
r=A*x-b
```

The resulting vector **r** should be very close to zero.

Variables: **r**

4. The Matlab backslash “\” operator is based on *LU*- or *PLU* decomposition of the matrix **A**. The main idea here is that any square non-singular matrix **A** (meaning that the inverse matrix  $\mathbf{A}^{-1}$  exists) can be written as  $\mathbf{A} = \mathbf{PLU}$  (actually, in Matlab it is  $\mathbf{P}^T\mathbf{LU}$ ), where **P** is a permutation matrix (or matrix obtained by permuting the rows of the identity matrix of the same size as **A**); **L** is a lower-triangular matrix (all the elements above the main diagonal are equal to zero), and **U** is an upper triangular matrix (all the elements below the main diagonal are equal zero). Matlab has a special `lu` function to generate a  $\mathbf{P}^T\mathbf{LU}$  decomposition of the matrix **A**. Type the following lines into your Matlab script:

```
%% LU-factorization
[L,U,P]=lu(A);
```

This will create a new code cell and compute the matrices **P**, **L**, and **U**. Now let us use these matrices to solve the linear system  $\mathbf{Ax} = \mathbf{b}$ :

```
x1=U\ (L\ (P*b))
r1=A*x1-b
err1=x-x1
```

Observe that the difference between the solutions `err1=x-x1` is exactly the zero vector<sup>3</sup>. This is due to the fact that we found the solution using exactly the same steps as Matlab. Observe also that at no point did we use the Matlab inverse matrix function `inv`. In general, there is almost never a good reason to compute an inverse matrix. Whenever you see an inverse matrix in a numerical linear algebra textbook it is usually shorthand for “solve a system of linear equations here”.

Variables: `L, U, P, x1, r1, err1`

5. Matlab also has a separate slash operator `/` to solve linear systems of the type  $\mathbf{yA} = \mathbf{b}$ . Try running the following command

```
y=b'/A
```

Variables: `y`

6. Next, let us use the inverse matrix to solve the same system of linear equations:

```
%% Inverse matrix
x2=inv(A)*b
r2=A*x2-b
err2=x-x2
```

Here `inv(A)` produces the inverse  $\mathbf{A}^{-1}$  of the matrix  $\mathbf{A}$ . This code will also produce the solution of the system  $\mathbf{Ax} = \mathbf{b}$ , compute the residual, and find the difference between the current solution and the solution obtained by the backslash operator. Observe that the vector of difference `err2` is not exactly zero unlike in the case when we used *LU*-decomposition. Observe also that the residual `r2` is likely to be larger than the one obtained with the help of the backslash operator.

Variables: `x2, err2, r2`

7. Finally, another way of solving the system  $\mathbf{Ax} = \mathbf{b}$  is based on the reduced row echelon form. The reduced row echelon form in Matlab can be produced by using the command `rref`. Type in the following code:

```
%% Reduced row echelon form
C=rref([A b]);
x3=C(:,6)
err3=x-x3
r3=A*x3-b
```

Observe that the residual `r3` is much larger than the residual `r`. Generally speaking, `rref` is also a much less efficient method of solving linear systems compared to the backslash operator and its use for this purpose is not advised.

Variables: `x3, err3, r3`

8. To compare the computational efficiency of all three methods, run the code cell:

```
clear;
%% Computational time
Num=500;
A=rand(Num,Num)+Num*eye(Num);
```

---

<sup>3</sup>This has been observed in both Matlab 2015a and 2019a. It has come to author’s attention that this might not be true in other versions of Matlab.

```

b=rand(Num,1);
tic;
x1=A\b;
toc;
tic;
x2=inv(A)*b;
toc;
tic;
C=rref([A b]);
x3=C(:,101);
toc;

```

This code compares all three methods in terms of the computational efficiency in the example of a large matrix  $\mathbf{A}$  with the dimensions  $\text{Num} \times \text{Num}$ . The parameter  $\text{Num}$  in this code is set to be 500. The matrix  $\mathbf{A}$  is chosen in such a way that it is diagonally row-dominant. Then by the theorem of linear algebra it is guaranteed that the matrix  $\mathbf{A}$  is non-singular. The Matlab function `tic ... toc` returns the computational time necessary to perform the Matlab operations between `tic` and `toc`. You can try to experiment with the parameter  $\text{Num}$  to see how the computational time changes with the size of the matrix  $\mathbf{A}$ . Observe that solving the system with the `rref` function may take a very noticeable amount of time (you probably don't want to set  $\text{Num}$  to be more than 1000).

Variables: `Num, A, b, C, x1, x2, x3`

- Finally, let us see how to use Matlab to solve rectangular systems of linear equations. Let  $\mathbf{A}$  be an  $m \times n$  matrix of the system and let  $\mathbf{b}$  be an  $m \times 1$  column vector. Let us start with the situation when  $m > n$ . In this case, the system is overdetermined. However, it is still possible to use the Matlab backslash operator. Consider the following code:

```

%% Overdetermined system
clear;
A=[1 2 3; 4 5 6; 7 8 10; 9 11 12]
b=[1; 2; 3; 4]
x=A\b
r=A*x-b

```

The system  $\mathbf{Ax} = \mathbf{b}$  is inconsistent. The resulting “solution”  $\mathbf{x}$  is not the solution of the original linear system  $\mathbf{Ax} = \mathbf{b}$  itself (observe that the residual  $\mathbf{r}$  is not close to zero), but rather the solution to the so-called least squares problem. Namely,  $\mathbf{x}$  minimizes the length of the residual vector  $\mathbf{r} = \mathbf{Ax} - \mathbf{b}$ . It is known that the solution to the least squares problem can also be obtained by solving the system of the so-called normal equations  $\mathbf{A}^T \mathbf{Ax} = \mathbf{A}^T \mathbf{b}$ . To verify that the vector  $\mathbf{x}$  is indeed the solution to this system, type in the following code:

```

y=(A'*A)\(A'*b)
err=x-y

```

and notice that  $\mathbf{x}-\mathbf{y}$  is indeed close to zero (it is rare to obtain an exact zero in numerical computations due to roundoff errors).

Variables: `A, b, x, r, y, err`

- Let us look at the last possible case when  $m < n$ . This is a case of an underdetermined system. If this system is consistent then it has an infinite number of solutions. Consider the following code:



```

%% Underdetermined system
clear;
A=[1 2 3; 4 5 6; 7 8 9; 10 11 12]';
b=[1; 3; 5]
x=A\b
r1=A*x-b

```

The backslash operator in this case produces only one particular solution to the system  $\mathbf{Ax} = \mathbf{b}$ . Another particular solution can be obtained by the following operations:

```

y=pinv(A)*b
r2=A*y-b

```

where the function `pinv` produces the so-called Moore-Penrose pseudoinverse (the precise definition of this matrix is outside of the scope of this project). It is known that the vector  $\mathbf{y}$  has the smallest possible length among all the solutions of the system  $\mathbf{Ax} = \mathbf{b}$ . Finally, to obtain the complete set of the solutions of the system above, consider the function `null`. This function produces the complete set of the linearly independent solutions to the homogeneous system  $\mathbf{Ax} = \mathbf{0}$ . The complete solution to the inhomogeneous system  $\mathbf{Ax} = \mathbf{b}$  then can be obtained by:

```

z=null(A)*C+x;

```

where the constant vector  $\mathbf{C} = [C_1, C_2]^T$  can be chosen arbitrarily (pick any numbers for  $C_1, C_2$  that you like!).

Variables: A, b, x, r1, y, r2, C, z

## Project 5: Systems of linear equations and college football team ranking (with an example of the Big 12)

**Goals:** To use systems of linear equations to develop a ranking for Big 12 college football teams based on 2016-2017 season data using Colley's and Massey's methods.

### To get started:

- Open a new Matlab script and save it as `lab05.m`.
- Download the files `Big12th.xls`, `Scores.mat`, and `Differentials.mat`. Observe that the file `Big12th.xls` is provided for reference only and won't be used explicitly in the lab. You can also use the scores from your favorite tournament. The scores here are based on a 10 team round-robin tournament.

**Matlab commands used:** `load`, `eye`, `diag`, `sum`, `abs`, `fprintf`, `for ... end`, `sort`

**What you have to submit:** The file `lab05.m`, which you will create during the lab session.

## INTRODUCTION

The present project deals with the ranking of competing sports teams. At first, this does not appear to be a difficult problem. For instance, the easiest way to rank sports teams' performance is to compute the percentage of games which each team has won. This method has several drawbacks which are especially noticeable for the college football championship. In particular, for the percentage ranking to be objective, it is necessary that every team plays with a majority of teams participating in the competition. However, the number of teams participating in the Bowl Championship Series (BCS) is very large and each team only plays a small number of games. This is what makes the ranking of college football significantly different from, say, the NFL, where a relatively small number of teams play a representative number of games against each other.

It is necessary to note that there is no one true answer to a ranking problem. The rank of a team will depend on the factors which we will take into account and on the weight we will give to those factors. There are several questions to ask here. For instance, do point differentials matter? Is a 60-20 win any better than a 34-33 win? Taking into account point differentials may have both pros and cons. A 34-33 win versus a strong opponent is arguably better than 60-20 versus an unranked team. At the same time, if all the teams have about the same strength, then 60-20 is clearly better. At the end of the day, whether to take into account point differentials or not is a matter of taste and preference.

In this lab we will look at the two methods of ranking currently used by BCS with an example of the Big 12 Conference data from 2016-2017 season. As you may know, 10 teams play in the conference: Baylor University, Iowa State University, University of Kansas, Kansas State University, University of Oklahoma, Oklahoma State University, Texas Christian University, University of Texas Austin, Texas Tech University, and West Virginia University. We will look at the games only within the Big 12 conference and ignore all other games outside of it.

## Colley's bias free ranking

The first method we will look at has been developed by astrophysicist Wesley Colley and is named after him: Colley's bias free ranking. This method does not take into account point differentials in the games (which again can be both a good and a bad thing).

There are two main ideas behind the method. First of all, we substitute the winning percentages by the Laplace's rule of succession:

$$r_i = \frac{1 + w_i}{2 + t_i},$$

where  $w_i$  is the number of winning games for the  $i$ th team,  $t_i$  is the number of total games, and  $r_i$  is the rank of the team. Laplace's rule of succession is used for events which have only two outcomes such as either winning or losing. The rule comes from the idea in probability that unless an event always happens or never happens, we should avoid assigning this event probabilities of 1 (always true) and 0 (never true). Indeed, after playing one game, a team could either win the game or lose it. If the team won the game, then its percentage ranking is 100%; if it lost the game, its ranking is 0%. Meaning that after one game some teams are *infinitely* better than others which is not very realistic. At the same time, the Laplace's rule of succession will give us approximately 2/3 and 1/3, meaning that the team which won one game is about two times better than the one which lost one game - a good starting point for further estimates.

The second idea is that the rank of opposing teams should matter just as much as the total number of wins and losses. To incorporate the opposing teams' rankings into our system, we can write:

$$w_i = \frac{w_i - l_i}{2} + \frac{w_i + l_i}{2} = \frac{w_i - l_i}{2} + \frac{t_i}{2},$$

where  $l_i$  is the number of losses for the  $i$ th team.

Now observe that with the use of Laplace's succession rule the rankings of the teams will hover around 1/2, so we can approximate:

$$\frac{t_i}{2} = t_i \frac{1}{2} \approx \sum_{j=1}^{t_i} r_j^i.$$

Basically, we substitute each 1/2 with the ranking of the  $r_j^i$  of the  $j$ th team which  $i$ th team played against. Now, let us substitute the last formula back into the Laplace's rule to obtain the following system of linear equations (check it!):

$$(2 + t_i)r_i - \sum_{j=1}^{t_i} r_j^i = 1 + \frac{w_i - l_i}{2}.$$

This system has  $N$  equations with  $N$  unknowns  $r_i$ , where  $N$  is the number of teams playing in the championship (in our case  $N = 10$ ). Observe that the matrix of this system has special properties - it is diagonally row dominant, meaning that the system will always have a unique solution.

## Massey's method

Massey's method was developed by Ken Massey in his undergraduate research project and is now used by the BCS (Oh, the places you can go!... if you know math). Unlike Colley's method,

Massey's takes into account point differentials as well as number of wins/losses. Again, denote as  $r_i$  the rankings of the competing teams. The data we obtain from the games can be written in the form of linear equations:

$$r_i - r_j = d_{ij},$$

meaning that  $i$ th team won (lost to)  $j$ th team by the  $d_{ij}$  point differential.

The data from the games produces a large system of linear equations of the form:

$$\mathbf{P}\mathbf{R} = \mathbf{B},$$

where  $\mathbf{P}$  is the matrix of the system such that each row has only two non-zero elements, 1 and  $-1$ ,  $\mathbf{R}$  is the column-vector of the rankings  $r_i$ , and the vector of the right-hand-side  $\mathbf{B}$  contains the corresponding point differentials.

Observe that, in general, we are likely to have a lot more equations than unknowns (more games than teams), so, in general, this system of equations is likely to be inconsistent and not have any solutions. To see this, assume that the first team beat the second team by 20 points and the third team beat the second team by 5 points. This data implies that the first team should beat the third team by 15 points. Of course, the point differential of the actual game (if it happens!) between the first team and the third team might be different.

So, instead, we will give up on trying to solve this system exactly, and we will look for the vector  $\mathbf{R}$  which satisfies the system as close as possible. Namely, we will employ the least squares method, meaning, that we want to find the vector  $\mathbf{R}$  which minimizes the vector norm:

$$\|\mathbf{P}\mathbf{R} - \mathbf{B}\|_2$$

where  $\|\cdot\|_2$  is a usual Euclidean norm: if  $\mathbf{x} = (x_1, x_2, \dots, x_n)$  is a vector with  $n$  components, then

$$\|\mathbf{x}\|_2 = \sqrt{x_1^2 + x_2^2 + \dots + x_n^2}.$$

It is well-known from linear algebra that this problem is equivalent to the so-called system of normal equations written as:

$$\mathbf{P}^T\mathbf{P}\mathbf{R} = \mathbf{P}^T\mathbf{B},$$

where  $\mathbf{P}^T$  is a transpose of the matrix  $\mathbf{P}$ . Now,  $\mathbf{P}^T\mathbf{P}$  is an  $N \times N$  matrix, where  $N$  is again the number of the teams, so this results in a square system of linear equations.

The last obstacle which we will need to overcome is that the matrix  $\mathbf{P}^T\mathbf{P}$  has a rank of at most  $N - 1$  (why?), and hence, in general, the last system does not have a unique solution. To obtain a unique solution we will change each element in the last row of the matrix  $\mathbf{P}^T\mathbf{P}$  to 1 and we will change the last element in the vector  $\mathbf{P}^T\mathbf{B}$  to zero. Practically this means that all the rankings  $r_i$  obtained in this way will need to sum up to zero.

Now that you learned the theory behind the ranking, let us create code in Matlab which will produce the rankings of teams in the Big 12. To do so, first of all save the data files `Scores.mat` and `Differentials.mat` in the directory you will be working in. Each of these files contains the results of the games from the Big 12 championship from the season 2016-2017 in the form of the table (matrix). The `Scores.mat` contains a  $10 \times 10$  matrix where each row corresponds to the scores of the Big 12 teams in the following order:

1. Baylor
2. Iowa State
3. University of Kansas

4. Kansas State
5. University of Oklahoma
6. Oklahoma State
7. Texas Christian
8. University of Texas Austin
9. Texas Tech
10. West Virginia

The element  $(i, j)$  of the matrix **Scores** is 1 if  $i$ th team won over  $j$ th team. Correspondingly, if  $i$ th team lost to  $j$ th team, then the  $(i, j)$  element is  $-1$ . The diagonal elements are taken to be 0.

The file **Differentials.mat** contains the matrix **Differentials** which is similar to **Scores**, only instead of  $\pm 1$  it has the point differentials of the corresponding games.

## TASKS

1. Start by loading both matrices into your Matlab script using the **load** command:

```
load('Scores.mat');
load('Differentials.mat');
```

**Variables: Scores, Differentials**

2. First, consider the Colley method. Create the matrix and the right-hand side of Colley's system of linear equations:

```
%% Colley's method
games=abs(Scores);
total=sum(games,2);
ColleyMatrix=2.*eye(10)+diag(total)-games;
RightSide=(1+0.5.*sum(Scores,2));
```

**Variables: ColleyMatrix, RightSide**

**Q1: What do the commands `eye` and `diag` do?**

3. Solve the linear system using the backslash operator:

```
RanksColley=ColleyMatrix\RightSide;
```

**Variables: RanksColley**

4. Display the results on the screen:

```
Teams={'Baylor', 'Iowa State', 'University of Kansas', 'Kansas State',...
      'University of Oklahoma', 'Oklahoma State', 'Texas Christian',...
      'University of Texas Austin', 'Texas Tech', 'West Virginia'};
[RanksDescend,Order]=sort(RanksColley,'descend');
fprintf('\n');
for j=1:10
    fprintf('%8.3f %-15s\n', RanksColley(Order(j)), Teams{Order(j)})
end;
```

Variables: Teams, Order, RanksDescend

Q2: Observe that the array **Teams** is in curly parentheses. What kind of array is that?

Q3: Explain what kind of formatting does the line `'%8.3f %-15s\n'` in the command `fprintf` produce?

Q4: What does `sort` function do?

5. Now let us use Massey's method to rank the teams. Create the matrix **P** and the right-hand-side **B** by using the following code:

```
%% Massey's method
l=0;
for j=1:9
    for k=j+1:10
        if Differentials(j,k)~=0
            l=l+1;
            P(l,j)=1;
            P(l,k)=-1;
            B(l)=Differentials(j,k);
        end;
    end;
end;
```

This code scans the part of the matrix **Differentials** which is located above the main diagonal. For each nonzero  $(i, j)$  (meaning that the teams  $i$  and  $j$  played a game) element of this matrix it creates a row of the matrix **P** which contains only two non-zero elements: 1 in the  $i$ th column of this row and  $-1$  in the  $j$ th column. The vector **B** of the right-hand-side is created simultaneously and has the corresponding point differential in the game between  $i$ th and  $j$ th team.

Variables: P, B

6. Next, create the normal system of linear equations:

```
A=P'*P;
D=P'*B';
```

Variables: A, D

7. Substitute the last row of the matrix as described above:

```
A(10,:)= [1 1 1 1 1 1 1 1 1 1];
D(10)=0;
```

8. Finally, solve the system:

```
RanksMassey=A\D;
```

Variables: RanksMassey

9. Display the results on the screen as before:

```
[RanksDescend,Order]=sort(RanksMassey,'descend');
fprintf('\n');
for j=1:10
    fprintf('%8.3f %-15s\n', RanksMassey(Order(j)), Teams{Order(j)})
end;
```

Variables: `RanksDescend, Order`

10. Compare the results of both Colley's and Massey's methods. Compare both the methods with the official results. Observe that while these two methods are used by BCS, they do not take into account many things, such as the timing of the games, strength of schedule, games played against opponents which are not in the Big 12, and so on. These factors may affect the rankings and make it deviate from the official. For the data considered in this project the comparison with the official results should be very good.
11. Create a new code cell. Switch the result of the game between the top two teams in Colley's ranking and run the Colley's algorithm again. Produce the rankings of the teams in this case.

Q5: Did the ranking of the top teams change?

12. Create a new code cell. Now, switch the result of the game between two top teams in Massey's ranking (those might be different teams from Colley's algorithm!) and run the Massey's algorithm again. Produce the rankings of the teams in this case.

Q6: Did the ranking of the top two teams change?

## Project 6: Convolution, inner product, and image processing revisited

**Goal:** In this project we will look at the convolution of two matrices and its application to image processing.

**To get started:**

- Create a new Matlab script file and save it as `lab06.m`
- Download the file `einstein.jpg` and put it in your working directory.

**Matlab commands used:** `load`, `rand`, `double`, `uint8`, `conv2`, `filter2`

**What you have to submit:** The file `lab06.m` which you will create during the lab session.

### INTRODUCTION

In this section we will look at the applications of matrix multiplication to the filtering of images. The most common operations in filtering are blurring/smoothing, sharpening, and edge detection. All of these are examples of so-called convolution filters. These filters work by changing the color of the pixels in the image using the colors of nearby pixels.

Let us look at how filters of this type work. Consider an image described by the matrix  $\mathbf{f} = (f_{ij})$  where  $(i, j)$  are integer coordinates of the pixel in an image, and the value  $f_{ij}$  returns the color of the image. Consider  $\mathbf{h} = (h_{kl})$  to be a convolution matrix (filter matrix). These matrices might have (and usually do have) different dimensions. The filter matrix is usually relatively small. Then a new matrix which is a convolution of  $\mathbf{f}$  and  $\mathbf{h}$  (written as  $\mathbf{g} = \mathbf{f} \star \mathbf{h}$ ) can be found by the following formula:

$$g_{ij} = \sum_k \sum_l f_{kl} h_{i-k+1, j-l+1},$$

where the summation runs over all the indices which make sense. In particular, if an element with the certain index does not exist, we simply assume the corresponding term to be zero. You can think of convolution being obtained by the filter matrix  $\mathbf{h}$  sliding over the original image  $\mathbf{f}$ .

Notice that the formula for convolution resembles the definition of an inner product if we treat both matrices as vectors (say, with columnwise orientation) and flip the second matrix both horizontally and vertically.

Matlab has two functions to compute convolutions of matrices: `conv2` and `filter2`. These two functions are closely related. In particular, the function `filter2` is equivalent to the function `conv2` with the matrix  $\mathbf{h}$  rotated 180 degrees.

### TASKS

1. To begin, let us load the file `einstein.jpg` and save the resulting matrix as `ImJPG`. Use the `size` function to find the dimensions  $m, n$  of the matrix `ImJPG`.

Variables: `ImJPG`, `m`, `n`

2. Introduce some noise into the image by adding random fluctuations of color to each point:

```
ImJPG_Noisy=double(ImJPG)+50*(rand(m,n)-0.5);
```



Observe that the command `rand(m,n)` produces a matrix of the dimension  $m \times n$  filled with pseudorandom numbers within the interval  $(0, 1)$ . The amplitude of the noise is equal to  $\pm 25$  shades of gray. Also, notice the function `double` which converts the variables from the type `uint8` to the type `double`.

- Two of the most common operations on images done with convolution filters include smoothing and sharpening. Let us start by using two average smoothing filters given by the matrices:

$$Kernel\_Average1 = \frac{1}{5} \begin{bmatrix} 0 & 1 & 0 \\ 1 & 1 & 1 \\ 0 & 1 & 0 \end{bmatrix}, \quad Kernel\_Average2 = \frac{1}{9} \begin{bmatrix} 1 & 1 & 1 \\ 1 & 1 & 1 \\ 1 & 1 & 1 \end{bmatrix}$$

- Using the appropriate Matlab syntax, type in the matrices `Kernel_Average1` and `Kernel_Average2`.

Variables: `Kernel_Average1`, `Kernel_Average2`

Q1: Why are the fractions  $1/5$  and  $1/9$  necessary?

- Apply the filters to the noisy image by using the commands

```
ImJPG_Average1=conv2(Kernel_Average1,ImJPG_Noisy);
ImJPG_Average2=conv2(Kernel_Average2,ImJPG_Noisy);
```

Display the resulting images in separate figure windows and observe the result. Don't forget to convert the results back to the integer format by using `uint8` function.

Variables: `ImJPG_Average1`, `ImJPG_Average2`

Q1: What is the size of the matrices `ImJPG_Average1`, `ImJPG_Average2`?

Q2: What is the size of the original matrix `ImJPG`? Use the `size` function.

Q3: Which filter blurs more? Why?

- An alternative blurring filter, Gaussian blur, is given by the matrix:

$$Kernel\_Gauss = \frac{1}{8} \begin{bmatrix} 0 & 1 & 0 \\ 1 & 4 & 1 \\ 0 & 1 & 0 \end{bmatrix}$$

which assigns a higher weight to the pixel color in the center. Type in the matrix `Kernel_Gauss` in your Matlab file using the appropriate syntax.

Variables: `Kernel_Gauss`

- Perform the convolution using the function `conv2` and `Kernel_Gauss` and save the resulting array as `ImJPG_Gauss`. Display the result in a new window.

Variables: `ImJPG_Gauss`

- Observe that we can “layer” filter effects. Perform another convolution with the Gaussian kernel on the image `ImJPG_Gauss`, save the result as `ImJPG_Gauss2`, and display the result in a new window.

Variables: `ImJPG_Gauss2`

Q4: Devise a matrix for a filter which is equivalent to applying Gaussian convolution twice.

Q5: What is the size of the matrix for this filter?

9. It is also possible to blur the image over a larger area:

$$Kernel\_Large = \frac{1}{80} \begin{bmatrix} 0 & 1 & 2 & 1 & 0 \\ 1 & 4 & 8 & 4 & 1 \\ 2 & 8 & 16 & 8 & 2 \\ 1 & 4 & 8 & 4 & 1 \\ 0 & 1 & 2 & 1 & 0 \end{bmatrix}$$

Apply the kernel `Kernel_Large` to the matrix `ImJPG`, save the result as `ImJPG_Large`, and display the figure in a new figure window.

Variables: `ImJPG_Large`

Q6: Compare with applying Gaussian blur applied twice. Which one blurs more?

10. An opposite action to blurring is the sharpening of an image with a convolution filter. Type the following kernels into your Matlab file:

$$Kernel\_Sharp1 = \begin{bmatrix} 0 & -1 & 0 \\ -1 & 5 & -1 \\ 0 & -1 & 0 \end{bmatrix}, \quad Kernel\_Sharp2 = \begin{bmatrix} -1 & -1 & -1 \\ -1 & 9 & -1 \\ -1 & -1 & -1 \end{bmatrix}.$$

Variables: `Kernel_Sharp1`, `Kernel_Sharp2`

11. Perform the convolution of the original image `ImJPG` with the kernels `Kernel_Sharp1`, `Kernel_Sharp2` using the function `conv2` and save the resulting arrays as `ImJPG_Sharp1` and `ImJPG_Sharp2`. Display the results in new figure windows.

Variables: `ImJPG_Sharp1`, `ImJPG_Sharp2`

12. Finally, it is possible to use convolution to detect edges in the image. Edge detection is used for image segmentation and data extraction in areas such as image processing, computer vision, and machine vision. Two of the most common filters used for these are the Sobel horizontal and vertical filters:

$$Kernel\_Sobel1 = \begin{bmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{bmatrix}, \quad Kernel\_Sobel2 = \begin{bmatrix} -1 & -2 & -1 \\ 0 & 0 & 0 \\ 1 & 2 & 1 \end{bmatrix}.$$

Sobel filters can be interpreted as discrete derivatives in the horizontal and vertical directions. Type in the matrices `Kernel_Sobel1`, `Kernel_Sobel2`.

Variables: `Kernel_Sobel1`, `Kernel_Sobel2`

13. Perform the convolution of the original image `ImJPG` with the Sobel kernels using the function `conv2` and save the resulting arrays as `ImJPG_Sobel1` and `ImJPG_Sobel2`. Display the results in new figure windows.

Variables: `ImJPG_Sobel1`, `ImJPG_Sobel2`

14. Create a combined image with both horizontal and vertical edges by summing up the matrices `ImJPG_Sobel1`, `ImJPG_Sobel2`. Display the result in a new figure window by using the code:

```
figure;
imshow(uint8(ImJPG_Sobel1+ImJPG_Sobel2));
```

15. Alternatively, Laplacian edge detection can be used with the following filter:

$$Kernel\_Laplace = \begin{bmatrix} 0 & -1 & 0 \\ -1 & 4 & -1 \\ 0 & -1 & 0 \end{bmatrix}.$$

Type in the matrix `Kernel_Laplace`. The laplace kernel is a discrete analogue of continuous Laplacian and may be interpreted as a sum of two discrete partial derivatives of the second order.

Variables: `Kernel_Laplace`

16. Perform the convolution of the original image `ImJPG` with the Laplace kernel using the function `conv2` and save the resulting array as `ImJPG_Laplace`. Display the results in a new window.

Variables: `ImJPG_Laplace`

## Project 7: Norms, angles, and your movie choices

**Goals:** Use a norm and an inner product to detect similarity between users' choices and to create a simple recommender system.

**To get started:**

- Open a new Matlab script and save it as `lab07.m`.
- Download the file `users_movies.mat`<sup>4</sup> which contains the arrays `movies`, `users_movies`, `users_movies_sort`, `index_small`, `trial_user` and place it in your working directory.

**Matlab commands used:** `load`, `size`, `for... end`, `if... end`, `fprintf`, `length`, `prod`, `norm`

**What you have to submit:** The file `lab07.m` which you will create during the lab session.

### INTRODUCTION

Being able to predict users' choices is big business. Many internet services are studying consumer choices and preferences to be able to offer you the products you might like (and correspondingly the products you are more likely to buy). Netflix, Pandora Internet radio, Spotify, Facebook targeted ads, Amazon, and even online dating websites are all examples of recommender systems. These services are sometimes willing to go to great lengths in order to improve their algorithms. In fact, in 2006, Netflix offered 1 million dollars to the team which would be able to improve their Cinematch algorithm predictions by just 10%.

Music Genome Project which is a trademark of Pandora Internet radio is another example of a recommender system. If you have ever listened to Pandora, you might have been amazed how uncannily spot-on their next song recommendation is (at least, the author of this project was certainly impressed by it). Every song is assigned a rating from 0 to 5 in 450 different categories ("genes"). Thus, any song is represented by a vector with 450 components. Mathematical comparison of these vectors allows the algorithm to suggest the next song to a user.

Observe that data about movies, songs, consumer products, etc, can be often arranged in a form of a vector. These vector representations can then be used in different algorithms to compare items and predict similarity. In this project, we will use linear algebra to establish similarities in tastes between different users. The main idea is that if we know the current ratings from a certain user, then by comparing them with the ratings of other users in a database, we can find users with similar tastes. Finally, we can look at the items highly rated by those users and suggest these items to the current user.

In this lab, we will look at the MovieLens database [10] which contains around 1 million ratings of 3952 movies by 6040 users. We will create a very simple recommender system by finding similarities between users' tastes. Our techniques will be based on using the Euclidean distance (norm), the standard scalar product, and finding correlation coefficients (or angles between the vectors of data).

---

<sup>4</sup>The original data is available at <https://grouplens.org/datasets/movielens/>.

## TASKS

1. Load the arrays `movies`, `users_movies`, `users_movies_sort`, `index_small`, `trial_user` from the file `users_movies.mat` using the `load` command. The matrix `users_movies` should be a  $6040 \times 3952$  matrix containing integer values between 0 and 5 with 1 meaning “strongly dislike” and 5 meaning “strongly like”. The 0 in the matrix means that the user did not rate the movie. The array `movies` contains all the titles of the movies. The matrix `users_movies_sort` contains an extract from the matrix `users_movies.mat` with rating for the 20 most popular movies selected. The indexes of these popular movies are recorded in the array `index_small`. Finally, ratings of these popular movies by yet another user (not any of the users contained in the database) are given by the vector `trial_user`. It is suggested to view all the variables and their dimensions by using the “Workspace” window of the Matlab environment:

```
%% Load the data
clear;
load('users_movies.mat','movies','users_movies','users_movies_sort',...
    'index_small','trial_user');
[m,n]=size(users_movies);
```

Variables: `movies`, `users_movies`, `users_movies_sort`, `index_small`, `trial_user`, `m`, `n`

2. Print the titles of the 20 most popular movies by using the following code:

```
fprintf('Rating is based on movies:\n')
for j=1:length(index_small)
    fprintf('%s \n',movies{index_small(j)})
end;
fprintf('\n')
```

Observe that the movie titles are called from the cell array `movies` (notice the curly parentheses) by using an intermediate array `index_small`.

3. Now let us select the users we will compare the `trial_user` to. Here, we want to select the people who rated all of the 20 movies under consideration. This means that there should not be any zeros in the corresponding rows of the matrix `users_movies_sort`. This can be accomplished by the following code:

```
%% Select the users to compare to
[m1,n1]=size(users_movies_sort);
ratings=[];
for j=1:m1
    if prod(users_movies_sort(j,:))~=0
        ratings=[ratings; users_movies_sort(j,:)];
    end;
end;
```

Observe the use of the command `prod` to compute the product of the elements of the rows of the array `users_movies_sort`. The array `ratings` contains all the users which have rated all the popular movies from the array `index_small`. In general, it may happen that this array is empty or too small to create any meaningful comparison - we won't consider these cases in this project.

Variables: `ratings`, `m1`, `n1`

Q1: What does the command `ratings=[]` do?

- Next, we can look at the similarity metric based on the Euclidean distance. The idea here is that we treat the array `trial_user` and the rows of the array `ratings` as vectors in 20-dimensional real space  $\mathbb{R}^{20}$ . Assume that all the vectors have the origin as the beginning point. We can find the distance between the end points of the vector `trial_user` and each of the vectors `ratings(j,:)`. In other words, we are looking for the user with the closest ratings to the `trial_user`. This can be accomplished by the following code:

```
%% Find the Euclidean distances
[m2,n2]=size(ratings);
for i=1:m2
    eucl(i)=norm(ratings(i,:)-trial_user);
end;
```

The vector `eucl` contains all the Euclidean distances between `trial_user` and the rows of the matrix `ratings`.

Variables: `eucl`

- Now let us select the user from the database with the smallest Euclidean distance from the `trial_user`. Instead of using the usual function `min` we will use a slightly more complicated approach. Let us sort the elements of the vector `eucl` by using the function `sort`. The advantage of this is that it allows us to find the second closest user, the third closest user, etc. There may only be a small difference between the several closest users and we might want to use their data as well.

```
[MinDist,DistIndex]=sort(eucl,'ascend');
closest_user_Dist=DistIndex(1)
```

Variables: `MinDist`, `DistIndex`, `closest_user_Dist`

- The similarity metric above is one of the simplest ones which can be used to compare two objects. However, when it comes to user ratings it has certain disadvantages. For instance, what if the users have similar tastes, but one of them consistently judges movies harsher than the other one? The metric above would rate those two users as dissimilar since the Euclidean distance between vectors of their opinions might be pretty large. To rectify this problem, we can look at a different similarity metric known in statistics as the Pearson correlation coefficient which can be defined as

$$r(\mathbf{x}, \mathbf{y}) = \frac{\sum_{i=1}^N (x_i - \bar{x})(y_i - \bar{y})}{\sqrt{\sum_{i=1}^N (x_i - \bar{x})^2} \sqrt{\sum_{i=1}^N (y_i - \bar{y})^2}},$$

where  $\mathbf{x} = (x_1, x_2, \dots, x_N)$ ,  $\mathbf{y} = (y_1, y_2, \dots, y_N)$  are two vectors with  $N$  components, and  $\bar{x} = \frac{1}{N} \sum_{i=1}^N x_i$ ,  $\bar{y} = \frac{1}{N} \sum_{i=1}^N y_i$  are the average (mean) values of the vectors  $\mathbf{x}$  and  $\mathbf{y}$ . If we look at the formula for  $r(\mathbf{x}, \mathbf{y})$ , we can see that it accounts for the average user opinion  $\bar{x}$ ,  $\bar{y}$ , and also for how harshly/enthusiastically they tend to judge their movies (defined by the magnitude of the vectors  $\mathbf{x} - \bar{\mathbf{x}}$  and  $\mathbf{y} - \bar{\mathbf{y}}$ , where  $\bar{\mathbf{x}}$  and  $\bar{\mathbf{y}}$  denote the vectors of the same size as  $\mathbf{x}$  and  $\mathbf{y}$  with all the components equal to  $\bar{x}$  and  $\bar{y}$  correspondingly). Geometrically, the correlation coefficient  $r(\mathbf{x}, \mathbf{y})$  corresponds to the cosine of the angle between the vectors  $\mathbf{x} - \bar{\mathbf{x}}$  and  $\mathbf{y} - \bar{\mathbf{y}}$  (the closer this angle is to zero, the more similar are the opinions of two people). To compute the Pearson correlation coefficient, let us centralize the columns of the matrix `ratings` and the vector `trial_user` first:

```
ratings_cent=ratings-mean(ratings,2)*ones(1,n2);
trial_user_cent=trial_user-mean(trial_user);
```

Variables: `ratings_cent`, `trial_user_cent`

7. Next, use the `for... end` loop to compute the Pearson correlation coefficients between the rows of the matrix `ratings` and the vector `trial_user`. Save the result as a vector `pearson`.

Variables: `pearson`

8. Finally, observe that the value  $r(\mathbf{x}, \mathbf{y})$  belongs to the interval  $(-1, 1)$ . The closer the coefficient is to 1, the more similar the tastes of the users are. Finally, let us sort the vector `pearson` as before using the `sort` function. Save the results of this function as `[MaxPearson, PearsonIndex]` and find the maximal correlation coefficient which will be the first element in the matrix `MaxPearson`. Save this element as `closest_user_Pearson`.

Variables: `MaxPearson`, `PearsonIndex`, `closest_user_Pearson`

9. Compare the elements of the vectors `DistIndex`, `PearsonIndex`.

Q2: Are the variables `closest_user_Pearson` and `closest_user_Dist` the same?

10. Now let us display the recommendations on the screen. Use the following list to create the list of movies which the `trial_user` has liked and the lists of recommendations for him/her based on the distance criterion and the Pearson correlation coefficient criterion:

```
%% Recommendations
recommend_dist=[];
for k=1:n
    if (users_movies(closest_user_Dist,k)==5)
        recommend_dist=[recommend_dist; k];
    end;
end;
recommend_Pearson=[];
for k=1:n
    if (users_movies(closest_user_Pearson,k)==5)
        recommend_Pearson=[recommend_Pearson; k];
    end;
end;
liked=[];
for k=1:20
    if (trial_user(k)==5)
        liked=[liked; index_small(k)];
    end;
end;
```

We use the rating equal to 5 both as the criterion for liking the movie and the criterion to recommend the movie. Of course, you can broaden this up and also include the movies ranked as 4.

Variables: `liked`, `recommend_dist`, `recommend_Pearson`

11. Finally, display the titles of the movies from the arrays `liked`, `recommend_dist`, `recommend_Pearson` on the screen by using the procedure similar to the one used in the step 2.

12. Take a look at the list of the popular movies displayed in step 2. Chances are you might have seen the majority of them before. Create your own vector of ratings and call it `myratings`. Ratings should be integers between 1 and 5. Again, assign the rating 1 if you really disliked the movie, and 5 if you really liked it. If you haven't seen a particular movie, pick its rating at random. The vector of ratings `myratings` should be a row-vector with 20 elements.

Variables: `myratings`

13. Create a new code cell. In this cell, repeat steps 4-11 of this project and substitute the vector `trial_user` by the vector `myratings`. This should produce a personal recommendation based on your own ratings.

Variables: `liked`, `recommend_dist`, `recommend_Pearson`



## Project 8: Interpolation, extrapolation, and climate change

**Goals:** To explore applications of linear algebra for the interpolation of data and prediction of future tendencies.

**To get started:**

- Create a new Matlab script and save it as `lab08.m`.
- Download the file `temperature.mat` which contains average temperatures for the past 136 years <sup>5</sup>. Temperature is given in Celsius.

**Matlab commands used:** `plot`, `axis`, `vander`, `interp1`, `orth`

**What you have to submit:** The file `lab08.m`, which you will create during the lab session.

### INTRODUCTION

In many practical applications it is necessary to obtain an intermediate value of a parameter from a few known data points. If this intermediate value lies in between two known measurements, we will call this process interpolation. If the value is outside of the sampled interval, then we will call this process extrapolation. In this project we will look at several procedures for interpolation and extrapolation and apply them to weather and climate change prediction.

The most common reason for using interpolation is not having enough of data points. These data points can be obtained from experimental observations which can be expensive or difficult to perform, or from past observations where the data is simply unavailable. Interpolation is also used when the function which produces the data is known but has a complicated structure. In this case it can be computationally prohibitive to use this function, especially if, in the course of computation, this function needs to be called multiple times. For instance, the function can be given by a slowly convergent series which requires computing many terms to obtain adequate accuracy. Approximating this complicated function by a simpler function may be helpful. Interpolation of a function can be sufficiently accurate provided that the function is smooth and there are enough data points available. Some of the usual types of interpolation include polynomial interpolation when the function is approximated by a polynomial, trigonometric approximation with a truncated Fourier series, or various piece-wise approximations. More intricate interpolation methods can be used if additional information is available about the approximated function.

Extrapolation, on the other hand, is a significantly more difficult procedure. The general rule is not to extrapolate too far beyond the range of the observed parameters. There are multiple reasons for this. For instance, some additional parameters may affect the behavior of a function outside of the observed range of values. Imagine trying to extrapolate physical laws applicable for regular macro-solids (such as things we use in our daily life) to nano-scale or to astro-scales. The obtained results will be meaningless since different laws apply at those scales. Greater uncertainty and errors while extrapolating also appear because small errors in data measurements can potentially lead to very large errors while extrapolating the data. These errors are somewhat bounded while interpolating between two known data points.

---

<sup>5</sup>The temperature data has been taken from NASA's website, <https://climate.nasa.gov/vital-signs/global-temperature/>

## TASKS

1. To begin, let us look at interpolation based on the polynomial approximation. Assume that we have  $n + 1$  data points (measurements)  $(x_i, y_i)$ ,  $i = 0, \dots, n$ . To obtain the values of the function in between these data points we can simply trace a polynomial of the  $n$ th degree through the data points and use this polynomial to estimate the values of a function in between measurements. Thus, the goal is to find a polynomial

$$p(x) = a_n x^n + a_{n-1} x^{n-1} + \dots a_1 x + a_0$$

with unknown coefficients  $a_i$ ,  $i = 0, \dots, n$ , which passes through the points  $(x_i, y_i)$ . Then the coefficients  $a_i$  must satisfy the following system of linear algebraic equations

$$a_n x_i^n + a_{n-1} x_i^{n-1} + \dots a_1 x_i + a_0 = 0, \quad i = 0, \dots, n.$$

We can write this system in the form

$$\mathbf{V}\mathbf{z} = \mathbf{y},$$

where  $V$  is the so-called Vandermonde matrix:

$$\mathbf{V} = \begin{bmatrix} 1 & x_0 & \dots & x_0^{n-1} & x_0^n \\ 1 & x_1 & \dots & x_1^{n-1} & x_1^n \\ \dots & \dots & \dots & \dots & \dots \\ 1 & x_n & \dots & x_n^{n-1} & x_n^n \end{bmatrix},$$

$\mathbf{z} = [a_0, a_1, \dots, a_n]^T$  is the unknown vector of polynomial coefficients and  $\mathbf{y} = [y_0, y_1, \dots, y_n]^T$  is the known vector of the function values at sample points.

We will start by approximating the data for the average temperatures in Kansas<sup>6</sup>:

Month	1	2	3	4	5	6	7	8	9	10	11	12
Temperature (High)	37	44	55	66	75	84	89	88	80	69	53	41
Temperature (Low)	17	23	33	43	53	63	68	65	56	44	32	22

In our computations, we will work with high temperatures only. To begin, let us generate a plot of the temperatures in Matlab.

```
clear;
WeatherHigh=[37 44 55 66 75 84 89 88 80 69 53 41]';
figure;
plot(1:12,WeatherHigh,'r-x');
axis([1,12 30 95]);
```

Observe that Matlab will automatically join the available data points by straight lines.

**Variables: WeatherHigh**

2. Assume for now, that we are only given the temperatures for four months: January, May, August, and December. Select the corresponding data from the variable **WeatherHigh** above and generate the Vandermonde matrix based on this data. Now solve the resulting system using Matlab's backslash linear system solver:

---

<sup>6</sup>Source for the temperature data: Wikipedia <https://en.wikipedia.org/wiki/Kansas#Climate>, temperature is given in Fahrenheit.

```
x=[1,5,8,12]; %Numbers for January, May, August, and December
V=vander(x);
CoefHigh=V\WeatherHigh(x);
```

The resulting vector **CoefHigh** contains coefficients of the cubic polynomial approximating the average high annual temperature in Kansas.

**Variables: V, CoefHigh**

3. Matlab's **polyval** function can be used to efficiently evaluate a polynomial at the given set of points. It takes two vectors as arguments. The first vector represents the coefficients of the polynomial and the second vector represents the points we want to evaluate the polynomial at. Remember that you can always type **help polyval** or **doc polyval** in the Matlab command window to see more options. Let us plot the graph of the polynomial obtained at the previous step by using the following code:

```
xc=1:0.1:12;
ycHigh=polyval(CoefHigh,xc);
plot(xc,ycHigh,'b-',1:12,WeatherHigh,'r-x');
axis([1,12 30 95]);
```

This graph shows the difference between the approximated values and the exact values. Observe that the command **axis([1,12 30 95])** sets the parameters for  $x$  and  $y$  axes. Namely, the  $x$  axis will range from  $x = 1$  to  $x = 12$  and the  $y$  axis will range from  $y = 30$  to  $y = 95$ .

**Variables: xc, ycHigh**

4. Repeat the steps above to produce a polynomial approximation based on the temperatures from the following six months: January, March, May, August, October, and December. You will need to use a polynomial of the fifth degree. Plot the resulting polynomial next to the exact values as in the step above. You will see that this approximation produces much closer results to the exact values.
5. Finally, let us use the temperatures from all twelve months. You will need to use a polynomial of the eleventh degree this time. Modify the code above and run it in Matlab. If you did everything correctly, you will obtain a picture which looks similar to the one in Fig. 7. You may notice that this approximation is not necessarily better than the

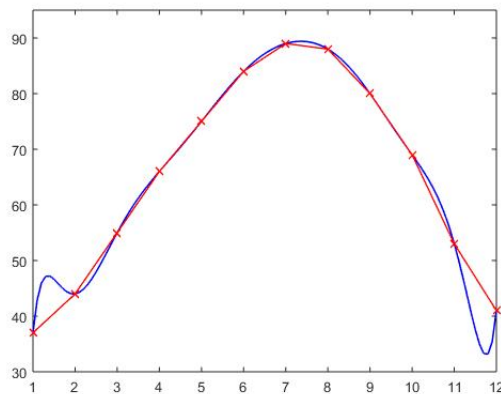


Figure 7: Approximation with the 11th degree polynomial

approximation with the 5th degree polynomial due to large fluctuations near the end points. This is typical for approximations with polynomials of a high degree on evenly spaced grids which tend to oscillate widely near the end points. This is one of the reasons why polynomials of a high degree are normally not used for interpolation on evenly spaced grids<sup>7</sup>. Additionally, if you look at the Matlab output in the command window, you will probably see something like this

```
Warning: Matrix is close to singular or badly scaled. Results may be inaccurate.
RCOND = 8.296438e-17.
```

This is due to the fact that Vandermonde matrix  $V$  is an ill-conditioned matrix, meaning that the system of linear equations with this matrix is difficult to solve with high accuracy. The matrix  $V$  becomes more ill-conditioned as the dimensions of the matrix increase. This is another reason why high degree polynomial approximations on evenly spaced grids are rarely used in interpolation.

6. If polynomials of a high degree on evenly spaced grids are not good for interpolation, what can be done instead? The most common method of interpolation is through the approximation of functions with piece-wise (typically polynomial) functions. This means that the function between data points is approximated by a small degree polynomial (polynomials of no higher than third degree are usually used). These polynomials are different on each of the intervals between the data points. Individual approximations are then glued together to achieve a desired degree of smoothness at the data points. Under smoothness here we understand the continuity of a function and its derivatives up to the some order. Smoothness of the resulting approximation depends on the degree of polynomials used and how the polynomial pieces are glued together. If a higher degree of smoothness is desired, it is necessary to use polynomials of a higher degree. For instance, if we use cubic polynomials, then it is possible to achieve continuity of the function itself and its first two derivatives at the most.

To investigate this method of interpolation, we will look at the built in Matlab function `interp1`. This function allows one to interpolate a one-dimensional function. It has several attributes which allow for different types of interpolation: “linear” which simply joins the data points with straight lines, “cubic” or “pchip” which approximates the function on each individual segment with the third degree polynomial, and “spline” which also uses third degree polynomials, but with different junction conditions. So what is the difference between “pchip” and “spline”? Simply put, the “pchip” method provides a shape-preserving piece-wise cubic polynomial function which is smooth together with its first derivative. The “spline” interpolation creates an “extra-smooth” piece-wise cubic polynomial function which is smooth together with its first and second derivatives. In general, “pchip” approximations have smaller oscillations near the ends and are easier to compute.

Compare these options by using the following code:

```
ycHigh1=interp1(x,WeatherHigh,xc,'linear');
ycHigh2=interp1(x,WeatherHigh,xc,'pchip');
ycHigh3=interp1(x,WeatherHigh,xc,'spline');
```

---

<sup>7</sup>However, approximations with the polynomials of high degree on non-evenly spaced grids may be extremely useful in computations and achieve very high precision. Chebyshev nodes and Chebyshev polynomials represent a good example of this phenomena.

Here the variables `x` and `WeatherHigh` give the vectors of initial data. The vector `xc` is a vector of points where we want to approximate the function and the vectors `ycHigh1`, `ycHigh2`, and `ycHigh3` are the values at the points `xc` resulting from the interpolation. Plot the vectors `ycHigh1`, `ycHigh2`, and `ycHigh3` versus `xc` on the same plot to observe the smoothness of the resulting data. You can see that `ycHigh2` and `ycHigh3` achieve better approximations than the polynomial approximation. The function `interp1` also has a possibility to extrapolate the results which we will explore shortly.

Variables: `ycHigh1`, `ycHigh2`, `ycHigh3`

- Now let us look at the extrapolation of the data. Extrapolation means finding an approximation to the data outside of the initial measurement interval. For our data, we will take the average global temperatures from the last 136 years starting from the year 1880<sup>8</sup>. To begin, load the matrix `temperature` from the file “temperature.mat” using the `load` command.

Variables: `temperature`

- The matrix `temperature` consists of two columns. The first column represents the years and the second the average temperatures at those years. Separate the data into two variables:

```
years=temperature(:,1);
temp=temperature(:,2);
```

Plot the graph of temperatures using the `plot` function. This will produce a graph of fluctuations of average temperatures since 1880. If you did everything correctly, you should obtain a graph similar to the one on the Fig. 8.

Variables: `years`, `temp`

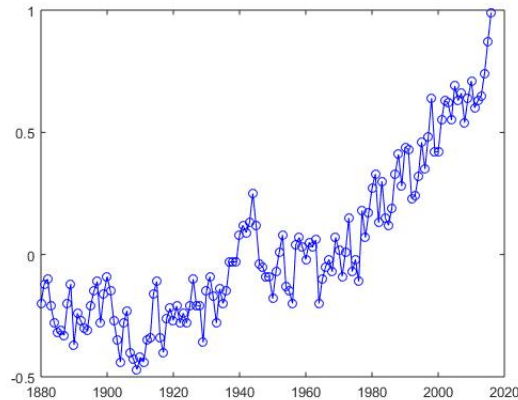


Figure 8: Average yearly temperatures.

- Let us attempt to extrapolate the data in the future. Assume that we want to know the temperature about 10 years into the future (again, it is advised not to attempt to extrapolate too far beyond the initial interval). We can do this using the following code:

```
futureyears=2016:2025;
```

---

<sup>8</sup>The source of data: NASA, <https://climate.nasa.gov/vital-signs/global-temperature/>

```

futuretemp1=interp1(years,temp,futureyears,'linear','extrap');
futuretemp2=interp1(years,temp,futureyears,'pchip','extrap');
futuretemp3=interp1(years,temp,futureyears,'spline','extrap');

figure;
plot(years,temp,'b-o',futureyears,futuretemp1,'g-o',...
futureyears,futuretemp2,'r-x',futureyears,futuretemp3,'k-d');

```

This produces three graphs based on the different interpolation/extrapolation strategies which were discussed above. If you did everything correctly, you should obtain three radically different predictions for the next 10 years!

Variables: `futureyears`, `futuretemp1`, `futuretemp2`, `futuretemp3`

10. So what happened in the previous example? The local fluctuations in temperature significantly affected the global predictions for the future. However, we are not interested in the local fluctuations but rather in global tendencies. So let us try to separate those two from each other and extrapolate based on global tendencies only. We can think of temperatures in the following terms:

$$t_i = t(\text{year}_i) + \text{noise},$$

where  $t(\text{year})$  would be the global tendency of temperature and *noise* represents relatively small year-to-year fluctuations.

We will use orthogonal projections to separate the global behavior from the “noise”. As a first approximation, let us consider how we can assess the overall average temperature. That is, we approximate the temperature  $t(\text{years})$  as a constant:

$$t(\text{years}) \approx \frac{1}{n} \sum_{i=1}^n t_i.$$

Enter the following command in the command window

```
>> sum(temp)/n
```

This allows us to see that the average temperature for the past 136 years was 0.0244C.

11. Here is another way of thinking of the same operation using orthogonal projections. If the data was truly flat, then there would be a real number  $c$  so that

$$\mathbf{t} = \begin{bmatrix} t_1 \\ t_2 \\ \vdots \\ t_n \end{bmatrix} = c \begin{bmatrix} 1 \\ 1 \\ \vdots \\ 1 \end{bmatrix} = c\mathbf{1}.$$

That is, the average temperature vector  $\mathbf{t}$  would be a scalar times the vector  $\mathbf{1}$  of all ones. Since the data is not flat, we can't find such a  $c$ . However, we can do the next best thing. We can find  $c$  to minimize the norm  $\|\mathbf{t} - c\mathbf{1}\|$ . The best choice for  $c$  is determined by the orthogonal projection on a vector  $\mathbf{1}$ :

$$c = \frac{\mathbf{1}^T \mathbf{t}}{\|\mathbf{1}\|^2} \quad \text{and for this choice of } c, \quad c\mathbf{1} = \frac{\mathbf{1}^T \mathbf{t}}{\|\mathbf{1}\|^2} \mathbf{1} = \text{proj}_{\mathbf{1}}(\mathbf{t}).$$

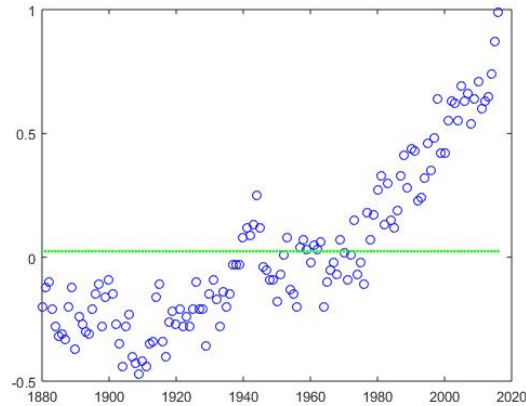


Figure 9: Approximation of data by a constant function.

Moreover, a little rearranging shows that

$$\text{proj}_{\mathbf{1}}(\mathbf{t}) = \left( \frac{1}{\|\mathbf{1}\|^2} \mathbf{1}\mathbf{1}^T \right) \mathbf{t} = \mathbf{P}\mathbf{t} \quad \text{where} \quad \mathbf{P} = \frac{1}{\|\mathbf{1}\|^2} \mathbf{1}\mathbf{1}^T \text{ is the projection matrix.}$$

Define the variable `b1` to be the `length(temp)`-dimensional `1` vector. In the space after that, define the variable `P1` to hold the projection matrix. You will either want to use Matlab's `norm` function, or remember that  $\|\mathbf{u}\| = \sqrt{\mathbf{u}^T \mathbf{u}}$ . Also, define the variable `temp1` to be the value `P1*temp`.

Variables: `P1`, `temp1`

12. Plot the resulting vector using the code:

```
plot(years,temp,'bo',years,temp1,'g. ');
hold on;
```

You should see a similar plot as before, but now with a set of green dots representing the projection (Fig. 9).

13. Now, examine the value of `temp1` in the Matlab command window.

Q1: How does the value of `temp1` relate to the average value of temperature?

14. Like vectors, matrices also have norms. The norm  $\|\mathbf{A}\|$  of a matrix `A` describes how far `A` is from the zero matrix in a certain sense. Enter the following command into Matlab

```
norm(P1*P1-P1)
```

Your answer should be something like `ans = 2.0134e-15`.

Q2: What does the result say about the relationship between the projection matrix `P` and its square `P2`?

15. While the previous approach of approximating the data with a constant function may be acceptable in some cases, it is clearly not a perfect match. Let us try instead to model the data as a linear function of `yearsi`:

$$t_i = b \text{ years}_i + c + \text{noise.}$$

If this formula was exact (with no noise), we would have that

$$\mathbf{t} = \begin{bmatrix} t_1 \\ t_2 \\ \vdots \\ t_n \end{bmatrix} = \begin{bmatrix} \text{years}_1 & 1 \\ \text{years}_2 & 1 \\ \vdots & \vdots \\ \text{years}_n & 1 \end{bmatrix} \begin{bmatrix} b \\ c \end{bmatrix} = b\mathbf{years} + c\mathbf{1},$$

so  $\mathbf{t}$  would be a vector in the subspace  $\text{span}\{\mathbf{years}, \mathbf{1}\}$ . However, as before, this is not exactly true. So, we will need to use an orthogonal projection again.

There is a nice formula for projecting onto a subspace  $\mathcal{S}$  generated by a span of several vectors  $\text{span}\{\mathbf{b}_1, \mathbf{b}_2, \dots, \mathbf{b}_k\}$ , but it requires that the vectors be *orthonormal*. That is  $\|\mathbf{b}_i\| = 1$  for each  $i$ , and  $\mathbf{b}_i^T \mathbf{b}_j = 0$  for all  $i \neq j$ .

One way to produce an orthonormal basis in Matlab is to use the Matlab `orth` function. Please, type in the code

```
m=length(years);
B2=[ones(m,1) years];
Q2=orth(B2);
```

This creates a matrix  $\mathbf{B} = [\mathbf{1} \ \mathbf{years}]$ , whose first column consists entirely of ones and whose second column is the vector  $\mathbf{years}$ , which provides the years of observation. The new array  $\mathbf{Q2}$  is defined to be the result of `orth(B2)`.

The function `orth` will create an orthonormal basis (defined by the columns of  $\mathbf{Q2}$ ) for the subspace  $\mathcal{S}$ . To check that the columns of  $\mathbf{Q2}$  form a basis, verify in the command window that the values of `rank(Q2)` and `rank([Q2,B2])` are both equal to 2.

Variables:  $\mathbf{Q2}$

Q3: What kind of matrix is the matrix  $\mathbf{Q2}^T \mathbf{Q2}$ ? Why?

16. Given an orthonormal basis defined by the columns of  $\mathbf{Q2}$  for the subspace  $\mathcal{S}$ , the formula for the projection matrix onto the subspace  $\mathcal{S}$  is simple:

$$\mathbf{P2} = \mathbf{Q2} \mathbf{Q2}^T.$$

Create a new cell and in this cell create a variable  $\mathbf{P2}$  to hold the value of the matrix of the orthogonal projection onto the subspace  $\mathcal{S}$ . Repeat the previous steps to produce the projection vector `temp2=P2*temp` and plot this vector in the previous figure window. This will produce the linear approximation to the given data. Run the following code in Matlab:

```
norm(P2*P2-P2)
```

You should get a number which is close to zero again, such as,  $5.6169\text{e-}16$ . There is a geometric explanation for this fact. The matrices  $\mathbf{P1}$ ,  $\mathbf{P2}$  are the projection matrices, and projecting a vector on a subspace produces a vector belonging to this subspace. Taking a second projection, thus, will simply leave the vector as is. The projection matrix behaves as an identity matrices on the corresponding subspace if the basis consisting of the columns of the matrix  $\mathbf{Q2}$  is selected.

Variables:  $\mathbf{P2}$ , `temp2`



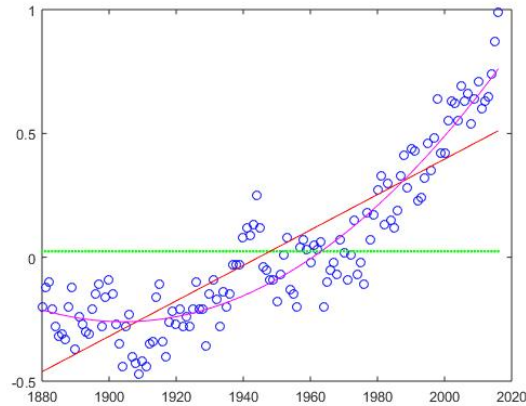


Figure 10: Approximation of data by a quadratic function.

17. Finally, let us approximate the data by using a quadratic function. Create a new cell and reproduce the code from the previous cell but with renamed variables (e.g., B3 instead of B2). Make sure to copy the plotting code and switch P2 to P3.

Aside from changing 2s to 3s, the only other change you'll need to make is to add a column to the B3 matrix. This column should be the vector of squares of the elements of the vector **years**, which can be produced in Matlab with the expression **years.^2** (notice the dot before the power sign!).

**Variables: B3, Q3, P3**

18. Plot the initial data and all the approximations above on a new plot (use the command **figure**). If correct, your code should produce a figure similar to the one on the Fig. 10. Observe that the quadratic function (shown by the magenta line) appears to follow the global tendencies of the temperature graph pretty well.
19. Now, let us make predictions for the future. Create a new cell and run the following code:

```
futureyears=2016:2116;
futuretemp3=interp1(years,temp3,futureyears,'spline','extrap');
plot(futureyears,futuretemp3,'g-');
```

This should create our prediction for an average temperature for the next 100 years. If you did everything correctly, provided the current tendencies hold, by the year 2116 the average temperature should be about 3.5C. Note that climate prediction is notoriously difficult, and it is very likely that there are factors which influence the data which are not accurately picked up by our simple model. Also, please note that in the past, temperatures tended to rise faster than any prediction. At the same time, NASA reports that the currently available models for climate change predict rise in temperatures anywhere between a 2C and 6C, so our simple model predictions are right in the middle!

## Project 9: Orthogonal matrices and 3D graphics

**Goals:** To explore applications of orthogonal matrices to rotation of 3D objects.

**To get started:** Download the files `lab09.m`<sup>9</sup>, `v.mat`, and `f.mat`. The last two files contain corresponding vertices and faces of a 3D model<sup>10</sup>. Save these files in your working directory.

**Matlab commands used:** `function`, `zeros`, `sin`, `cos`, `if... end`, `for... end`, `line`, `figure`, `hold on`, `hold off`, `bucket`, `size`, `load`

**What you have to submit:** The files `rotation.m` and `lab09.m` which you will create during the lab session.

### INTRODUCTION

Recall that the square matrix  $\mathbf{Q}$  is called orthogonal if  $\mathbf{Q}^T \mathbf{Q} = \mathbf{Q} \mathbf{Q}^T = \mathbf{I}$ , where  $\mathbf{I}$  is an identity matrix of the same size as  $\mathbf{Q}$ . In other words, this means that  $\mathbf{Q}^T = \mathbf{Q}^{-1}$ . Rows (columns) of an orthogonal matrix generate an orthonormal set of vectors. Namely, if  $\mathbf{Q} = [\mathbf{q}_1, \mathbf{q}_2, \dots, \mathbf{q}_n]$ , where  $\mathbf{q}_j$  are the columns of the matrix  $\mathbf{Q}$ , then  $\mathbf{q}_j \cdot \mathbf{q}_k = \delta_{jk}$ , and  $\delta_{jk}$  denotes the Kronecker symbol:  $\delta_{jk} = 1$ , if  $j = k$ , and  $\delta_{jk} = 0$ , if  $j \neq k$ .

Another property of orthogonal matrices is that  $\mathbf{x} \cdot \mathbf{y} = \mathbf{Q}\mathbf{x} \cdot \mathbf{Q}\mathbf{y}$  for any two vectors  $\mathbf{x}$  and  $\mathbf{y}$ , meaning that multiplication by an orthogonal matrix  $\mathbf{Q}$  does not change the scalar product of two vectors. Geometrically, this means that a linear transformation  $T(\mathbf{x}) = \mathbf{Q}\mathbf{x}$  of the space  $\mathbb{R}^n$  is equivalent to the rotation of the space together with a possible reflection along some hyperplane. Whether a reflection of the space is involved can be seen by looking at the determinant of the matrix  $\mathbf{Q}$ . If  $\det \mathbf{Q} = 1$ , then the transformation is a pure rotation. If  $\det \mathbf{Q} = -1$ , then the transformation is a rotation together with a reflection.

Orthogonal matrices can be used in 3D graphics. Let us consider a three-dimensional object which we want to show on a computer screen. The obvious difficulty is that the object is three-dimensional and the screen is only two-dimensional. Thus, we need to generate a projection to show the object on the screen. One of the easiest ways to do this is to simply drop one of the coordinates. However, that allows us to view the object only from three different angles which can make it difficult to understand the real shape of the object. Using orthogonal matrices allows us to look at the object from many different angles. To accomplish this, we need three matrices which represent rotations around each of the coordinate axes  $Ox$ ,  $Oy$ , and  $Oz$ :

$$\mathbf{R}_x = \begin{bmatrix} 1 & 0 & 0 \\ 0 & \cos \theta_x & -\sin \theta_x \\ 0 & \sin \theta_x & \cos \theta_x \end{bmatrix}, \quad \mathbf{R}_y = \begin{bmatrix} \cos \theta_y & 0 & -\sin \theta_y \\ 0 & 1 & 0 \\ \sin \theta_y & 0 & \cos \theta_y \end{bmatrix}, \quad \mathbf{R}_z = \begin{bmatrix} \cos \theta_z & -\sin \theta_z & 0 \\ \sin \theta_z & \cos \theta_z & 0 \\ 0 & 0 & 1 \end{bmatrix}.$$

Here  $\theta_x$ ,  $\theta_y$ , and  $\theta_z$  are the angles of rotation around the axes  $Ox$ ,  $Oy$ , and  $Oz$  correspondingly in the counterclockwise direction. Multiplying these matrices will result in a combined rotation around all three axes:

$$\mathbf{R} = \mathbf{R}_x \mathbf{R}_y \mathbf{R}_z.$$

In this project we will use the matrix  $\mathbf{R}$  to create a two-dimensional projection of a three-dimensional object.

---

<sup>9</sup>The printout of the file `lab09.m` can be found in appendices of this book.

<sup>10</sup>The original data for this 3D model can be found here: <https://www.thingiverse.com/thing:906692>. The model is licensed under Creative Commons Attribution Share Alike license.

In computer graphics, most objects are not actually smooth, but rather represented by a polytope. In other words, they are obtained by identifying vertices on the surface of an object and connecting those vertices with straight lines (edges). Thus, the surface of an object actually consists of flat faces. In 3D graphics the faces are usually triangular. There is a simple mathematical reason for this: there always exists a plane passing through any three points in a space but not necessary through four or more points. If the number of faces is large, then the object observed from far away appears to be smooth.

Two things are necessary to create a three-dimensional computer model of an object: coordinates of the vertices and the edges showing the connections between the vertices. The former can be given in a form of an  $n \times 3$  array  $V$ , where  $n$  is the number of vertices in a 3D model. The latter can be given in several different ways, for instance, by an  $n \times n$  adjacency matrix  $E$  (hence, a 3D model can be treated as a graph!) or by an  $m \times 3$  array containing numbers of vertices in each face of the model where  $m$  is the number of faces. We will look at both of these descriptions.

Observe that this project is intended as a demonstration of techniques used to create a projection of a three dimensional object on a computer screen. Of course, Matlab has built in procedures to work with 3D objects. We will explore these procedures at the end of the project as well.

## TASKS

1. First of all, let us create a user defined function which will return a rotation matrix based on the user-specified angles  $\theta_x$ ,  $\theta_y$ , and  $\theta_z$  which correspond to the rotations around the axes  $0x$ ,  $0y$ , and  $0z$ . Create a new Matlab script and save it as `rotation.m` in your working directory. Define the header of this function as:

```
function rotmat=rotation(theta_x,theta_y,theta_z)
```

This function takes in three variables `theta_x`, `theta_y`, `theta_z` and returns the  $3 \times 3$  rotation matrix `rotmat`.

Variables: `theta_x`, `theta_y`, `theta_z`

2. Within the function, create the matrices `Rx`, `Ry`, and `Rz` as defined above by using standard Matlab syntax and the trigonometric functions `sin`, `cos`.

Variables: `Rx`, `Ry`, `Rz`

3. Compute the product of these matrices and assign the result to the output variable `rotmat`.

Variables: `rotmat`

4. Save the resulting function in the file `rotation.m` in your working directory. Now that the user-defined function is created, we can use it to generate projections.

5. Let us start by creating a projection of a simple object such as a cube. Open the file `m551lab11.m` and notice the following code in the first code cell `% Cube`:

```
Vertices=[1 1 1;...
          -1 1 1;...
           1 -1 1;...
           1 1 -1;...
          -1 -1 1;...
          -1 1 -1;...]
```

```

1 -1 -1;...
-1 -1 -1];

Edges=zeros(8,8);
Edges(1,2)=1;
Edges(1,3)=1;
Edges(1,4)=1;
Edges(2,5)=1;
Edges(2,6)=1;
Edges(3,5)=1;
Edges(3,7)=1;
Edges(4,6)=1;
Edges(4,7)=1;
Edges(5,8)=1;
Edges(6,8)=1;
Edges(7,8)=1;

Edges=Edges+Edges';

```

This code will create the matrix of the vertices of the cube and the corresponding adjacency matrix.

Variables: **Vertices, Edges**

- Assign the following values to the angles of the rotations  $\theta_x = \pi/3$ ,  $\theta_y = \pi/4$ , and  $\theta_z = \pi/6$ . Call the function `rotation` to generate the rotation matrix with these angles:

```
rotmat=rotation(theta1,theta2,theta3);
```

Variables: **rotmat**

- Transform the coordinates of the vertices with the rotation matrix.

```
VertRot=Vertices*rotmat';
```

Variables: **VertRot**

- Create a new figure window. Moving through the elements of the matrix **Edges**, draw a projection of the cube:

```

figure;
axis equal;
hold on;
for j=1:8
    for k=j+1:8
        if (Edges(j,k)==1)
            line([VertRot(j,1),VertRot(k,1)],[VertRot(j,2),VertRot(k,2)]);
        end;
    end;
end;
hold off;

```

The function `line` draws a line through two points given by their coordinates. The points can have two or three coordinates. Observe that to generate the projection, we dropped the last coordinate of the vector **VertRot**.

Q1: Why does the second `for` cycle start with `j+1`?

9. Modify the code above to obtain the projection generated by dropping the second coordinate of the vector **VertRot**.
10. Let us consider a more complicated object. Matlab has a built in buckyball (**bucky**) procedure which produces an array containing the vertices of a buckyball and the adjacency matrix showing how to connect the vertices:

```
[Edges2,Vertices2] = bucky;
```

Variables: **Edges2**, **Vertices2**

11. Using the same procedure as above, produce a projection of the buckyball onto the screen. You can use the same rotation matrix **rotmat** as before. Find the number of vertices in the array **Vertices2** using the **size** function.
12. To draw a three dimensional buckyball, use the following code:

```
figure;
axis equal;
hold on;

for j=1:60
    for k=j+1:60
        if (Edges2(j,k)==1)
            line([Vertices2(j,1),Vertices2(k,1)],[Vertices2(j,2),Vertices2(k,2)],...
                [Vertices2(j,3),Vertices2(k,3)]);
        end;
    end;
end;

hold off;
```

Observe that you can use the “3D rotation” button in the top panel of the figure window to rotate the figure. If you use the same button on the previous figure, you will notice that the generated projection is flat. If you did everything correctly then you will see something similar to the Fig. 11.

**Q2: Why do we use 60 as the end limit for our for cycles?**

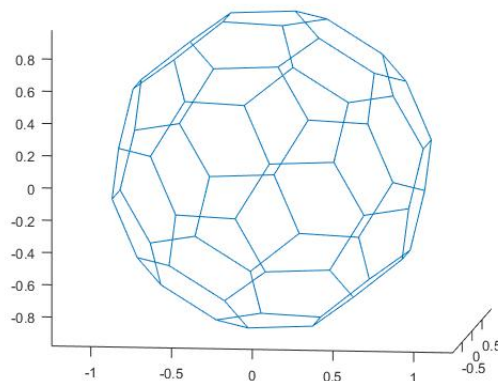


Figure 11: Buckyball in Matlab window.

13. Finally, this can be applied to any three dimensional construction. To illustrate this consider a model for 3D printing<sup>11</sup>. Load the matrices **v** and **f** into Matlab from the files **v.mat** and **f.mat** using **load** command. The array **v** contains the vertices of the model, and the array **f** contains the triangular faces of the model. Each row of the face array **f** contains the numbers which identify the vertices in the face. The corresponding coordinates of the vertices can then be found by referring to the array **v**.

Variables: **v**, **f**

14. Determine the size of the matrix **f** by using **size** function. Save the dimensions of this matrix as **mFaces**, **nFaces**. The number of rows **mFaces** in the array **f** will give you the number of faces in the model, while the number of columns **nFaces** will give you the number of vertices in each face (it should be equal to three).

Variables: **mFaces**, **nFaces**.

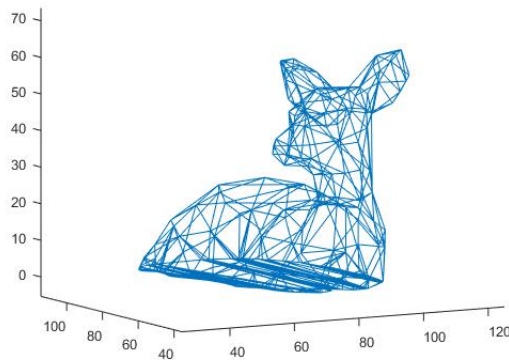


Figure 12: 3D model of a fawn.

15. To generate the three dimensional model, use the following code:

```
figure;
axis equal;
hold on;
for j=1:mFaces
    line([v(f(j,1),1),v(f(j,2),1)], [v(f(j,1),2),v(f(j,2),2)], ...
        [v(f(j,1),3),v(f(j,2),3)]);
    line([v(f(j,1),1),v(f(j,3),1)], [v(f(j,1),2),v(f(j,3),2)], ...
        [v(f(j,1),3),v(f(j,3),3)]);
    line([v(f(j,2),1),v(f(j,3),1)], [v(f(j,2),2),v(f(j,3),2)], ...
        [v(f(j,2),3),v(f(j,3),3)]);
end;
hold off;
```

If you did everything correctly you will see a figure similar to Fig. 12.

16. Using a similar procedure as above, generate a two dimensional projection of the three dimensional model. You can use the same rotation matrix **rotmat** as before. Store the

<sup>11</sup>The original data for the model is available from <https://www.thingiverse.com/thing:906692/#files>

coordinates of the projection in the matrix `VertRot`.

Variables: `VertRot`

17. Another way to generate an image like that in Fig. 17 is as follows.

```
% create rotation matrix
rotmat2 = rotation(-pi/3,0,pi/4);
% rotate vertices
vRot = v*rotmat2';
% project to xy plane
vPrj = vRot(:,1:2);
% draw
figure;
axis equal;
hold on;
for j=1:mFaces
line([vPrj(f(j,1),1),vPrj(f(j,2),1)], [vPrj(f(j,1),2),vPrj(f(j,2),2)]);
line([vPrj(f(j,1),1),vPrj(f(j,3),1)], [vPrj(f(j,1),2),vPrj(f(j,3),2)]);
line([vPrj(f(j,2),1),vPrj(f(j,3),1)], [vPrj(f(j,2),2),vPrj(f(j,3),2)]);
end;
hold off;
```

Variables: `rotmat2`, `vRot`, `vPrj`

## Project 10: Discrete dynamical systems, linear transformations of the plane, and the Chaos Game

**Goals:** To use matrix mappings of the plane  $\mathbb{R}^2$  in the form of a Chaos Game in order to produce fractals.

**To get started:** Download the Matlab script `lab10.m`<sup>12</sup> and save it in your working directory.

**Matlab commands used:** `linspace`, `for... end`, `rand`, `randi`, `plot`, `hold on...`, `hold off`, `cos`, `sin`

**What you have to submit:** The file `lab10.m` which you will modify during the lab session.

### INTRODUCTION

Many processes can be described by using dynamical systems. The main idea here is that the state of a system in the current moment may be derived by knowing the state of the system at the previous moment(s) of time. For instance, if we know that the velocity of a car increases with the acceleration  $a = 2\text{ m/s}^2$  and the initial velocity of the car is  $v_0 = 15\text{ m/s}$ , then we can easily compute the velocity of the car after  $t$  seconds by using the formula  $v = at + v_0$ . This is one of the simplest examples of dynamical systems. Motion of celestial bodies and population dynamics of species are some of the examples of dynamical systems in nature. If the time  $t$  is measured continuously then we are talking about continuous dynamical systems. If the time is measured discretely (only at certain points), then we are talking about discrete dynamical systems. For instance, the population of certain species at any moment of time depends on its population in the previous years. Since offspring for many species is produced only once a year, we can look at this population dynamics as a discrete dynamical system with time measured in years.

In this project we will explore the connection between fractals and matrix (linear) mappings in the form of the so-called Chaos Game. The Chaos Game is an algorithm first proposed by M. Barnsley in 1998 in his book “Fractals everywhere”.

The basic idea of the Chaos Game is simple. Pick  $n$  points on the plane (say, vertices of a regular  $n$ -gon), and a number  $r$ ,  $0 < r < 1$ . Pick a random point inside the polygon. Then repeat the following steps: pick a polygon vertex at random and draw the next point at a fraction  $r$  of the distance between the current point and the chosen vertex of the polygon. Repeat the process with the new point.

This process can be written in the form of the following linear mapping. Let  $\mathbf{x} = (x_1, x_2)$  be the current point, and let  $\mathbf{v} = (v_1, v_2)$  be the selected vertex of the polygon. Then the next point can be generated by using the linear mapping  $T$ :

$$T(\mathbf{x}) = \begin{bmatrix} T_{11} & T_{12} \\ T_{21} & T_{22} \end{bmatrix} (\mathbf{x} - \mathbf{v}) + \mathbf{v}.$$

Here  $T(\mathbf{x})$  is the next point. Repeated many times, this process sometimes (but not always) produces a fractal. The mapping  $T$  can change from iteration to iteration or can be picked at random with different probabilities (we will look at the examples of this type later). In the tasks below, we will look at some limiting sets of points which can be obtained by this procedure.

---

<sup>12</sup>The printout of the file `lab10.m` can be found in the appendices of this book.



## TASKS

1. Let us start simple by creating an image of the Sierpinski Triangle using the Chaos Game. Locate the cell titled `%% Sierpinski Triangle`. In this cell, define the vertices of an equilateral triangle by using the following commands

```
clear;
t=linspace(0,2*pi,4);
t(4)=[];
v=[cos(t); sin(t)];
```

This code creates a set of vertices for equilateral triangle. Each vertex is a distance of 1 unit away from the origin and the first vertex is located on the  $Ox$  axis.

Variables: `t, v`

Q1: What does the command `t(4)=[]` do?

2. Next, define the starting point and the matrix of the linear transformation:

```
T=[1/2 0; 0 1/2];
x(:,1)=[rand-0.5; rand-0.5];
```

The linear transformation with the matrix `T` is equivalent to multiplying the vector by  $1/2$ , and the starting point `x(:,1)` is picked at random.

Variables: `T, x`

3. Plot the vertices of the triangle on the plane and select the number of iterations of the process:

```
plot(v(1,:),v(2:,:), 'k*', x(1,1), x(2,1), 'b. ');
axis equal;
hold on;
Num=10000;
```

All the points which will be obtained during the iterative process will be stored in the  $2 \times N$  array `x` where  $N$  is the number of iterations. After computing all the iterations we need, we will plot all the points from the array `x` using the command `plot`. Observe that while it is possible to immediately plot the point we obtain at each step, multiple calls to the `plot` function take a significant amount of computational time and, thus, doing so is impractical.

Variables: `Num`

4. The following code creates the iterated points and plots them:

```
for j=1:Num
    k=randi(3);
    x(:,j+1)=T*(x(:,j)-v(:,k))+v(:,k);
end;
plot(x(1,:),x(2,:), 'b. ');
hold off;
```

In this code, we select an integer number `k` from 1 to 3 at random by using the command `randi`. We then do a linear transformation with the matrix `T`, the selected vertex `v(:,k)`, and the point `x(:,j)` obtained on the previous step. Running this code with a large number of points may take some time. If you did everything correctly, you should obtain

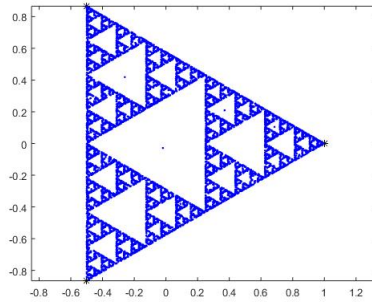


Figure 13: The Sierpinski triangle

an image similar to Fig. 13. The resulting picture approximates a fractal called the Sierpinski triangle.

There is a different procedure to generate this fractal starting with the interior of an equilateral triangle. At each step we will connect the mid-points of the sides of the triangle by straight lines. This subdivides the triangle into four triangles with the sides twice smaller than the original. Now we will throw out the middle triangle. Then we repeat this process with each of the leftover triangles, and so on. Doing this infinitely many times will produce the Sierpinski triangle as well.

**Variables:  $\mathbf{x}$**

5. Let us try some variations of the previous procedure. First of all, let us change the ratio in which the segment is divided. Locate the code cell titled `%% Sierpinski triangle with a change of ratio` and run the previous code with  $T = \begin{bmatrix} 1/3 & 0 \\ 0 & 1/3 \end{bmatrix}$ . Don't forget to clear the variables first (use the command `clear`)!
6. We can also add rotation to each of the iterations. Locate the code cell titled `%% Sierpinski triangle with rotation`. Use the original algorithm for Sierpinsky triangle but change the matrix  $T$  to the following::

```
theta=pi/18;
T=1/2*[cos(theta) -sin(theta); sin(theta) cos(theta)];
```

and run the procedure again. Observe that the matrix  $T$  is equivalent to two transformations: the shrinking of the plane by  $1/2$  and the counterclockwise rotation of the plane around the origin with the angle of rotation `theta`.

7. Next, let us look at the fractal generated starting from four vertices of a square. Locate the code cell titled `%% Square` and run the following code:

```
T=[1/2 0; 0 1/2];
t=linspace(0,2*pi,5);
t(5)=[];
v=[cos(t); sin(t)];
```

Now repeat the iterative procedure allowing the vertex at each step to be randomly selected from the four vertices of the square. Generate the points and plot them on a new figure. Observe that no fractal has been created in this case.

8. Locate the code cell titled `%% Square with different ratio`. Change the ratio in which the segment is divided to  $1/3$  and run the code again (don't forget to clear the variables!) with the modified matrix `T`:

```
T=[1/3 0; 0 1/3];
```

**Q2: Does the code generate a fractal this time?**

9. Let us try to do more sophisticated things. Locate the code cell titled `%% Square with vertex preference` (the new vertex cannot be the same as the previous) which contains the following code:

```
clear;
T=[1/2 0; 0 1/2];
t=linspace(0,2*pi,5);
t(5)=[];
v=exp(1i*t);
x(:,1)=[rand-0.5; rand-0.5];
plot(real(v),imag(v),'k*',x(1,1),x(2,1),'b. ');
axis equal;
hold on;
Num=5000;
k1=0;
for j=1:Num
    k=randi(3);
    if (k>=k1)
        k=k+1;
    end;
    w=[real(v(k));imag(v(k))];
    x(:,j+1)=T*(x(:,j)-w)+w;
    k1=k;
end;
plot(x(1,:),x(2,:),'b. ');
hold off;
```

This code repeats the process for the square except that it does not allow the same vertex to be selected twice in a row.

10. Try the next code cell titled `%% Square with vertex preference` (the new vertex cannot be opposite of the previous)

```
%% Square with vertex preference (new vertex cannot be opposite of the previous):
clear;
T=[1/2 0; 0 1/2];
t=linspace(0,2*pi,5);
t(5)=[];
v=exp(1i*t);
x(:,1)=[rand-0.5; rand-0.5];
plot(real(v),imag(v),'k*',x(1,1),x(2,1),'b. ');
axis equal;
hold on;
Num=5000;
w=[real(v(1));imag(v(1))];
k1=0;
```

```

for j=1:Num
    k=randi(4)
    if (k~=k1+2)&&(k1~=k+2)
        w=[real(v(k));imag(v(k))];
        x(:,j+1)=T*(x(:,j)-w)+w;
        k1=k;
    else
        x(:,j+1)=x(:,j);
    end;
end;
plot(x(1,:),x(2,:), 'b. ');
hold off;

```

This code repeats the process above except that it does not allow for the new vertex to lie opposite the old one.

11. Run the next code cell titled `%% Barnsley fern`. The code in this cell creates the so-called Barnsley fern which is a fractal with striking resemblance to a fern leaf (a lot of plants and animals in nature use fractals to their advantage!):

```

%% Barnsley fern
clear;

T1=[0.85,0.04;-0.04,0.85];
T2=[-0.15,0.28;0.26,0.24];
T3=[0.2,-0.26;0.23,0.22];
T4=[0,0;0,0.16];
Q1=[0;1.64];
Q2=[-0.028;1.05];
Q3=[0;1.6];
Q4=[0,0];
P1=0.85;
P2=0.07;
P3=0.07;
P4=0.01;

Num=15000;
x(:,1)=rand(2,1);
plot(x(1,1),x(2,:), 'b. ')
axis equal;
hold on;
for j=1:Num
    r=rand;
    if r<=P1
        x(:,j+1)=T1*x(:,j)+Q1;
    elseif r<=P1+P2
        x(:,j+1)=T2*x(:,j)+Q2;
    elseif r<=P1+P2+P3
        x(:,j+1)=T3*x(:,j)+Q3;
    else
        x(:,j+1)=T4*x(:,j)+Q4;
    end
end

plot(x(1,:),x(2,:), 'b. ');

```

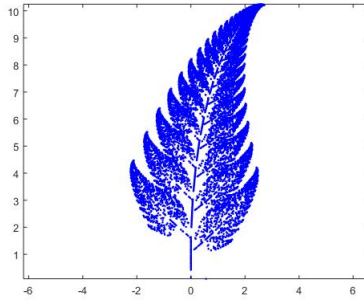


Figure 14: Barnsley Fern

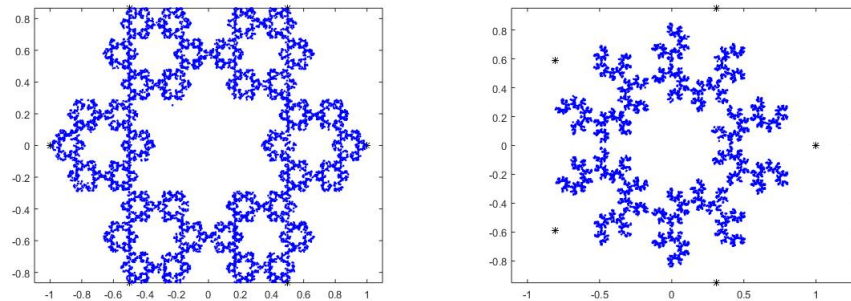


Figure 15: Fractals

```
hold off;
```

This code alternates between four different linear mappings given by  $y = T_j * x + Q_j$ ,  $j = 1, 2, 3, 4$ , where the matrices  $T_j$  and the vectors  $Q_j$  are given in the code. The probabilities of each mapping being selected are not equal as well and are given by the numbers P1, P2, P3, and P4 in the code. If you did everything correctly, you should see the image similar to the one in Fig. 14.

12. Modify the initial code for the Sierpinski triangle to create a fractal based on a regular hexagon. Take the matrix T to be

```
T=[1/4 0; 0 1/4];
```

If you did everything correctly, you should see the figure similar to the first image in Fig. 15.

Q3: Experiment with the ratio. What is the largest ratio you can find such that the little hexagons don't touch (two digits after the decimal point are sufficient)?

13. Finally, create another Chaos Game. Use the matrix  $T=[2/5 \ 0; \ 0 \ 2/5]$  for the linear mapping and the vertices of a regular pentagon with the restriction that each new vertex cannot be the same as the previous one. Plot the result in a new figure window. If you did everything correctly, you should see the figure similar to the second image in Fig. 15.

## Project 11: Projections, eigenvectors, Principal Component Analysis, and face recognition algorithms

**Goals:** To use eigenvalues and eigenvectors of a matrix to create a simple face recognition algorithm.

**To get started:**

- Download the file `lab11.m` and put it in your working directory<sup>13</sup>.
- Download the archive file `database.zip` and unpack it into your working directory. This will produce a folder “database”, which contains 33 image files in the format .pgm<sup>14</sup>.

**Matlab commands used:** `imread`, `size`, `reshape`, `num2str`, `mean`, `uint8`, `double`, `ones`, `eig`, `diag`

**What you have to submit:** The file `lab11.m` which you will modify during the lab session.

### INTRODUCTION

In this project we will apply principal component analysis (PCA) to create a simple face recognition algorithm. The PCA algorithm works by identifying patterns in the data which allows one to reduce the dimension of the dataset. We will use the following steps to apply the PCA algorithm to face recognition:

- Download the database of  $N$  images (in our case,  $N = 30$ ). All images must have the same dimensions  $m \times n$ .
- Input the images `person1.pgm-person30.pgm` into Matlab as arrays. Reshape each of the arrays as a column vector  $\mathbf{X}_i$ ,  $i = 1, \dots, N$ .
- Compute the mean of all images  $\mathbf{X}_{mean} = \frac{1}{N} \sum_{i=1}^N \mathbf{X}_i$ .
- Subtract the mean image from each of the vectors  $\mathbf{X}_i$ :  $\tilde{\mathbf{X}}_i = \mathbf{X}_i - \mathbf{X}_{mean}$  and create a matrix  $\mathbf{P}$ , the  $i$ th column of which is equal to the vector  $\tilde{\mathbf{X}}_i$ . The matrix  $\mathbf{P}$  has 30 columns and  $mn$  rows where  $m, n$  are the dimensions of the images.
- We are interested in the eigenvalues and eigenvectors of the matrix  $\mathbf{P}\mathbf{P}^T$ . However, the matrix  $\mathbf{P}\mathbf{P}^T$  is usually very large. For instance, if our database consists of the images with dimensions  $m \times n = 100 \times 100$ , then the matrix  $\mathbf{P}\mathbf{P}^T$  has the dimensions  $10000 \times 10000$ , and computation of eigenvalues and eigenvectors becomes unfeasible. Instead, we will use the fact that only at most  $N - 1$  eigenvalues of the matrix  $\mathbf{P}\mathbf{P}^T$  are non-zero (why?), and those non-zero eigenvalues are the same as non-zero eigenvalues of the matrix  $\mathbf{P}^T\mathbf{P}$  (why?). The matrix  $\mathbf{P}^T\mathbf{P}$  has the dimensions  $N \times N$  which are usually much smaller than the dimensions of the matrix  $\mathbf{P}\mathbf{P}^T$ . For instance, if there are 30 images in the database (like in our case), then the dimensions of  $\mathbf{P}^T\mathbf{P}$  are only  $30 \times 30$ . Thus, it is much more efficient to solve the eigenvalue problem for the matrix  $\mathbf{P}^T\mathbf{P}$ .

---

<sup>13</sup>The printout of the file `lab11.m` can be found in appendices of this book.

<sup>14</sup>These files represent a subset of the database of faces from AT&T Laboratories Cambridge [www.cl.cam.ac.uk/research/dtg/attarchive/facedatabase.html](http://www.cl.cam.ac.uk/research/dtg/attarchive/facedatabase.html).

- If  $\mathbf{u}_i$  is an eigenvector of the matrix  $\mathbf{P}^T\mathbf{P}$  corresponding to a non-zero eigenvalue  $\lambda_i$ , then  $\mathbf{v}_i = \mathbf{P}\mathbf{u}_i$  is the eigenvector of the matrix  $\mathbf{P}\mathbf{P}^T$  corresponding to  $\lambda_i$ . The eigenvectors of the matrix  $\mathbf{P}\mathbf{P}^T$  constitute the set of components of the data with the largest variance between them (which are called principal components).
- Compute the projections of your sample data onto the linear subspace generated by the vectors  $\mathbf{v}_i$ . The power of the algorithm is in the dimension reduction. Our initial images of faces “live” in the  $mn$ -dimensional real space. For instance, if the images are  $100 \times 100$  pixels, then the dimension of the space is equal to  $100^2 = 10000$ . If we would try to compare these images to each other pixel by pixel, we would need to compare 10000 pixels. However, not all of these pixels are significant. Principal Component Analysis allows us to identify the important information about the image and compare only that information. For instance, if we are working with a database of 30 images, then we will need to compare only 29 numbers. That is a dimensional reduction from 10000 to only 29! The PCA algorithm is a work horse behind many modern techniques in data mining, including face recognition algorithms.

## TASKS

1. Open the file lab11.m and run the first code cell titled %% Input database files into the Matlab:

```
%%Input database files into Matlab
clear;
Database_Size=30;
%Reading images from the database. The image files should be located in the
%subfolder ‘‘database’’
for j=1:Database_Size
    image_read=imread(['database\person' num2str(j) '.pgm']);
    [m,n]=size(image_read);
    P(:,j)=reshape(image_read,m*n,1);
end;
```

This code will input the database images into Matlab, reshape each image from the rectangular array to a column vector, and create the matrix P. Observe the use of the string function `num2str` in the `imread` command and the use of the `reshape` function to change the dimensions of the arrays containing the image data. Instead of the rectangular arrays with the elements corresponding to the pixel colors we now have a set of long vectors where columns of each image have been “stacked” on top of each other.

Variables: Database\_Size, m, n, P

2. Compute the mean image by applying the `mean` function to the the matrix P row-wise. Display the mean image by using the function `imshow`.

```
%% Computing and displaying the mean face
mean_face=mean(P,2);
imshow(uint8(reshape(mean_face,m,n)))
```

Variables: mean\_face

3. Subtract the mean of the images from each of the columns of the matrix P:

```

%% Subtract the mean face
P=double(P);
P=P-mean_face*ones(1,Database_Size);

```

Notice the conversion to the double type of the array P.

4. Compute the covariance matrix  $\mathbf{P}^T\mathbf{P}$  and compute the eigenvalues and the eigenvectors of this matrix  $\mathbf{P}^T\mathbf{P}$  by using the built in function `eig`. The function `eig` returns two arrays, one of which has eigenvalues on the main diagonal, and the other has eigenvectors as columns. Observe that one of the eigenvalues is very close to zero.

```

%% Compute the covariance matrix of the set and its eigenvalues and
eigenvectors
[Vectors,Values]=eig(P'*P);
EigenVectors=P*Vectors;

```

Variables: `Vectors`, `Values`, `EigenVectors`

Q1: Why does the matrix  $\mathbf{P}^T\mathbf{P}$  have a zero eigenvalue?

5. Display the set of eigenfaces on the same figure by running the code cell %% Display the set of eigenfaces:

```

%% Display the set of eigenfaces
for j=2:Database_Size;
    if j==2
        EigenFaces=reshape(EigenVectors(:,j)+mean_face,m,n);
    else
        EigenFaces=[EigenFaces reshape(EigenVectors(:,j)+mean_face,m,n)];
    end;
end
EigenFaces=uint8(EigenFaces);
figure;
imshow(EigenFaces);

```

If you did everything correctly, you will see an image containing 29 eigenfaces, the first five of which are shown in Fig. 16. Observe that the mean face has been added back to the eigenfaces.

Variables: `EigenFaces`

6. The matrix  $\mathbf{P}\mathbf{P}^T$  is symmetric. It follows from the theorems of linear algebra that the eigenvectors corresponding to the distinct eigenvalues are orthogonal. Write code to verify the last statement. You will need to create the matrix

```
Products=EigenVectors'*EigenVectors
```

and observe that it is a diagonal matrix. This matrix contains all possible dot products of the eigenvectors of the matrix  $\mathbf{P}\mathbf{P}^T$ . The fact that the eigenvectors are orthogonal will be used in the following sections to find orthogonal projections.

Variables: `Products`

Q2: Why is the matrix  $\mathbf{P}\mathbf{P}^T$  symmetric?

7. In this project we will use the obtained eigenfaces to perform several operations. Our first task will be to “unmask” an altered face (sunglasses have been digitally added to the image). The main idea behind this code is in computing the projection of the vector of data (obtained from the altered image) onto the subspace spanned by the eigenvectors. This





Figure 16: The first five eigenfaces

projection is then displayed on the screen. Now run the code cell titled `%% Recognition of an altered image (sunglasses)`:

```
%% Recognition of an altered image (sunglasses)
image_read=imread(['database\person30altered1.pgm']);
U=reshape(image_read,m*n,1);
NormsEigenVectors=diag(Products);
W=(EigenVectors'*(double(U)-mean_face));
W=W./NormsEigenVectors;
U_approx=EigenVectors*W+mean_face;
image_approx=uint8(reshape(U_approx,m,n));
figure;
imshow([image_read,image_approx])
```

In this code, we start by uploading the image `person30altered1.pgm` into Matlab. The image is then reshaped as a long vector similarly to the way it was done before. Then we compute the projection of the new vector with the subtracted vector `mean_face` onto the space spanned by the set of the eigenvectors and add the vector `mean_face` back. Finally, the reconstructed image is displayed. Observe that we get a very good approximation of the initial image without sunglasses.

**Variables: `image_read`, `U`, `NormsEigenVectors`, `W`, `U_approx`**

8. Modify the code above to unmask the image `person30altered2.pgm`.
9. Now let us compute the approximation of the face which is not in the database. To do this, modify the code above again to approximate the image `person31.pgm`. Observe, that the results of the last two approximations are somewhat worse than the very first one. This can be explained by the fact that there is an insufficient number of images in our database which do not span a wide enough variety of faces.

## Project 12: Matrix eigenvalues and the Google's PageRank algorithm

**Goals:** To apply matrix eigenvalues and eigenvectors to ranking of webpages in the World Wide Web.

**To get started:**

- Download the file `lab12.m` and put it in your working directory<sup>15</sup>.
- Download the file `AdjMatrix.mat` which contains the adjacency matrix of a so-called “wiki-vote” network with 8297 nodes<sup>16</sup> [9].

**Matlab commands used:** `load`, `size`, `numel`, `nnz`, `for... end`, `gplot`

**What you have to submit:** The file `lab12.m` which you will modify during the lab session.

### INTRODUCTION

According to social polls, the majority of users only look at the first few results of the online search and very few users look past the first page of results. Hence, it is crucially important to rank the pages in the “right” order so that the most respectable and relevant results will come first. The simplest way to determine the rank of a webpage in a network is to look at how many times it has been referred to by other webpages. This simple ranking method leaves a lot to be desired. In particular, it can be easily manipulated by referring to a certain webpage from a lot of “junk” webpages. The quality of the webpages referring to the page we are trying to rank should matter too. This is the main idea behind the Google PageRank algorithm.

The Google PageRank algorithm is the oldest algorithm used by Google to rank the web pages which are preranked offline. The PageRank scores of the webpages are recomputed each time Google crawls the web. Let us look at the theory behind the algorithm. As it turns out it, it is based on the theorems of linear algebra!

The main assumption of the algorithm is that if you are located on any webpage then with equal probability you can follow any of the hyperlinks from that page to another page. This allows to represent a webpage network as a directed graph with the webpages being the nodes, and the edges being the hyperlinks between the webpages. The adjacency matrix of such a network is built in the following way: the  $(i, j)$ th element of this matrix is equal to 1 if there is a hyperlink from the webpage  $i$  to the webpage  $j$  and is equal to 0 otherwise. Then the row sums of this matrix will represent numbers of hyperlinks from each webpage and the column sums will represent numbers of times each webpage has been referred to by other webpages.

Even further, we can generate a matrix of probabilities  $\mathbf{S}$  such that the  $(i, j)$ th element of this matrix is equal to the probability of traveling from  $i$ th webpage to  $j$ th webpage in the network. This probability is equal to zero if there is no hyperlink from  $i$ th page to  $j$ th page and is equal to  $1/N_i$  if there is a hyperlink from  $i$ th page to  $j$ th page, where  $N_i$  is the total number of hyperlinks from  $i$ th page. For instance, consider a sample network of only four webpages shown on the Fig. 17. The matrix  $\mathbf{S}$  for this network can be written as:

<sup>15</sup>The printout of the file `lab12.m` is given in the appendices of this book.

<sup>16</sup>The original network data is available here: <https://snap.stanford.edu/data/>.

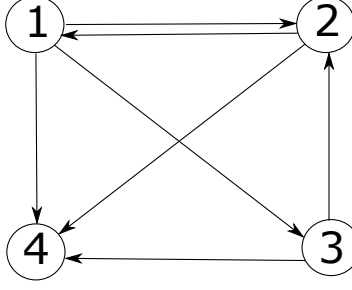


Figure 17: Sample network of four webpages

$$\mathbf{S} = \begin{bmatrix} 0 & 1/3 & 1/3 & 1/3 \\ 1/2 & 0 & 0 & 1/2 \\ 0 & 1/2 & 0 & 1/2 \\ 0 & 0 & 0 & 0 \end{bmatrix} \quad (1)$$

There are several issues which make working with the matrix  $\mathbf{S}$  inconvenient. First of all, there are webpages that do not have any hyperlinks - the so-called “dangling nodes” (such as the node 4 in Fig. 17). These nodes will correspond to zero rows of the matrix  $\mathbf{S}$ . Moreover, the webpages in the network may not be connected to each other and the graph of the network may consist of several disconnected components. These possibilities lead to undesirable properties of the matrix  $\mathbf{S}$  which make computations with it complicated and not even always possible.

The problem of the dangling nodes can be solved by assigning all elements of the matrix  $\mathbf{S}$  in the rows corresponding to the dangling nodes equal probabilities  $1/N$ , where  $N$  is the number of the nodes in the network. This can be understood in the following way: if we are at the dangling node we can with equal probability jump to any other page in the network. To solve the potential disconnectedness problem, we assume that a user can follow hyperlinks on any page with a probability  $1 - \alpha$  and can jump (or “teleport”) to any other page in the network with a probability  $\alpha$ . The number  $\alpha$  is called a damping factor. The value of  $\alpha = 0.15$  is usually taken in practical applications. The “teleport” surfing of the network can be interpreted as a user manually typing the webpage address in the browser or using a saved hyperlink from their bookmarks to move from one page onto another. The introduction of the damping factor allows us to obtain the Google matrix  $\mathbf{G}$  in the form:

$$\mathbf{G} = (1 - \alpha)\mathbf{S} + \alpha\mathbf{E},$$

where  $\mathbf{E}$  is a matrix with all the elements equal to  $1/N$ , where  $N$  is a number of webpages in the network.

The matrix  $\mathbf{G}$  has nice properties. In particular, it has only positive entries and all of its rows sum up to 1. In mathematical language, this matrix is *stochastic* and *irreducible* (you can look up the precise definitions of these terms if you are interested). The matrix  $\mathbf{G}$  satisfies the following **Perron-Frobenius theorem**:

**Theorem 1 (Perron-Frobenius)** *Every square matrix with positive entries has a unique unit eigenvector with all positive entries. The eigenvalue corresponding to this eigenvector is real and positive. Moreover, this eigenvalue is simple and is the largest in absolute value among all the eigenvalues of this matrix.*

Let us apply this theorem to the matrix  $\mathbf{G}$ . First of all, observe that the row sums of the

matrix  $\mathbf{G}$  are equal to 1. Consider the vector  $\mathbf{v}_1 = (1, 1, \dots, 1)^T/N$ . It is easy to see that

$$\mathbf{G}\mathbf{v}_1 = \mathbf{v}_1.$$

But then it follows that  $\mathbf{v}_1$  is the unique eigenvector with all positive components, and, therefore, by the Perron-Frobenius theorem,  $\lambda_1 = 1$  is the largest eigenvalue!

We are interested in the left eigenvector for the eigenvalue  $\lambda_1 = 1$ :

$$\mathbf{u}_1^T \mathbf{G} = \mathbf{u}_1^T.$$

Again, by the Perron-Frobenius theorem, the vector  $\mathbf{u}_1$  is the unique unit eigenvector with all positive components corresponding to the largest in absolute value eigenvalue  $\lambda_1 = 1$ . We will use the components of this vector for the ranking of webpages in the network.

Let us look at the justification behind this algorithm. We have already established that the vector  $\mathbf{u}_1$  exists. Consider the following iterative process. Assume that at the beginning a user can be on any webpage in the network with equal probability:

$$\mathbf{w}_0 = (1/N, 1/N, \dots, 1/N).$$

After 1 step (one move from one webpage to another using hyperlinks or teleporting), the probability vector of being on the  $i$ th webpage is determined by the  $i$ th component of the vector

$$\mathbf{w}_1 = \mathbf{w}_0 \mathbf{G}.$$

After two moves the vector of probabilities becomes

$$\mathbf{w}_2 = \mathbf{w}_1 \mathbf{G} = \mathbf{w}_0 \mathbf{G}^2,$$

and so on.

We hope that after a large number of steps  $n$ , the vector  $\mathbf{w}_n = \mathbf{w}_0 \mathbf{G}^n$  starts approaching some kind of limit vector  $\mathbf{w}_*$ ,  $\mathbf{w}_n \rightarrow \mathbf{w}_*$ . It turns out that due to the properties of the matrix  $\mathbf{G}$  this limit vector  $\mathbf{w}_*$  indeed exists and it is exactly the eigenvector corresponding to the largest eigenvalue, namely,  $\lambda_1 = 1$ . Moreover, numerical computation of matrix eigenvalues is actually based on taking the powers of the matrix (it is called the Power method) and not on solving the characteristic equation!

Let us assume that the vector  $\mathbf{w}_*$  is a non-negative vector whose entries sum to 1. Then the components of this vector represent the probabilities of being on each webpage in the network after a very large number of moves along the hyperlinks. Thus, it is perfectly reasonable to take these probabilities as ranking of the webpages in the network.

## TASKS

1. Open the file `lab12.m`. In the code cell titled `%%Load the network data` load the data from the file `AdjMatrix.mat` into Matlab by using the `load` command. Save the resulting matrix as `AdjMatrix`. Observe that the adjacency matrices of real networks are likely to be very large (may contain millions of nodes or more) and sparse. Check the sparsity of the matrix `AdjMatrix` using the functions `numel` and `nnz`. Denote the ratio of non-zero elements as `nnzAdjMatrix`. If you did everything correctly you should obtain that only 0.15% of the elements of the matrix `AdjMatrix` are non-zero.

Variables: `AdjMatrix`, `nnzAdjMatrix`

2. Check the dimensions of the matrix `AdjMatrix` using the `size` function. Save the dimensions as new variables `m` and `n`.

Variables: `m`, `n`

3. Observe that while the network described by the matrix `AdjMatrix` is not large at all from the viewpoint of practical applications, computations with this matrix may still take a noticeable amount of time. To save time, we will cut a subset out of this network and use it to illustrate the Google PageRank algorithm. Introduce a new variable `NumNetwork` and set its value to 500. Then cut a submatrix `AdjMatrixSmall` out of the matrix `AdjMatrix` and plot the graph represented by the matrix `AdjMatrixSmall` by running the following code cell:

```
%% Display a small amount of network
NumNetwork=500;
AdjMatrixSmall=AdjMatrix(1:NumNetwork,1:NumNetwork);
for j=1:NumNetwork
    coordinates(j,1)=NumNetwork*rand;
    coordinates(j,2)=NumNetwork*rand;
end;
gplot(AdjMatrixSmall,coordinates,'k-*');
```

This will plot the subgraph of the first 500 nodes in the network with random locations of the nodes. Notice the use of the function `gplot` to produce this graph. Observe that Matlab has special functions `graph` and `digraph` for working with graphs, but those functions are a part of the special package “Graph and Network Algorithms” which may not be immediately available. Simpler methods, as shown above, will be sufficient for our purposes.

Variables: `AdjMatrixSmall`, `coordinates`, `NumNetwork`

4. Set the parameter  $\alpha = 0.15$ . Introduce the vector  $w_0 = (1, 1, \dots, 1) / \text{NumNetwork}$ , and compute the consequent vectors  $w_1 = w_0 G$ ,  $w_2 = w_1 G$ ,  $w_3 = w_2 G$ ,  $w_5 = w_0 G^5$ ,  $w_{10} = w_0 G^{10}$ . Compute the difference  $\delta w = w_{10} - w_5$ . Observe that the sequence  $w_n$  converges to a certain limit vector  $w_*$  very fast.

Variables: `w0`, `w1`, `w2`, `w3`, `w5`, `w10`, `deltaw`

5. Compute the eigenvalues and the left and the right eigenvectors of the matrix  $G$  using the function `eig`. Observe that the right eigenvector corresponding to the eigenvalue  $\lambda_1 = 1$  is proportional to the vector  $v_1 = (1, 1, \dots, 1)$ . To compute the left eigenvectors, use the function `eig` on the matrix  $G'$ . Select the left eigenvector corresponding to the eigenvalue  $\lambda_1 = 1$  and denote it as `u1`.

Variables: `u1`

6. Observe that by default the vector `u1` is not scaled to have all positive components (even though all the components of the vector `u1` will have the same sign). Normalize this vector by using the code:

```
u1=abs(u1)/norm(u1,1);
```

This will create a vector with all positive components whose entries sum to 1 (called a probability vector).

7. Use the function `max` to select the maximal element and its index in the array.

Variables: `MaxRank`, `PageMaxRank`

8. Find out whether the highest ranking webpage is the same as the page with the most hyperlinks pointed to it. To do so, create the vector of column sums of the matrix **G** and save it as **MostLinks**. Use the function **max** again to select the element with the maximal number of links.

Variables: **MostLinks**, **MaxLinks**, **PageMaxLinks**

9. Compare if **MaxRank** and **MaxLinks** are the same.

Q1: What is the number of hyperlinks pointing to the webpage **MaxRank**?

(look at the vector **MostLinks** to find out).

## Project 13: Social networks, clustering, and eigenvalue problems

**Goals:** To apply linear algebra techniques to data clustering.

**To get started:**

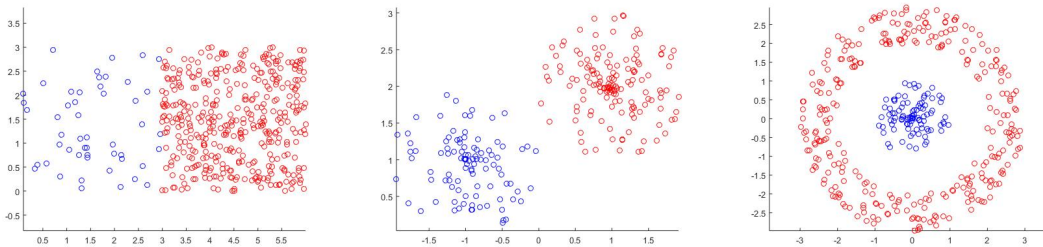
- Download the file `lab13.m` and put it in your working directory<sup>17</sup>.
- Download the file `social.mat` which contains the adjacency matrix of a simulated social network<sup>18</sup>.

**Matlab commands used:** `sum`, `diag`, `eig`, `for...end`, `if... then... end`, `load`, `clear`, `spy`, `eye`

**What you have to submit:** The file `lab13.m` which you will modify during the lab session.

### INTRODUCTION

Many practical applications deal with large amounts of data. One of the standard problems in data science is to group this data in clusters containing similar items. In many cases, clustering can be easily accomplished by the human brain (take a look at the attached pictures).



In general, creating an algorithm which allows us to separate the data into clusters is a difficult task. The data in many cases is multidimensional and cannot be visualized easily. Even deciding how many clusters to select is not necessary an easy task. There are currently several widely used algorithms which allow for the separation of the data points by similarity. All these algorithms have their own advantages and disadvantages.

In this project, we will use the spectral clustering algorithm based on the Fiedler's vector. We will look at the graph representing an individual's profile on a social network. The profile has a total of 351 "friends". Some of these friends know each other and are friends with each other as well. This friendship network can be represented as a graph with nodes being friends and edges showing the friendship connection between the individuals. The goal of clustering is to separate this graph into components which are tightly connected within themselves and have fewer connections to the outside (the so-called communities). The method explored here is based on the theorem by M. Fiedler, who has shown that for a connected graph (meaning that there is a path between any two vertices) the eigenvector corresponding to the second smallest eigenvalue of the Laplacian of the graph allows the graph to be split into two maximally intraconnected components.

---

<sup>17</sup>The printout of the file `lab13.m` can be found in the appendices of this book.

<sup>18</sup>This data has been artificially created.

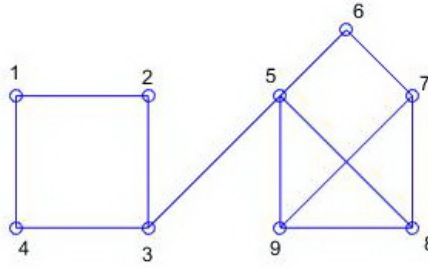


Figure 18: An example of a simple graph

## TASKS

1. We will start by considering an example of a simple graph: Create a new code cell with the title `%% Small graph`. Type in the adjacency matrix of this graph into Matlab and save it in a variable `AdjMatrix`.

Variables: `AdjMatrix`

2. Find the row sums of the matrix `AdjMatrix` (use the `sum` function) and save it as `RowSums`.

Variables: `RowSums`

3. Compute the Laplacian of the graph using the following command:

```
LaplaceGraph=diag(RowSums)-AdjMatrix;
```

Observe that the `diag(RowSums)` command creates a diagonal matrix with the vector `RowSums` on the main diagonal and zeros everywhere else. Observe also that the matrix `LaplaceGraph` is a symmetric positive semi-definite singular matrix. Recall that a matrix  $\mathbf{A}$  is positive semi-definite if  $\mathbf{xAx}^T \geq 0$  for all column vectors  $\mathbf{x}$ . To check that it is singular, multiply the matrix `LaplaceGraph` on the right by the column vector  $[1, 1, 1, 1, 1, 1, 1, 1, 1]^T$ .

Variables: `LaplaceGraph`

4. Find the eigenvalues and eigenvectors of the matrix `LaplaceGraph` using the `eig` function:

```
[V,D]=eig(LaplaceGraph);
```

Here the matrix `V` contains all the right eigenvectors of the matrix `LaplaceGraph` and the matrix `D` is a diagonal matrix with eigenvalues of the matrix `LaplaceGraph` on the main diagonal. Since the Laplacian is a symmetric graph, its eigenvalues are real. To simplify our task, let us sort the eigenvalues from smallest to largest. We will need to sort the eigenvectors along with them. Here is how to do that.

```
% sort eigenvalues and eigenvectors
[d,ind] = sort(diag(D));
D = D(ind,ind);
V = V(:,ind);
```



Variables: V, D

5. View the elements of the matrices **V** and **D** using the “Workspace” window of the Matlab environment. Observe that one of the eigenvalues of the matrix **LaplaceGraph** is zero (this is due to the fact that the matrix **LaplaceGraph** is singular). The eigenvector for this eigenvalue is proportional to the vector  $[1, 1, 1, 1, 1, 1, 1, 1, 1]^T$ . All other eigenvalues of the matrix **LaplaceGraph** are positive. This follows from the fact that the matrix **LaplaceGraph** is positive semi-definite and the graph corresponding to this Laplacian is connected. Find the smallest positive eigenvalue of the matrix **LaplaceGraph** (since we sorted the eigenvalues, it is the second diagonal element of the matrix **D**).
6. Identify the eigenvector corresponding to the second smallest eigenvalue:

```
V2=V(:,2);
```

In the next step, we will use the entries of **V2** to group the elements. To make sure we do this consistently, let us make sure that **V2** is the unit eigenvector with positive first entry:

```
% make sure the first entry is positive
if V2(1) < 0
    V2 = -V2;
end
```

Variables: V2

7. Separate the elements of the eigenvector **V2** onto positive and negative:

```
pos=[];
neg=[];
for j=1:9
    if V2(j)>0
        pos=[pos,j];
    else
        neg=[neg,j];
    end;
end;
```

If everything is done correctly, the arrays **pos** and **neg** will contain correspondingly 1, 2, 3, 4 and 5, 6, 7, 8, 9 which would be an intuitive clustering choice.

Variables: pos, neg

8. Now let us apply this technique to a larger graph when the results cannot be visualized so easily. To do this, load the matrix **Social** which is an adjacency matrix of a network of 351 Facebook friends. The  $(i, j)$ th element is equal to 1 if the  $i$ th and  $j$ th friends are also friends with each other and is equal to 0 otherwise:

```
%% Load the data
clear;
load('social.mat','Social');
spy(Social);
```

The command **spy(Social)** will plot the sparsity pattern of the adjacency matrix **Social** on a figure, putting blue dots for non-zero elements of the matrix (you should see something similar to the first image in Fig. 19). Observe that it is difficult to distinguish any kind of underlying pattern in this matrix. Our goal will be to cluster this set of data into two

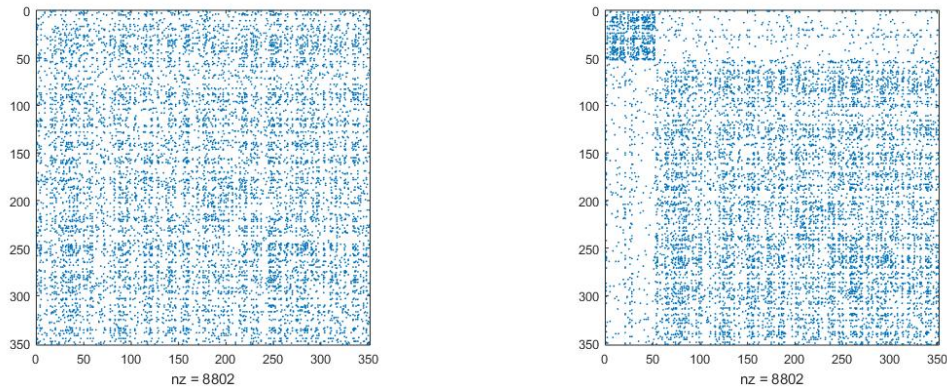


Figure 19: Adjacency matrices of the unsorted and sorted data.

maximally intraconnected groups.

Variables: **Social**

9. To do this, repeat the procedure we performed on a smaller matrix. Define the matrices **DiagSocial** and **LaplaceSocial** which are correspondingly the vector of row sums of the matrix **Social** and the Laplacian of this matrix.

Variables: **DiagSocial, LaplaceSocial**

10. Compute the eigenvalues and the eigenvectors:

```
[V,D]=eig(LaplaceSocial);
```

Variables: **V,D**

11. As before, identify the second smallest eigenvalue and the corresponding eigenvector **V2** and sort the vertices of the graph by the positive and negative components of the vector **V2**. Make the necessary adjustment to ensure that **V2(1)>0**.

Variables: **V2**

12. Permute the adjacency matrix **Social** using the permutation generated by the lists **pos**, **neg**:

```
order=[pos,neg];
[m,n]=size(Social);
iden=eye(m,m);
for j=1:351;
    for k=1:351;
        P(j,k)=iden(order(j),k);
    end;
end;
SocialOrdered=P*Social*P'
```

Variables: **order, P, SocialOrdered**

Q1: What does the command **P\*Social\*P'** do to the rows and columns of the matrix **Social**?

13. Plot the matrix `SocialOrdered` using the function `spy` again. If you did everything correctly, you will see the matrix separating into two denser connected subsets similarly to what is pictured in the second image in Fig. 19.
14. What if we want more clusters? There are two ways to proceed. First, we can explore the eigenvector `V3` corresponding to the third smallest eigenvalue. We will group the nodes in the four clusters: the ones which have positive components in both eigenvectors `V2`, `V3` (`++` group), the ones which have a positive component in `V2` and a negative component in `V3` (`+-`), the ones which have a negative component in `V2` and a positive component in `V3` (`-+`), and, finally, the ones with negative components in both vectors (`--`). This generates 4 distinct groups and can be accomplished by the code in the following code cell:

```
%% Cluster in 4 groups
V3=V(:,3);
if V3(1) < 0
    V3 = -V3;
end
pp=[];
pn=[];
np=[];
nn=[];
for j=1:m
    if (V2(j)>0)
        if (V3(j)>0)
            pp=[pp,j];
        else
            pn=[pn,j];
        end;
    else
        if (V3(j)>0)
            np=[np,j];
        else
            nn=[nn,j];
        end;
    end;
end;
order=[pp,pn,np,nn];
iden=eye(m,m);
for j=1:351;
    for k=1:351;
        P(j,k)=iden(order(j),k);
    end;
end;
SocialOrdered=P*Social*P'
figure;
spy(SocialOrdered)
```

15. An alternative way to obtain more clusters is to use the Fiedler vector procedure iteratively, meaning that we will apply it again to the clusters obtained in the previous step. This can be accomplished by running the following code:

```
%% Second order of Fiedler
SocialPos=Social(pos,pos);
SocialNeg=Social(neg,neg);
rowsumpos=sum(SocialPos,2);
```

```

DiagSocialPos=diag(rowsumpos);
LaplaceSocialPos=DiagSocialPos-SocialPos;
[VPos,DPos]=eig(LaplaceSocialPos);
V2Pos=VPos(:,2);
[mpos,npos]=size(V2Pos);
posp=[];
posn=[];
for j=1:mpos
    if (V2Pos(j)>0)
        posp=[posp,pos(j)];
    else
        posn=[posn,pos(j)];
    end;
end;
rowsumneg=sum(SocialNeg,2);
DiagSocialNeg=diag(rowsumneg);
LaplaceSocialNeg=DiagSocialNeg-SocialNeg;
[VNeg,DNeg]=eig(LaplaceSocialNeg);
V2Neg=VNeg(:,2);
[mneg,nneg]=size(V2Neg);
negp=[];
negn=[];
for j=1:mneg
    if (V2Neg(j)>0)
        negp=[negp,neg(j)];
    else
        negn=[negn,neg(j)];
    end;
end;
ordergen=[posp,posn,negp,negn];
iden=eye(m,m);
for j=1:351;
    for k=1:351;
        P(j,k)=iden(ordergen(j),k);
    end;
end;
SocialOrderedGen=P*Social*P'
figure;
spy(SocialOrderedGen)

```

Observe that the densely connected components obtained by the last two methods might be different. In the context of a social network, the dense clusters may represent groups of people who have common connections in real life. For instance, your family, your friends from high school, your colleagues, etc.

## Project 14: Singular Value Decomposition and image compression

**Goals:** In this project we will discuss a singular value decomposition of a matrix and its applications to image compression and noise filtering.

**To get started:**

- Download the file `lab14.m` and put it in your working directory<sup>19</sup>.
- Download the files `einstein.jpg`<sup>20</sup> and `checkers.pgm`<sup>21</sup> and put them in your working directory.

**Matlab commands used:** `clear`, `linspace`, `subplot`, `quiver`, `plot`, `hold on...`, `hold off`, `svd`, `axis`, `title`, `imread`, `imshow`, `size`, `min`, `for...`, `end`, `cos`, `sin`, `'`, `rand`, `double`, `ones`

**What you have to submit:** The file `lab14.m` which you will modify during the lab session.

### INTRODUCTION

Any  $m \times n$  matrix  $\mathbf{A}$  possesses a singular value decomposition of the form:

$$\mathbf{A} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^T,$$

where  $\mathbf{U}$  is an orthogonal  $m \times m$  matrix satisfying the condition  $\mathbf{U}^T\mathbf{U} = \mathbf{I}_m$ ,  $\mathbf{\Sigma}$  is an  $m \times n$  rectangular diagonal matrix with nonnegative values  $\sigma_1, \sigma_2, \dots, \sigma_{\min(m,n)}$  on the main diagonal, and  $\mathbf{V}$  is an orthogonal  $n \times n$  matrix satisfying the condition  $\mathbf{V}^T\mathbf{V} = \mathbf{I}_n$ . The nonnegative numbers  $\sigma_1, \sigma_2, \dots, \sigma_{\min(m,n)}$  are called the singular values of the matrix  $\mathbf{A}$ . They are arranged in the decreasing order:  $\sigma_1 \geq \sigma_2 \geq \dots \geq \sigma_{\min(m,n)} \geq 0$ . Observe that the singular value decomposition of a matrix is not unique.

Singular value decomposition is important for implementation of many numerical algorithms in linear algebra. In this project we will look at the geometric meaning of singular value decomposition and two applications related to image processing and noise filtering. Observe that more efficient image compression methods exist. The examples in this project are used mainly to show the reduction in the amount of data (so-called dimension reduction) which can be accomplished with singular value decomposition. Dimension reduction is an important tool in many practical applications dealing with large amounts of data, such as statistics, data science, and machine learning.

### TASKS

1. We will start with an illustration of the geometric meaning of singular value decomposition. Let us look at a singular value decomposition of a  $2 \times 2$  matrix. Open the file `lab16.m`, locate the code cell `%% 2x2 matrix`, and add the following commands

---

<sup>19</sup>The printout of the file `lab14.m` can be found in the appendices of this book.

<sup>20</sup>The image is available at: [https://commons.wikimedia.org/wiki/File:Albert\\_Einstein\\_Head.jpg](https://commons.wikimedia.org/wiki/File:Albert_Einstein_Head.jpg)

<sup>21</sup>Image from page 446 of "The standard Hoyle; a complete guide and reliable authority upon all games of chance or skill now played in the United States, whether of native or foreign introduction" (1909), image appears on Flickr Commons.

```

%% 2x2 matrix
clear;
t=linspace(0,2*pi,100);
X=[cos(t);sin(t)];
subplot(2,2,1);
hold on;
plot(X(1,:),X(2,:), 'b');
quiver(0,0,1,0,0, 'r');
quiver(0,0,0,1,0, 'g');
axis equal
title('Unit circle')
hold off;

```

This code will create a  $2 \times 100$  matrix  $\mathbf{X} = [\mathbf{x}_1 \ \mathbf{x}_2 \ \cdots \ \mathbf{x}_{100}]$  whose columns  $\mathbf{x}_i$  are unit vectors pointing in various directions. The plot will show a blue circle corresponding to the endpoints of these vectors and two vectors of the standard basis on the plane  $\mathbf{e}_1 = (1, 0)^T$  and  $\mathbf{e}_2 = (0, 1)^T$ . We use the function `subplot` which will create a plot containing four subplots arranged in two rows and two columns. The plot above will occupy the first “cell” of this plot. Observe the command `quiver` which draws a vector with the beginning point given by the first two arguments  $((0,0)$  in this case) and an ending point given by the next two arguments  $((1,0)$  or  $(0,1)$  in the code above).

Variables: **X**

- Now in the M-file, define a variable **A** holding the matrix

```
A = [ 2, 1; -1, 1 ];
```

and compute the singular value decomposition of this matrix using the `svd` function:

```
[U,S,V] = svd(A);
```

Using the workspace window of the Matlab main environment, check out the matrices **U**, **S**, **V**. Perform the commands

```
U'*U
V'*V
```

to ascertain that the matrices **U** and **V** are orthogonal. The output should produce  $2 \times 2$  identity matrices.

Variables: **A, U, S, V**

- Next, let us observe the geometric meaning of the individual matrices **U**,  **$\Sigma$** , **V** (**U**, **S**, **V** in our Matlab code) in the singular value decomposition. To do this, observe the transformations induced by these matrices on a unit circle and the vectors of the standard basis  $\mathbf{e}_1$ ,  $\mathbf{e}_2$ . Let us start by multiplying the coordinates of the points of the circle and the vectors  $\mathbf{e}_1$ ,  $\mathbf{e}_2$  by the matrix **V**. Execute the following code:

```

VX=V'*X;
subplot(2,2,2)
hold on;
plot(VX(1,:),VX(2,:), 'b');
quiver(0,0,V(1,1),V(1,2),0, 'r');
quiver(0,0,V(2,1),V(2,2),0, 'g');
axis equal
title('Multiplied by matrix V^T')
hold off;

```

Observe that the matrix  $VX$  contains the vectors of the matrix  $X$  transformed by the multiplication by the matrix  $V^T$ . Since the matrices  $V$  and  $V^T$  are orthogonal, multiplication by the matrix  $V^T$  is equivalent to rotation of a plane, possibly in combination with a reflection along some straight line. This allows us to conjecture that the image of the unit circle under this mapping will still be a unit circle, but the vectors of the basis will be rotated and possibly switched in orientation.

Variables:  $VX$

Q1: Did the multiplication by the transpose of the matrix  $V$  resulted in a reflection of the plane?

- Now, let us multiply the result from the previous step by the matrix  $\Sigma$  ( $S$  in the Matlab code). Observe that since the matrix  $\Sigma$  is diagonal, then multiplication by this matrix geometrically means stretching of the plain in two directions. To verify this, execute the following Matlab code:

```
SVX = S*VX;
subplot(2,2,3);
hold on;
plot(SVX(1,:),SVX(2,:), 'b');
quiver(0,0,S(1,1)*V(1,1),S(2,2)*V(1,2),0, 'r');
quiver(0,0,S(1,1)*V(2,1),S(2,2)*V(2,2),0, 'g');
axis equal
title('Multiplied by matrix \Sigma V^T')
hold off;
```

Observe that, as expected, the unit circle is stretched and becomes an ellipsis. The images of the standard basis vectors are stretched as well.

Variables:  $SVX$

- Finally, multiply the results from the last step by the matrix  $U$  to obtain:

```
AX = U*SVX;
subplot(2,2,4)
hold on;
plot(AX(1,:),AX(2,:), 'b');
quiver(0,0,U(1,1)*S(1,1)*V(1,1)+U(1,2)*S(2,2)*V(1,2),U(2,1)*S(1,1)*V(1,1)+...
      U(2,2)*S(2,2)*V(1,2),0, 'r');
quiver(0,0,U(1,1)*S(1,1)*V(2,1)+U(1,2)*S(2,2)*V(2,2),U(2,1)*S(1,1)*V(2,1)+...
      U(2,2)*S(2,2)*V(2,2),0, 'g');
axis equal
title('Multiplied by matrix U\Sigma V^T=A')
hold off;
```

Observe that the result is equivalent to multiplying the initial vector  $X$  by the matrix  $A$ . Since the matrix  $U$  is orthogonal, then the multiplication by this matrix should result in a rotation of the plane possibly combined with a reflection. Confirm this by observing the images of the basis vectors.

Variables  $AX$

Q2: Did multiplication by the matrix  $U$  produce a reflection of the plane?

- If you answered yes to both Q1 and Q2 above, can you modify the matrices  $U$  and  $V$  in such a way that no reflections of the plane occur? Produce the modified matrices  $U1$  and  $V1$  and confirm that  $U1*S*V1'=A$ . Observe that this shows that singular value decomposition

is not unique.

Variables **U1, V1**

7. Finally, observe that

$$[\mathbf{A}\mathbf{v}_1 \ \mathbf{A}\mathbf{v}_2] = \mathbf{A}\mathbf{V} = \mathbf{U}\mathbf{S}\mathbf{V}^T\mathbf{V} = \mathbf{U}\mathbf{S} = [\mathbf{u}_1 \ \mathbf{u}_2] \begin{bmatrix} \sigma_1 & 0 \\ 0 & \sigma_2 \end{bmatrix} = [\sigma_1\mathbf{u}_1 \ \sigma_2\mathbf{u}_2],$$

showing that

$$\mathbf{A}\mathbf{v}_1 = \sigma_1\mathbf{u}_1 \quad \text{and} \quad \mathbf{A}\mathbf{v}_2 = \sigma_2\mathbf{u}_2.$$

Check this fact numerically by computing the expression  $\mathbf{A}*\mathbf{V}-\mathbf{U}*\mathbf{S}$ .

8. Now we will look at image compression using SVD. Add the following commands to your M-file:

```
%% Image compression
clear;
ImJPG=imread('einstein.jpg');
figure;
imshow(ImJPG);
[m,n]=size(ImJPG);
```

This code loads an image as a `uint8` matrix and displays it on the screen. Each entry in the matrix corresponds to a pixel on the screen and takes a value somewhere between 0 (black) and 255 (white).

Variables: **ImJPG**

9. Perform a singular value decomposition of the matrix `ImJPG` and save the output in matrices `UIm`, `SIm`, and `VIm`. Since `ImJPG` is integer-valued, you will need to use `svd(double(ImJPG))`.

Variables: **UIm, SIm, VIm**

10. Plot the singular values using the code:

```
figure;
plot(1:min(m,n),diag(SIm));
```

This shows the singular values (the diagonal entries of the `SIm` matrix) for the image matrix `ImJPG`. Notice that the diagonal entries of `SIm` are ordered in the decreasing order  $\sigma_1 \geq \sigma_2 \geq \dots \geq \sigma_{\min(m,n)} \geq 0$ .

11. The idea behind compression of data with singular value decomposition is in the following. Using the singular value decomposition, the matrix  $\mathbf{A}$  can be written in the form:

$$\mathbf{A} = \sigma_1\mathbf{u}_1\mathbf{v}_1^T + \sigma_2\mathbf{u}_2\mathbf{v}_2^T + \dots + \sigma_r\mathbf{u}_r\mathbf{v}_r^T$$

where  $r$  is the rank of  $\mathbf{A}$  and  $\mathbf{u}_i$  and  $\mathbf{v}_i$  are the  $i$ th columns of  $\mathbf{U}$  and  $\mathbf{V}$  respectively.

Observe that since the singular values are arranged in the decreasing order, the first terms in this sum provide a larger contribution to the matrix  $\mathbf{A}$  than the subsequent terms. It may happen that for some  $k < r$ ,  $\sigma_{k+1}$  is small compared to  $\sigma_1$ , and correspondingly does not affect the matrix  $\mathbf{A}$  too much. We should then expect

$$\mathbf{A} \approx \sigma_1\mathbf{u}_1\mathbf{v}_1^T + \sigma_2\mathbf{u}_2\mathbf{v}_2^T + \dots + \sigma_k\mathbf{u}_k\mathbf{v}_k^T$$



to be a good approximation to the matrix  $\mathbf{A}$  (this is called a truncated SVD). In fact, there is a theorem in linear algebra which states that SVD truncated to the  $k$  terms is the best approximation to the matrix  $\mathbf{A}$  among the matrices of the rank at most  $k$  in the sense of Frobenius norm (which is equivalent to the regular Euclidean norm if we consider the matrix  $\mathbf{A}$  as an  $mn$ -dimensional vector).

This idea can be used for image compression as follows. Instead of storing the whole  $m \times n$  matrix  $\mathbf{A}$ , we can instead store the  $m \times k$  and  $n \times k$  matrices

$$\mathbf{C} = [\mathbf{u}_1 \ \mathbf{u}_2 \ \cdots \ \mathbf{u}_k] \quad \text{and} \quad \mathbf{D} = [\sigma_1 \mathbf{v}_1 \ \sigma_2 \mathbf{v}_2 \ \cdots \ \sigma_k \mathbf{v}_k].$$

If  $k$  is much smaller than  $\min(m, n)$ , then storing  $\mathbf{C}$  and  $\mathbf{D}$  will take much less space than storing  $\mathbf{A}$ . Moreover, if we wish to display the image, we can reconstruct the approximation of  $\mathbf{A}$  as  $\mathbf{A} \approx \mathbf{C}\mathbf{D}^T$ .

Add the following code to your M-file:

```
% Approximation of the original matrix with 50, 100, and 150 singular values
for k=50:50:150
    ImJPG_comp=uint8(UIm(:,1:k)*SIm(1:k,1:k)*(VIm(:,1:k))');
    figure;
    imshow(ImJPG_comp)
    % compression percentage
    pct = 1 - (numel(UIm(:,1:k))+numel(VIm(:,1:k)*SIm(1:k,1:k)))/numel(ImJPG);
    fprintf('Compression percentage for %2.0f singular values: %8.3f\n',k, pct);
end;
```

This code compresses the image as described above using  $k = 50$ ,  $k = 100$ , and  $k = 150$  singular values and displays the reconstructed image. Compare the reconstructed images with the original. The code also displays the compression percentage as `pct`.

Variables: `ImJPG_comp`, `pct`

12. Observe that the singular value decomposition can also be used to smooth noisy data, especially if the data contains patterns. Data smoothing is often necessary because all measurements contain small errors resulting in a “noise”. This noise usually determines the smallest singular values of the matrix. Dropping these small values, thus, not only saves the storage space, but also allows to eliminate noise from the data.

Start a new code cell. Load the file `checkers.pgm` into Matlab and add some noise to the resulting image matrix using the following code:

```
% Noise filtering
clear;
ImJPG=imread('checkers.pgm');
[m,n]=size(ImJPG);
ImJPG_Noisy=double(ImJPG)+50*(rand(m,n)-0.5*ones(m,n));
figure;
imshow(ImJPG);
figure;
imshow(uint8(ImJPG_Noisy));
```

Variables: `ImJPG`, `ImJPG_Noisy`

13. Compute the SVD of the matrix `ImJPG_Noisy` and save the resulting decomposition matrices as `UIm`, `SIm`, and `VIm`.

Variables: `UIm`, `SIm`, `VIm`

14. Compute the approximations of the initial image with  $k = 10$ ,  $k = 30$ , and  $k = 50$  singular values. Display the resulting approximations and compare them to the “noisy” image. Observe that SVD significantly reduces the noise. Compare the images to the initial image without noise. Observe also that even though SVD reduces the noise, it also somewhat blurs the image.

# Appendices

Printout of the file lab09.m

```
%%
clear;
Vertices=[1 1 1;...
          -1 1 1;...
           1 -1 1;...
           1 1 -1;...
          -1 -1 1;...
          -1 1 -1;...
           1 -1 -1;...
          -1 -1 -1];

Edges=zeros(8,8);
Edges(1,2)=1;
Edges(1,3)=1;
Edges(1,4)=1;
Edges(2,5)=1;
Edges(2,6)=1;
Edges(3,5)=1;
Edges(3,7)=1;
Edges(4,6)=1;
Edges(4,7)=1;
Edges(5,8)=1;
Edges(6,8)=1;
Edges(7,8)=1;

Edges=Edges+Edges';

theta1=pi/3;
theta2=pi/4;
theta3=pi/6;

rotmat=rotation(theta1,theta2,theta3);

VertRot=Vertices*rotmat;

figure;
axis equal;
hold on;

for j=1:8
    for k=j+1:8
        if (Edges(j,k)==1)
            line([VertRot(j,1),VertRot(k,1)], [VertRot(j,2),VertRot(k,2)]);
        end;
    end;
end;

hold off;

%% Buckyball
```

```

[Edges,Vertices] = bucky;
figure;
axis equal;
hold on;

theta1=pi/3;
theta2=pi/4;
theta3=pi/6;

rotmat=rotation(theta1,theta2,theta3);

VertRot=Vertices*rotmat;

for j=1:60
    for k=j+1:60
        if (Edges(j,k)==1)
            line([VertRot(j,1),VertRot(k,1)],[VertRot(j,2),VertRot(k,2)]);
        end;
    end;
end;

hold off;

figure;
axis equal;
hold on;

for j=1:60
    for k=j+1:60
        if (Edges(j,k)==1)
            line([Vertices(j,1),Vertices(k,1)],[Vertices(j,2),Vertices(k,2)],...
                [Vertices(j,3),Vertices(k,3)]);
        end;
    end;
end;

hold off;

%% 3D fawn
clear;

load('v.mat','v');
load('f.mat','f');

[mVert,nVert]=size(v);

[mFace,nFace]=size(f);

figure;
axis equal;
hold on;
for j=1:mFace
    line([v(f(j,1),1),v(f(j,2),1)],[v(f(j,1),2),v(f(j,2),2)],[v(f(j,1),3),v(f(j,2),3)]);

```

```

        line([v(f(j,1),1),v(f(j,3),1)], [v(f(j,1),2),v(f(j,3),2)], [v(f(j,1),3),v(f(j,3),3)]);
        line([v(f(j,2),1),v(f(j,3),1)], [v(f(j,2),2),v(f(j,3),2)], [v(f(j,2),3),v(f(j,3),3)]);
    end;
    hold off;

    figure;
    axis equal;
    hold on;

    theta1=pi/3;
    theta2=pi/4;
    theta3=pi/2;

    rotmat=rotation(theta1,theta2,theta3);

    VertRot=v*rotmat;

    for j=1:mFace
        line([VertRot(f(j,1),1),VertRot(f(j,2),1)], [VertRot(f(j,1),2),VertRot(f(j,2),2)]);
        line([VertRot(f(j,1),1),VertRot(f(j,3),1)], [VertRot(f(j,1),2),VertRot(f(j,3),2)]);
        line([VertRot(f(j,2),1),VertRot(f(j,3),1)], [VertRot(f(j,2),2),VertRot(f(j,3),2)]);
    end;

```

## Printout of the file lab10.m

```
% The Chaos Game

%% Sierpinski triangle

%% Sierpinski triangle with a change of ratio

%% Sierpinski triangle with rotation

%% Square

%% Square with different ratio

%% Square with vertex preference (new vertex cannot be the same as previous)
clear;

T=[1/2 0; 0 1/2];

t=linspace(0,2*pi,5);
t(5)=[];
v=[cos(t); sin(t)];

x(:,1)=[rand-0.5; rand-0.5];

plot(v(1,:),v(2,:), 'k*', x(1,1), x(2,1), 'b. ');
axis equal;
hold on;

Num=5000;

k1=0;
for j=1:Num
    k=randi(3);
    if (k>=k1)
        k=k+1;
    end;
    x(:,j+1)=T*(x(:,j)-v(:,k))+v(:,k);
    k1=k;
end;

plot(x(1,:),x(2,:), 'b. ');
hold off;

%% Square with vertex preference (new vertex cannot be opposite of the previous)
clear;

T=[1/2 0; 0 1/2];

t=linspace(0,2*pi,5);
t(5)=[];
v=[cos(t); sin(t)];

x(:,1)=[rand-0.5; rand-0.5];
```

```

plot(v(1,:),v(2,:), 'k*', x(1,1), x(2,1), 'b. ');
axis equal;
hold on;

Num=5000;

w=[real(v(1));imag(v(1))];
k1=0;

for j=1:Num
    k=randi(4)
    if (k~=k1+2)&&(k1~=k+2)
        x(:,j+1)=T*(x(:,j)-v(:,k))+v(:,k);
        k1=k;
    else
        x(:,j+1)=x(:,j);
    end;
end;

plot(x(1,:),x(2,:), 'b. ');
hold off;

%% Barnsley fern
clear;

T1=[0.85,0.04;-0.04,0.85];
T2=[-0.15,0.28;0.26,0.24];
T3=[0.2,-0.26;0.23,0.22];
T4=[0,0;0,0.16];
Q1=[0;1.64];
Q2=[-0.028;1.05];
Q3=[0;1.6];
Q4=[0,0];
P1=0.85;
P2=0.07;
P3=0.07;
P4=0.01;

Num=15000;
x(:,1)=rand(2,1);
plot(x(1,:),x(2,:), 'b. ')
axis equal;
hold on;
for j=1:Num
    r=rand;
    if r<=P1
        x(:,j+1)=T1*x(:,j)+Q1;
    elseif r<=P1+P2
        x(:,j+1)=T2*x(:,j)+Q2;
    elseif r<=P1+P2+P3
        x(:,j+1)=T3*x(:,j)+Q3;
    else
        x(:,j+1)=T4*x(:,j);
    end;
end;

```

```

        end
    end

    plot(x(1,:),x(2,:), 'b. ');
    hold off;

    %% Hexagon
    clear;

    T=[1/3 0; 0 1/3];

    t=linspace(0,2*pi,7);
    t(7)=[];
    v=[cos(t); sin(t)];

    x(:,1)=[rand-0.5; rand-0.5];

    plot(v(1,:),v(2,:), 'k*', x(1,1), x(2,1), 'b. ');
    axis equal;
    hold on;

    Num=10000;

    for j=1:Num
        k=randi(6);
        x(:,j+1)=T*(x(:,j)-v(:,k))+v(:,k);
    end;
    plot(x(1,:),x(2,:), 'b. ');
    hold off;

    %% Pentagon with a skipped vertex

    clear;
    T=[1/2.5 0; 0 1/2.5];

    t=linspace(0,2*pi,6);
    t(6)=[];
    v=[cos(t); sin(t)];

    x(:,1)=[rand-0.5; rand-0.5];

    plot(v(1,:),v(2,:), 'k*', x(1,1), x(2,1), 'b. ');
    axis equal;
    hold on;

    Num=10000;

    k1=0;

    for j=1:Num
        k=randi(5);
        if k~=k1
            x(:,j+1)=T*(x(:,j)-v(:,k))+v(:,k);

```



```
        k1=k;  
    else  
        x(:,j+1)=x(:,j);  
    end;  
end;  
plot(x(1,:),x(2,:), 'b. ');  
hold off;
```

## Printout of the file lab11.m

```
% Simple face recognition algorithm
%% Input database files into the Matlab
clear;
Database_Size=30;

%Reading images from the database. The image files should be located in the
%subfolder "database"
for j=1:Database_Size
    image_read=imread(['database\person' num2str(j) '.pgm']);
    [m,n]=size(image_read);
    P(:,j)=reshape(image_read,m*n,1);
end;

%% Computing and displaying the mean face

%% Subtract the mean face

%% Compute the covariance matrix of the set and its eigenvalues and eigenvectors

%% Display the set of eigenvaces
for j=2:Database_Size;
    if j==2
        EigenFaces=reshape(EigenVectors(:,j)+mean_face,m,n);
    else
        EigenFaces=[EigenFaces reshape(EigenVectors(:,j)+mean_face,m,n)];
    end;
end

EigenFaces=uint8(EigenFaces);

figure;
imshow(EigenFaces);

%% Orthogonality and symmetry

%% Recognition of an altered image (sunglasses)
image_read=imread(['database\person30altered1.pgm']);
U=reshape(image_read,m*n,1);

NormsEigenVectors=diag(Products);
W=(EigenVectors'*(double(U)-mean_face));
W=W./NormsEigenVectors;

U_approx=EigenVectors*W+mean_face;

image_approx=uint8(reshape(U_approx,m,n));
figure;
imshow([image_read,image_approx])

%% Recognition of an altered image (lower part of the face)

%% Recognition of a person not in the database
```

## Printout of the file lab12.m

```
% Google PageRank algorithm on the example of random network

%% Load the network data
clear;

%% Display a small amount of network
NumNetwork=500;
AdjMatrixSmall=AdjMatrix(1:NumNetwork,1:NumNetwork);

for j=1:NumNetwork
    coordinates(j,1)=NumNetwork*rand;
    coordinates(j,2)=NumNetwork*rand;
end;

gplot(AdjMatrixSmall,coordinates,'k-*');

%% Check the amount of links originating from each webpage
NumLinks=sum(AdjMatrixSmall,2);

%% Create a matrix of probabilities (Google matrix)
% Element (i,j) of the matrix shows the probability of moving from i-th
% page of the network to jth page. It is assumed that the user can follow
% any link on the page with a total probability of 85% (all hyperlinks are
% equal), and jump (teleport) to any other page in the network with a total
% probability of 15% (again, all pages are equal).

alpha=0.15;
GoogleMatrix=zeros(NumNetwork,NumNetwork);
for i=1:NumNetwork
    if NumLinks(i)~=0
        GoogleMatrix(i,:)=AdjMatrixSmall(i,:)./NumLinks(i);
    else
        GoogleMatrix(i,:)=1./NumNetwork;
    end;
end;

GoogleMatrix=(1-alpha)*GoogleMatrix+alpha*ones(NumNetwork,NumNetwork)./NumNetwork;

%% Check that all the rows in the GoogleMatrix matrix sum to 1
SumGoogleMatrix=sum(GoogleMatrix,2);

%% Finding an eigenvector corresponding to 1 (why is there such an eigenvector)?
w0=ones(1,NumNetwork)./sqrt(NumNetwork);

w1=w0*GoogleMatrix;
w2=w1*GoogleMatrix;
w3=w2*GoogleMatrix;

w100=w0*(GoogleMatrix)^100;
w90=w0*(GoogleMatrix)^90;
```

```

% Check the difference between v30 and v20. Observe that it is pretty
% small
deltaw=w100-w90;

%% Compute the eigenvalues and the right eigenvectors
[VectRight,LamRight]=eig(GoogleMatrix);
% Explain the result
LamRight=diag(LamRight);

%% Compute the eigenvalues and the left eigenvectors
[VectLeft,LamLeft]=eig(GoogleMatrix');
LamLeft=diag(LamLeft);

%% Separate the eigenvector corresponding to the eigenvalue 1 and scale it
u1=VectLeft(:,1);

u1=abs(u1)/norm(u1);

%% Select the maximum element and the corresponding element.
%Which page is the most important in the network?

[MaxRank,PageMaxRank]=max(u1);

%% Check if it's the most popular (most linked to page):

MostLinks=sum(AdjMatrixSmall);
[MaxLinks,PageMaxLinks]=max(MostLinks);

```

### Printout of the file lab13.m

```
% Facebook and graph partitioning

%% Simple graph

%% Load the data
clear;
load('social.mat','Social');
spy(Social);

%% Laplacian

%% Spectral decomposition

%% Clusteer in two groups

%% Cluster in 4 groups

Diagonal(indsmall)=large+1000;
[small,indsmall]=min(Diagonal);

V3=V(:,indsmall);

pp=[];
pn=[];
np=[];
nn=[];

for j=1:m
    if (V2(j)>0)
        if (V3(j)>0)
            pp=[pp,j];
        else
            pn=[pn,j];
        end;
    else
        if (V3(j)>0)
            np=[np,j];
        else
            nn=[nn,j];
        end;
    end;
end;

order=[pp,pn,np,nn];

iden=eye(m,m);

for j=1:351;
    for k=1:351;
        P(j,k)=iden(order(j),k);
    end;
end;
```

```

SocialOrdered=P*Social*P'

figure;
spy(SocialOrdered)

%% Second order of Fiedler

SocialPos=Social(pos,pos);
SocialNeg=Social(neg,neg);

rowsumpos=sum(SocialPos,2);

DiagSocialPos=diag(rowsumpos);

LaplaceSocialPos=DiagSocialPos-SocialPos;

[VPos,DPos]=eig(LaplaceSocialPos);

V2Pos=VPos(:,2);

[mpos,npos]=size(V2Pos);

posp=[];
posn=[];

for j=1:mpos
    if (V2Pos(j)>0)
        posp=[posp,pos(j)];
    else
        posn=[posn,pos(j)];
    end;
end;

rowsumneg=sum(SocialNeg,2);

DiagSocialNeg=diag(rowsumneg);

LaplaceSocialNeg=DiagSocialNeg-SocialNeg;

[VNeg,DNeg]=eig(LaplaceSocialNeg);

V2Neg=VNeg(:,2);

[mneg,nneg]=size(V2Neg);

negp=[];
negn=[];

for j=1:mneg
    if (V2Neg(j)>0)
        negp=[negp,neg(j)];
    else
        negn=[negn,neg(j)];
    end;
end;

```

```

        end;
    end;

    ordergen=[posp,posn,negp,negn];

    iden=eye(m,m);

    for j=1:351;
        for k=1:351;
            P(j,k)=iden(ordergen(j),k);
        end;
    end;

    SocialOrderedGen=P*Social*P'
    figure;
    spy(SocialOrderedGen)

```

## Printout of the file lab14.m

```

%% 2x2 matrix

clear;
t=linspace(0,2*pi,100);
X=[cos(t);sin(t)];
subplot(2,2,1);
hold on;
plot(X(1,:),X(2:,:), 'b');
quiver(0,0,1,0,0, 'r');
quiver(0,0,0,1,0, 'g');
axis equal
title('Unit circle')
hold off;

A = [ 2, 1; -1, 1 ];
[U,S,V] = svd(A);
U'*U
V'*V

VX=V'*X;
subplot(2,2,2)
hold on;
plot(VX(1,:),VX(2:,:), 'b');
quiver(0,0,V(1,1),V(1,2),0, 'r');
quiver(0,0,V(2,1),V(2,2),0, 'g');
axis equal
title('Multiplied by matrix V^T')
hold off;

SVX = S*VX;
subplot(2,2,3);
hold on;
plot(SVX(1,:),SVX(2:,:), 'b');
quiver(0,0,S(1,1)*V(1,1),S(2,2)*V(1,2),0, 'r');
quiver(0,0,S(1,1)*V(2,1),S(2,2)*V(2,2),0, 'g');
axis equal
title('Multiplied by matrix \Sigma V^T')
hold off;

AX = U*SVX;
subplot(2,2,4)
hold on;
plot(AX(1,:),AX(2:,:), 'b');
quiver(0,0,U(1,1)*S(1,1)*V(1,1)+U(1,2)*S(2,2)*V(1,2),U(2,1)*S(1,1)*V(1,1)+...
    U(2,2)*S(2,2)*V(1,2),0, 'r');
quiver(0,0,U(1,1)*S(1,1)*V(2,1)+U(1,2)*S(2,2)*V(2,2),U(2,1)*S(1,1)*V(2,1)+...
    U(2,2)*S(2,2)*V(2,2),0, 'g');
axis equal
title('Multiplied by matrix U\Sigma V^T=A')
hold off;

%% Modified SVD

```



```

U1(:,1)=U(:,1);
U1(:,2)=-U(:,2);
V1(:,1)=V(:,1);
V1(:,2)=-V(:,2);

U1*S*V1'-A

%% Check
A*V-U*S

%% Image compression

% Creates a two-dimensional array with the dimensions equal to the dimensions of
% the image
clear;
ImJPG=imread('einstein.jpg');
figure;
imshow(ImJPG);

[m,n]=size(ImJPG);

% Compute an SVD

[UIm,SIm,VIm]=svd(double(ImJPG));

% plot the singular values
figure;
plot(1:min(m,n),diag(SIm));

%% Create approximations to the image

% With 50, 100, and 150 singular values
for k=50:50:150
    ImJPG_comp=uint8(UIm(:,1:k)*SIm(1:k,1:k)*(VIm(:,1:k))');
    figure;
    imshow(ImJPG_comp)
    % compression percentage
    pct = 1 - (numel(UIm(:,1:k))+numel(VIm(:,1:k)*SIm(1:k,1:k)))/numel(ImJPG);
    fprintf('Compression percentage for %2.0f singular values: %8.3f\n',k, pct);
end;

%% Noise filtering
clear;
ImJPG=imread('checkers.pgm')
[m,n]=size(ImJPG);

% Add some noise to the image
ImJPG_Noisy=double(ImJPG)+50*(rand(m,n)-0.5*ones(m,n));
figure;
imshow(ImJPG);

figure;
imshow(uint8(ImJPG_Noisy));

```

```

[UIm,SIm,VIm]=svd(ImJPG_Noisy);

figure;
plot(1:min(m,n),diag(SIm),'ko');

for k=10:20:50
    ImJPG_comp=uint8(UIm(:,1:k)*SIm(1:k,1:k)*(VIm(:,1:k))');
    figure;
    imshow(ImJPG_comp)
    % compression percentage
    pct = 1 - (numel(UIm(:,1:k))+numel(VIm(:,1:k)*SIm(1:k,1:k)))/numel(ImJPG);
    fprintf('Compression percentage for %2.0f singular values: %8.3f\n',k, pct);
end;

```

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