

MATH 2740 – 06 Matrix methods

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The University of Manitoba campuses are located on original lands of Anishinaabeg, Ininew, Anisininew, Dakota and Dene peoples, and on the National Homeland of the Red River Métis.

We respect the Treaties that were made on these territories, we acknowledge the harms and mistakes of the past, and we dedicate ourselves to move forward in partnership with Indigenous communities in a spirit of Reconciliation and collaboration.

Outline

Least squares problems

QR factorisation

Singular values decomposition (SVD)

Principal component analysis (PCA)

Support vector machines





Grabing the Canadian census data

We want to consider the evolution of the population of Canada through time

For this, we grab the Canadian census data

Search for (Google) "Canada historical census data csv", since csv (comma separated values) is a very easy format to use with ${\sf R}$

Here, we find a csv for 1851 to 1976

We follow the link to Table A2-14, where we find another link, this time to a csv file. This is what we use in R

Grabing the Canadian census data

1

The function read.csv reads in a file (potentially directly from the web) Assign the result to the variable data. We then use the function head to show the first few lines in the result.

```
data_old = read.csv("https://www150.statcan.gc.ca/n1/en/pub/11-516-x/secti
head(data_old)
##
     X Series.A2.14.
```

1 NA Year ## 2 NA

3 NA ## 4 NA

5 NA

##

6 NA

Population.of.Canada..by.province..census.dates..1851.to.1976 X.1

NA

Canada NA New - __Least-squares problems

Obviously, this does not make a lot of sense. This is normal: take a look at the first

```
few lines in the file. They take the form
head(data_old)
       X Series. A2.14.
##
## 1 NA
## 2 NA
                   Year
## 3 NA
## 4 NA
```

6 NA Population.of.Canada..by.province..census.dates..1851.to.1976 X.1

5 NA

NA New NA NA

NA

NA NA

Canada

2 ## 3 ## 4

5 ## 6 - Least squares problems

##

1

The first line here does this; it is easy to deal with this: the function read.csv takes the optional argument skip=, which indicates how many lines to skip at the beginning The second line is also empty, so let us skip it too

data_old = read.csv("https://www150.statcan.gc.ca/n1/en/pub/11-516-x/secti

skip = 2)

NA

NΑ

5 NA 1976 22,992,604

3 - Least squares problems

```
head(data_old)
                                                                            Que
##
      X Year
                  Canada X.1 Newfound. Prince X.2
                                                        Nova
                                                                    New
## 1 NA
                          NA
                                   land
                                         Edward
                                                  NA
                                                      Scotia Brunswick
                                         Island
## 2 NA
                          NA
                                                  NA
                                                  NA
## 3 NA
                          NΑ
                                                            5
                                                                      6
##
  4 NA
                          NA
                                                  NA
```

6 NA NANA Manitoba X.3 Saskat. X.4 Alberta X.5 British X.6 Y. ## Ontario Columbia NA Terri ## 1 NA chewan NA NA ## 2 NA NA NA NA

10

557,725 118,229

NΑ

NA 828,571

NA

11

677,250 6,234,

NΑ

12

Here, there is the further issue that to make things legible, the table authors used 3 rows (from 2 to 4) to encode for long names (e.g., Prince Edward Island is written over 3 rows). Note, however, that 'read.csv' has rightly picked up on the first row being the column names.

(You could also use the function 'read_csv' from the package 'readr' to read in the file. This function is a bit more flexible than 'read.csv' and can handle such cases more easily. However, it is not part of the base R package, so you would need to install it first.)

Because we are only interested in the total population of the country and the year, let us simply get rid of the first 4 rows and of all columns except the second (Year) and third (Canada)

```
data_old = data_old[5:dim(data_old)[1], 2:3]
head(data_old, n=4)

## Year Canada
## 5 1976 22,992,604
## 6

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## 7 1971 21,568,311
```

Still not perfect:

- there are some empty rows; - the last few rows need to be removed too, they contain remarks about the data; - the population counts contain commas; - it would be better if years were increasing.

Let us fix these issues.

For 1 and 2, this is easy: remark that the Canada column is empty for both issues.

Now remark as well that below Canada (and Year, for that matter), it is written <chr>. This means that entries in the column are characters. Looking for empty content therefore means looking for empty character chains.

So to fix 1 and 2, we keep the rows where Canada does not equal the empty chain.

To get rid of commas, we just need to substitute an empty chain for ",".

To sort, we find the order for the years and apply it to the entire table.

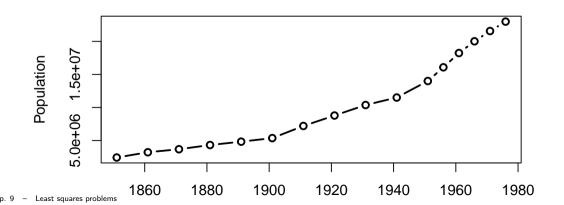
Finally, as remarked above, for now, both the year and the population are considered as character chains. This means that in order to plot anything, we will have to indicate that these are numbers. not characters.

```
data_old = data_old[which(data_old$Canada != ""),]
data_old$Canada = gsub(",", "", data_old$Canada)
order_data = order(data_old$Year)
data old = data old[order data.]
data_old$Year = as.numeric(data_old$Year)
data_old$Canada = as.numeric(data_old$Canada)
data old
##
      Year
             Canada
## 23 1851
            2436297
## 22 1861
            3229633
## 21 1871
            3689257
## 20 1881
            4324810
            4833239
## 19 1891
## 17 1901
            5371315
           7206643
## 16 1911
## 15 1921
            8787949
      1931 10376786
```

Row numbers are a little weird, so let us fix this.

```
row.names(data old) = 1:dim(data old)[1]
data_old
##
      Year
             Canada
## 1
      1851
            2436297
## 2
      1861
            3229633
## 3
      1871
            3689257
## 4
      1881
            4324810
## 5
      1891
            4833239
            5371315
## 6
      1901
## 7
      1911
            7206643
      1921
            8787949
## 8
      1931 10376786
## 9
  10 1941 11506655
  11 1951 14009429
## 12 1956 16080791
```

```
plot(data_old$Year, data_old$Canada,
    type = "b", lwd = 2,
    xlab = "Year", ylab = "Population")
```



download csv link in there, let us see where this leads us. The table is 720KB, so surely there must be more to this than just the population. To get a sense of that, we dump the whole data.frame, not just its head.

But wait, this is only to 1976..! Looking around, we find another table here. There's a

```
data_new = read.csv("https://www12.statcan.gc.ca/census-recensement/2011/d
head(data_new, 10)
      GEOGRAPHY.NAME
                                CHARACTERISTIC YEAR.S. TOTAL FLAG TOTAL
##
```

Canada Population (in thousands) 1956 16081 Canada Population (in thousands) 1961 18238 ## 2 ## 3 Canada Population (in thousands) 1966 20015

Canada Population (in thousands) ## 4 1971 21568 Canada Population (in thousands) ## 5 1976 22993

6 Canada Population (in thousands) 1981 24343

Canada Population (in thousands) ## 7 1986 25309

Canada Population (in thousands) ## 8 1991 27297

Canada Population (in thousands) ## 9 1996 28847 Canada Population (in thousands) 2001 30007

1

Haha, this looks quite nice but has way more information than we need: we just want the population of Canada and here we get 9960 rows. Also, the population of Canada is expressed in thousands, so once we selected what we want, we will need to multiply by 1,000.

There are many ways to select rows. Let us proceed as follows: we want the rows where the geography is "Canada" and the characteristic is "Population (in thousands)". Let us find those indices of rows that satisfy the first criterion, those that satisfy the second; if we then intersect these two sets of indices, we will have selected the rows we want.

```
idx_CAN = which(data_new$GEOGRAPHY.NAME == "Canada")
idx_char = which(data_new$CHARACTERISTIC == "Population (in thousands)")
idx_keep = intersect(idx_CAN, idx_char)
head(idx_keep, n = 8)
## [1] 1 2 3 4 5 6 7 8
```

Yes, this looks okay, so let us keep only these

```
data_new = data_new[idx_keep,]
head(data new. n = 8)
##
    GEOGRAPHY NAME
                              CHARACTERISTIC YEAR.S. TOTAL FLAG TOTAL
## 1
            Canada Population (in thousands) 1956 16081
## 2
            Canada Population (in thousands) 1961 18238
## 3
            Canada Population (in thousands) 1966 20015
## 4
            Canada Population (in thousands) 1971 21568
## 5
            Canada Population (in thousands) 1976 22993
## 6
            Canada Population (in thousands) 1981 24343
## 7
            Canada Population (in thousands) 1986 25309
            Canada Population (in thousands) 1991 27297
## 8
```

We want to concatenate this data.frame with the one from earlier

To do this, we need the two data frames to have the same number of columns and, actually, the same column names and entry types (notice that YEAR.S. in data_new is a column of characters)

What remains to do

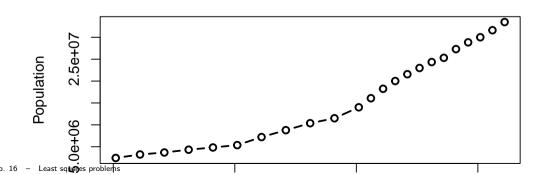
- ▶ Rename the columns in the pruned old data (data_pruned) to year and population. Personally, I prefer lowercase column names.. and population is more informative than Canada
- ► Keep only the relevant columns in data_new, rename them accordingly and multiply population by 1,000 there
- Transform year in data_new to numbers
- ▶ We already have data up to and including 1976 in data_old, so get rid of that in data_new
- Append the rows of data_new to those of data_pruned

```
colnames(data_old) = c("year", "population")
data_new = data_new[,c("YEAR.S.","TOTAL")]
colnames(data_new) = c("year", "population")
data_new$year = as.numeric(data_new$year)
data_new = data_new[which(data_new$year>1976),]
data_new$population = data_new$population*1000

data = rbind(data_old.data_new)
```

Let us plot the result

```
plot(data$year, data$population,
    type = "b", lwd = 2,
    xlab = "Year", ylab = "Population")
```



Save the processed data

In case we need the data elsewhere, we save the data to a csv file

```
write.csv(data, file = "../CODE/Canada_census.csv")
```

Using readr saves the data without row numbers (by default), so we can do this instead

```
readr::write_csv(data, file = "../CODE/Canada_census.csv")
```



The least squares problem (simplest version)

Definition 1

Given a collection of points $(x_1, y_1), \ldots, (x_n, y_n)$, find the coefficients a, b of the line y = a + bx such that

$$\|\mathbf{e}\| = \sqrt{\varepsilon_1^2 + \dots + \varepsilon_n^2} = \sqrt{(y_1 - \tilde{y}_1)^2 + \dots + (y_n - \tilde{y}_n)^2}$$

is minimal, where $\tilde{y}_i = a + bx_i$ for $i = 1, \dots, n$

We just saw how to solve this by brute force using a genetic algorith to minimise ||e||, let us now see how to solve this problem "properly"

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For a data point $i = 1, \ldots, n$

$$\varepsilon_i = y_i - \tilde{y}_i = y_i - (a + bx_i)$$

So if we write this for all data points,

$$arepsilon_1 = y_1 - (a + bx_1)$$
 $dots$
 $arepsilon_p = y_p - (a + bx_p)$

In matrix form

$$oldsymbol{e} = oldsymbol{b} - Aoldsymbol{x}$$

with

$$m{e} = \begin{pmatrix} arepsilon_1 \\ dots \\ arepsilon_n \end{pmatrix}, A = \begin{pmatrix} 1 & x_1 \\ dots & dots \\ 1 & x_n \end{pmatrix}, m{x} = \begin{pmatrix} a \\ b \end{pmatrix} \text{ and } m{b} = \begin{pmatrix} y_1 \\ dots \\ y_n \end{pmatrix}$$

The least squares problem (reformulated)

Definition 2 (Least squares solutions)

Consider a collection of points $(x_1, y_1), \ldots, (x_n, y_n)$, a matrix $A \in \mathcal{M}_{mn}$, $\mathbf{b} \in \mathbb{R}^m$. A least squares solution of $A\mathbf{x} = \mathbf{b}$ is a vector $\tilde{\mathbf{x}} \in \mathbb{R}^n$ s.t.

$$\forall \mathbf{x} \in \mathbb{R}^n, \quad \|\mathbf{b} - A\tilde{\mathbf{x}}\| \leq \|\mathbf{b} - A\mathbf{x}\|$$

Needed to solve the problem

Definition 3 (Best approximation)

Let V be a vector space, $W \subset V$ and $\mathbf{v} \in V$. The **best approximation** to \mathbf{v} in W is $\tilde{\mathbf{v}} \in W$ s.t.

$$\forall \mathbf{w} \in W, \mathbf{w} \neq \tilde{\mathbf{v}}, \quad \|\mathbf{v} - \tilde{\mathbf{v}}\| < \|\mathbf{v} - \mathbf{w}\|$$

Theorem 4 (Best approximation theorem)

Let V be a vector space with an inner product, $W \subset V$ and $\mathbf{v} \in V$. Then $\operatorname{proj}_W(\mathbf{v})$ is the best approximation to \mathbf{v} in W

Let us find the least squares solution

 $\forall x \mathbb{R}^n$, Ax is a vector in the **column space** of A (the space spanned by the vectors making up the columns of A)

Since
$$\mathbf{x} \in \mathbb{R}^n$$
, $A\mathbf{x} \in \text{col}(A)$

 \implies least squares solution of Ax = b is a vector $\tilde{y} \in col(A)$ s.t.

$$\forall \mathbf{y} \in \mathsf{col}(A), \quad \|\mathbf{b} - \tilde{\mathbf{y}}\| \leq \|\mathbf{b} - \mathbf{y}\|$$

This looks very much like Best approximation and Best approximation theorem

Putting things together

We just stated: The least squares solution of Ax = b is a vector $\tilde{y} \in col(A)$ s.t.

$$\forall \mathbf{y} \in \operatorname{col}(A), \quad \|\mathbf{b} - \tilde{\mathbf{y}}\| \leq \|\mathbf{b} - \mathbf{y}\|$$

We know (reformulating a tad):

Theorem 5 (Best approximation theorem)

Let V be a vector space with an inner product, $W \subset V$ and $\mathbf{v} \in V$. Then $\operatorname{proj}_W(\mathbf{v}) \in W$ is the best approximation to \mathbf{v} in W, i.e.,

$$\forall \mathbf{w} \in W, \mathbf{w} \neq \operatorname{proj}_W(\mathbf{v}), \quad \|\mathbf{v} - \operatorname{proj}_W(\mathbf{v})\| < \|\mathbf{v} - \mathbf{w}\|$$

$$\implies W = \operatorname{col}(A), \ \boldsymbol{v} = \boldsymbol{b} \text{ and } \tilde{\boldsymbol{y}} = \operatorname{proj}_{\operatorname{col}(A)}(\mathbf{b})$$

So if \tilde{x} is a least squares solution of Ax = b, then

$$\tilde{\pmb{y}} = A\tilde{\pmb{x}} = \mathsf{proj}_{\mathsf{col}(A)}(\mathbf{b})$$

We have

$$\boldsymbol{b} - A\tilde{\boldsymbol{x}} = \boldsymbol{b} - \operatorname{proj}_{\operatorname{col}(A)}(\mathbf{b}) = \operatorname{perp}_{\operatorname{col}(A)}(\mathbf{b})$$

and it is easy to show that

$$\mathsf{perp}_{\mathsf{col}(A)}(\mathbf{b}) \perp \mathsf{col}(A)$$

So for all columns \mathbf{a}_i of A

$$\mathbf{a}_i \cdot (\mathbf{b} - A\tilde{\mathbf{x}}) = 0$$

which we can also write as $\mathbf{a}_{i}^{T}(\mathbf{b} - A\tilde{\mathbf{x}}) = 0$

For all columns a_i of A,

$$\boldsymbol{a}_{i}^{T}(\boldsymbol{b}-A\tilde{\boldsymbol{x}})=0$$

This is equivalent to saying that

$$A^{T}(\boldsymbol{b} - A\tilde{\boldsymbol{x}}) = \boldsymbol{0}$$

We have

$$A^{T}(\boldsymbol{b} - A\tilde{\boldsymbol{x}}) = \boldsymbol{0} \iff A^{T}\boldsymbol{b} - A^{T}A\tilde{\boldsymbol{x}} = \boldsymbol{0}$$
$$\iff A^{T}\boldsymbol{b} = A^{T}A\tilde{\boldsymbol{x}}$$
$$\iff A^{T}A\tilde{\boldsymbol{x}} = A^{T}\boldsymbol{b}$$

The latter system constitutes the **normal equations** for \tilde{x}

Least squares theorem

Theorem 6 (Least squares theorem)

 $A \in \mathcal{M}_{mn}$, $\boldsymbol{b} \in \mathbb{R}^m$. Then

- 1. Ax = b always has at least one least squares solution \tilde{x}
- 2. $\tilde{\mathbf{x}}$ least squares solution to $A\mathbf{x} = \mathbf{b} \iff \tilde{\mathbf{x}}$ is a solution to the normal equations $A^T A \tilde{\mathbf{x}} = A^T \mathbf{b}$
- 3. A has linearly independent columns \iff A^TA invertible. In this case, the least squares solution is unique and

$$\tilde{\mathbf{x}} = \left(A^T A\right)^{-1} A^T \mathbf{b}$$

We have seen 1 and 2, we will not show 3 (it is not hard)



Suppose we want to fit something a bit more complicated..

For instance, instead of the affine function

$$y = a + bx$$

suppose we want to do the quadratic

$$y = a_0 + a_1 x + a_2 x^2$$

or even

$$y=k_0e^{k_1x}$$

How do we proceed?

Fitting the quadratic

We have the data points $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$ and want to fit

$$y = a_0 + a_1 x + a_2 x^2$$

At (x_1, y_1) ,

$$\tilde{y}_1 = a_0 + a_1 x_1 + a_2 x_1^2$$

:

At (x_n, y_n) ,

$$\tilde{y}_n = a_0 + a_1 x_n + a_2 x_n^2$$

In terms of the error

$$\varepsilon_1 = y_1 - \tilde{y}_1 = y_1 - (a_0 + a_1 x_1 + a_2 x_1^2)$$

 \vdots
 $\varepsilon_n = y_n - \tilde{y}_n = y_n - (a_0 + a_1 x_n + a_2 x_n^2)$

i.e.,

$$e = b - Ax$$

where

$$\boldsymbol{e} = \begin{pmatrix} \varepsilon_1 \\ \vdots \\ \varepsilon_n \end{pmatrix}, A = \begin{pmatrix} 1 & x_1 & x_1^2 \\ \vdots & \vdots & \vdots \\ 1 & x_n & x_n^2 \end{pmatrix}, \boldsymbol{x} = \begin{pmatrix} a_0 \\ a_1 \\ a_2 \end{pmatrix} \text{ and } \boldsymbol{b} = \begin{pmatrix} y_1 \\ \vdots \\ y_n \end{pmatrix}$$

Theorem 6 applies, with here $A \in \mathcal{M}_{n3}$ and $\boldsymbol{b} \in \mathbb{R}^n$

Fitting the exponential

Things are a bit more complicated here

If we proceed as before, we get the system

$$y_1 = k_0 e^{k_1 x_1}$$

$$\vdots$$

$$y_n = k_0 e^{k_1 x_n}$$

 $e^{k_1x_i}$ is a nonlinear term, it cannot be put in a matrix

However: take the In of both sides of the equation

$$\ln(y_i) = \ln(k_0 e^{k_1 x_i}) = \ln(k_0) + \ln(e^{k_1 x_i}) = \ln(k_0) + k_1 x_i$$

If $y_i, k_0 > 0$, then their In are defined and we're in business...

$$ln(y_i) = ln(k_0) + k_1 x_i$$

So the system is

$$\mathbf{y} = A\mathbf{x} + \mathbf{b}$$

with

$$A = \begin{pmatrix} x_1 \\ \vdots \\ x_n \end{pmatrix}, \boldsymbol{x} = (k_1), \boldsymbol{b} = (\ln(k_0)) \text{ and } \boldsymbol{y} = \begin{pmatrix} \ln(y_1) \\ \vdots \\ \ln(y_n) \end{pmatrix}$$





Matrix factorisations

Matrix factorisations are popular because they allow to perform some computations more easily

There are several different types of factorisations. Here, we study just the QR factorisation, which is useful for many least squares problems



Definition 7 (Orthogonal set of vectors)

The set of vectors $\{\mathbf{v}_1, \dots, \mathbf{v}_k\} \in \mathbb{R}^n$ is an **orthogonal set** if

$$\forall i, j = 1, \dots, k, \quad i \neq j \implies \mathbf{v}_i \bullet \mathbf{v}_j = 0$$

Theorem 8

 $\{\mathbf{v}_1,\ldots,\mathbf{v}_k\}\in\mathbb{R}^n$ with $\forall i,\ \mathbf{v}_i\neq\mathbf{0}$, orthogonal set $\implies\{\mathbf{v}_1,\ldots,\mathbf{v}_k\}\in\mathbb{R}^n$ linearly independent

Definition 9 (Orthogonal basis)

Let S be a basis of the subspace $W \subset \mathbb{R}^n$ composed of an orthogonal set of vectors. We say S is an **orthogonal basis** of W

Proof of Theorem 8

Assume $\{v_1,\ldots,v_k\}$ orthogonal set with $v_i\neq \mathbf{0}$ for all $i=1,\ldots,k$. Recall $\{v_1,\ldots,v_k\}$ is LI if

$$c_1 \mathbf{v}_1 + \cdots + c_k \mathbf{v}_k = \mathbf{0} \iff c_1 = \cdots = c_k = \mathbf{0}$$

So assume $c_1, \ldots, c_k \in \mathbb{R}$ are s.t. $c_1 \mathbf{v}_1 + \cdots + c_k \mathbf{v}_k = \mathbf{0}$. Recall that $\forall \mathbf{x} \in \mathbb{R}^k$, $\mathbf{0}_k \bullet \mathbf{x} = \mathbf{0}$. So for some $\mathbf{v}_i \in \{\mathbf{v}_1, \ldots, \mathbf{v}_k\}$

$$0 = \mathbf{0} \bullet \mathbf{v}_{i}$$

$$= (c_{1}\mathbf{v}_{1} + \dots + c_{k}\mathbf{v}_{k}) \bullet \mathbf{v}_{i}$$

$$= c_{1}\mathbf{v}_{1} \bullet \mathbf{v}_{i} + \dots + c_{k}\mathbf{v}_{k} \bullet \mathbf{v}_{i}$$

$$(1)$$

As $\{\mathbf{v}_1,\ldots,\mathbf{v}_k\}$ orthogonal, $\mathbf{v}_i \bullet \mathbf{v}_i = 0$ when $i \neq j$, (1) reduces to

$$c_i \mathbf{v}_i \bullet \mathbf{v}_i = 0 \iff c_i ||\mathbf{v}_i||^2 = 0$$

As $\mathbf{v}_i \neq 0$ for all i, $\|\mathbf{v}_i\| \neq 0$ and so $c_i = 0$. This is true for all i, hence the result

Example – Vectors of the standard basis of \mathbb{R}^3

For \mathbb{R}^3 , we denote

$$m{i} = egin{pmatrix} 1 \\ 0 \\ 0 \end{pmatrix}, \quad m{j} = egin{pmatrix} 0 \\ 1 \\ 0 \end{pmatrix} \ ext{and} \ m{k} = egin{pmatrix} 0 \\ 0 \\ 1 \end{pmatrix}$$

 $(\mathbb{R}^k \text{ for } k > 3, \text{ we denote them } \boldsymbol{e}_i)$

Clearly, $\{i,j\}$, $\{i,k\}$, $\{j,k\}$ and $\{i,j,k\}$ orthogonal sets. The standard basis vectors are also $\neq \mathbf{0}$, so the sets are LI. And

$$\{i,j,k\}$$

is an orthogonal basis of \mathbb{R}^3 since it spans \mathbb{R}^3 and is LI

$$c_1 oldsymbol{i} + c_2 oldsymbol{j} + c_3 oldsymbol{k} = c_1 egin{pmatrix} 1 \ 0 \ 0 \end{pmatrix} + c_2 egin{pmatrix} 0 \ 1 \ 0 \end{pmatrix} + c_3 egin{pmatrix} 0 \ 0 \ 1 \end{pmatrix} = egin{pmatrix} c_1 \ c_2 \ c_3 \end{pmatrix}$$

Orthonormal version of things

Definition 10 (Orthonormal set)

The set of vectors $\{v_1, \dots, v_k\} \in \mathbb{R}^n$ is an **orthonormal set** if it is an orthogonal set and furthermore

$$\forall i=1,\ldots,k, \quad \|\mathbf{v}_i\|=1$$

Definition 11 (Orthonormal basis)

A basis of the subspace $W\subset\mathbb{R}^n$ is an **orthonormal basis** if the vectors composing it are an orthonormal set

 $\{\mathbf{v}_1,\ldots,\mathbf{v}_k\}\in\mathbb{R}^n$ is orthonormal if

$$\mathbf{v}_i \bullet \mathbf{v}_j = \begin{cases} 1 & \text{if } i = j \\ 0 & \text{otherwise} \end{cases}$$

Projections

Definition 12 (Orthogonal projection onto a subspace)

 $W \subset \mathbb{R}^n$ a subspace and $\{u_1, \dots, u_k\}$ an orthogonal basis of W. $\forall v \in \mathbb{R}^n$, the **orthogonal projection** of v **onto** W is

$$\operatorname{proj}_{\mathcal{W}}(\mathbf{v}) = rac{\mathbf{u}_1 ullet \mathbf{v}}{\|\mathbf{u}_1\|^2} \mathbf{u}_1 + \dots + rac{\mathbf{u}_k ullet \mathbf{v}}{\|\mathbf{u}_k\|^2} \mathbf{u}_k$$

Definition 13 (Component orthogonal to a subspace)

 $W \subset \mathbb{R}^n$ a subspace and $\{u_1, \ldots, u_k\}$ an orthogonal basis of W. $\forall v \in \mathbb{R}^n$, the component of v orthogonal to W is

$$\mathsf{perp}_W(oldsymbol{v}) = oldsymbol{v} - \mathsf{proj}_W(oldsymbol{v})$$

What this aims to do is to construct an orthogonal basis for a subspace $W\subset \mathbb{R}^n$

To do this, we use the Gram-Schmidt orthogonalisation process, which turn s a basis of W into an orthogonal basis of W

Gram-Schmidt process

Theorem 14

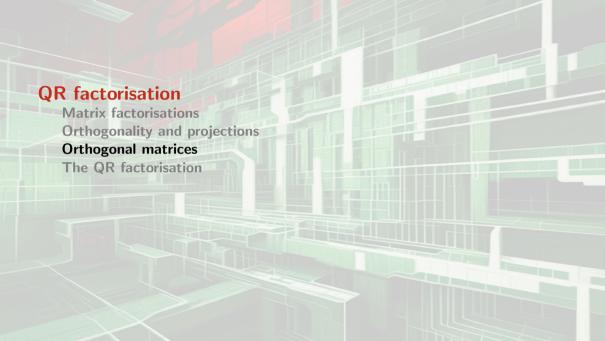
$$W \subset \mathbb{R}^n$$
 a subset and $\{x_1, \dots, x_k\}$ a basis of W . Let

$$\mathbf{v}_1 = \mathbf{x}_1$$
 $\mathbf{v}_2 = \mathbf{x}_2 - \frac{\mathbf{v}_1 \bullet \mathbf{x}_2}{\|\mathbf{v}_1\|^2} \mathbf{v}_1$
 $\mathbf{v}_3 = \mathbf{x}_3 - \frac{\mathbf{v}_1 \bullet \mathbf{x}_3}{\|\mathbf{v}_1\|^2} \mathbf{v}_1 - \frac{\mathbf{v}_2 \bullet \mathbf{x}_3}{\|\mathbf{v}_2\|^2} \mathbf{v}_2$
 \vdots

 $W_1 = \mathsf{span}(extbf{ extit{x}}_1), \, W_2 = \mathsf{span}(extbf{ extit{x}}_1, extbf{ extit{x}}_2), \ldots, \, W_k = \mathsf{span}(extbf{ extit{x}}_1, \ldots, extbf{ extit{x}}_k)$

 $\mathbf{v}_k = \mathbf{x}_k - \frac{\mathbf{v}_1 \bullet \mathbf{x}_k}{\|\mathbf{v}_k\|^2} \mathbf{v}_1 - \dots - \frac{\mathbf{v}_{k-1} \bullet \mathbf{x}_k}{\|\mathbf{v}_k\|^2} \mathbf{v}_{k-1}$

Then $\forall i = 1, ..., k$, $\{\mathbf{v}_1, ..., \mathbf{v}_i\}$ orthogonal basis for W_i



Theorem 15

Let $Q \in \mathcal{M}_{mn}$. The columns of Q form an orthonormal set if and only if

$$Q^TQ=\mathbb{I}_n$$

Definition 16 (Orthogonal matrix)

 $Q \in \mathcal{M}_n$ is an **orthogonal matrix** if its columns form an orthonormal set

So
$$Q \in \mathcal{M}_n$$
 orthogonal if $Q^TQ = \mathbb{I}$, i.e., $Q^T = Q^{-1}$

Theorem 17 (NSC for orthogonality)

$$Q \in \mathcal{M}_n$$
 orthogonal $\iff Q^{-1} = Q^T$

Theorem 18 (Orthogonal matrices "encode" isometries)

Let $Q \in \mathcal{M}_n$. TFAE

- 1. Q orthogonal
- 2. $\forall \mathbf{x} \in \mathbb{R}^n$, $||Q\mathbf{x}|| = ||\mathbf{x}||$
- 3. $\forall x, y \in \mathbb{R}^n$, $Qx \bullet Qy = x \bullet y$

Theorem 19

Let $Q \in \mathcal{M}_n$ be orthogonal. Then

- 1. The rows of Q form an orthonormal set
- 2. Q^{-1} orthogonal
- 3. $\det Q = \pm 1$
- 4. $\forall \lambda \in \sigma(Q), |\lambda| = 1$
- 5. If $Q_2 \in \mathcal{M}_n$ also orthogonal, then QQ_2 orthogonal

Proof of 4 in Theorem 19

All statements in Theorem 19 are easy, but let's focus on 4

Let λ be an eigenvalue of $Q \in \mathcal{M}_n$ orthogonal, i.e., $\exists \mathbb{R}^n \ni \mathbf{x} \neq \mathbf{0}$ s.t.

$$Q\mathbf{x} = \lambda \mathbf{x}$$

Take the norm on both sides

$$\|Q\mathbf{x}\| = \|\lambda\mathbf{x}\|$$

From 2 in Theorem 18, $\|Qx\| = \|x\|$ and from the properties of norms, $\|\lambda x\| = |\lambda| \|x\|$, so we have

$$\|Q\mathbf{x}\| = \|\lambda\mathbf{x}\| \iff \|\mathbf{x}\| = |\lambda| \|\mathbf{x}\| \iff 1 = |\lambda|$$

(we can divide by ||x|| since $x \neq 0$ as an eigenvector)

The QR factorisation

Theorem 20

Let $A \in \mathcal{M}_{mn}$ with LI columns. Then A can be factored as

$$A = QR$$

where $Q \in \mathcal{M}_{mn}$ has orthonormal columns and $R \in \mathcal{M}_n$ is nonsingular upper triangular

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Back to least squares

So what was the point of all that..?

Theorem 21 (Least squares with QR factorisation)

 $A \in \mathcal{M}_{mn}$ with LI columns, $\mathbf{b} \in \mathbb{R}^m$. If A = QR is a QR factorisation of A, then the unique least squares solution $\tilde{\mathbf{x}}$ of $A\mathbf{x} = \mathbf{b}$ is

$$\tilde{\mathbf{x}} = R^{-1}Q^T\mathbf{b}$$

Proof of Theorem 21

A has LI columns so

- least squares $A\mathbf{x} = \mathbf{b}$ has unique solution $\tilde{\mathbf{x}} = (A^T A)^{-1} A^T \mathbf{b}$
- ▶ by Theorem 20, A can be written as A = QR with $Q \in \mathcal{M}_{mn}$ with orthonormal columns and $R \in \mathcal{M}_n$ nonsingular and upper triangular

So

$$A^{T}A\tilde{\mathbf{x}} = A^{T}\mathbf{b} \implies (QR)^{T}QR\tilde{\mathbf{x}} = (QR)^{T}\mathbf{b}$$

$$\implies R^{T}Q^{T}QR\tilde{\mathbf{x}} = R^{T}Q^{T}\mathbf{b}$$

$$\implies R^{T}\mathbb{I}_{n}R\tilde{\mathbf{x}} = R^{T}Q^{T}\mathbf{b}$$

$$\implies R^{T}R\tilde{\mathbf{x}} = R^{T}Q^{T}\mathbf{b}$$

$$\implies (R^{T})^{-1}R\tilde{\mathbf{x}} = (R^{T})^{-1}R^{T}Q^{T}\mathbf{b}$$

$$\implies R\tilde{\mathbf{x}} = Q^{T}\mathbf{b}$$

$$\implies \tilde{\mathbf{x}} = R^{-1}Q^{T}\mathbf{b} \quad \Box$$

Least squares problems **QR** factorisation Singular values decomposition (SVD) Principal component analysis (PCA) Support vector machines

Matrix factorisations (continued)

The singular value decomposition (known mostly by its acronym, SVD) is yet another type of factorisation/decomposition..

Singular values

Definition 22 (Singular value)

Let $A \in \mathcal{M}_{mn}(\mathbb{R})$. The **singular values** of A are the real numbers

$$\sigma_1 \geq \sigma_2 \geq \cdots \sigma_n \geq 0$$

that are the square roots of the eigenvalues of A^TA

Singular values are real and nonnegative?

Recall that $\forall A \in \mathcal{M}_{mn}$, $A^T A$ is symmetric

Claim 1. Real symmetric matrices have real eigenvalues

Proof. $A \in \mathcal{M}_n(\mathbb{R})$ symmetric and (λ, \mathbf{v}) eigenpair of A, i.e, $A\mathbf{v} = \lambda \mathbf{v}$. Taking the complex conjugate, $\overline{A\mathbf{v}} = \overline{\lambda} \overline{\mathbf{v}}$

Since
$$A \in \mathcal{M}_n(\mathbb{R})$$
, $\overline{A} = A$ $(z = \overline{z} \iff z \in \mathbb{R})$

So

$$A\bar{\mathbf{v}} = \overline{A}\bar{\mathbf{v}} = \overline{A\mathbf{v}} = \overline{\lambda}\bar{\mathbf{v}} = \overline{\lambda}\bar{\mathbf{v}}$$

i.e., if (λ, \mathbf{v}) eigenpair, $(\bar{\lambda}, \bar{\mathbf{v}})$ also eigenpair

Still assuming $A \in \mathcal{M}_n(\mathbb{R})$ symmetric and (λ, \mathbf{v}) eigenpair of A and using what we just proved (that $(\bar{\lambda}, \bar{\mathbf{v}})$ also eigenpair), take transposes

$$A\bar{\mathbf{v}} = \bar{\lambda}\bar{\mathbf{v}} \iff (A\bar{\mathbf{v}})^T = (\bar{\lambda}\bar{\mathbf{v}})^T$$

$$\iff \bar{\mathbf{v}}^T A^T = \bar{\lambda}\bar{\mathbf{v}}^T$$

$$\iff \bar{\mathbf{v}}^T A = \bar{\lambda}\bar{\mathbf{v}}^T \qquad [A \text{ symmetric}]$$

Let us now compute $\lambda(\bar{\mathbf{v}} \bullet \mathbf{v})$. We have

$$\lambda(\bar{\boldsymbol{v}} \bullet \boldsymbol{v}) = \lambda \bar{\boldsymbol{v}}^T \boldsymbol{v} = \bar{\boldsymbol{v}}^T (\lambda \boldsymbol{v})$$

$$= \bar{\boldsymbol{v}}^T (A \boldsymbol{v}) = (\bar{\boldsymbol{v}}^T A) \boldsymbol{v}$$

$$= (\bar{\lambda} \bar{\boldsymbol{v}}^T) \boldsymbol{v} = \bar{\lambda} (\bar{\boldsymbol{v}} \bullet \boldsymbol{v})$$

$$\iff (\lambda - \bar{\lambda}) (\bar{\boldsymbol{v}} \bullet \boldsymbol{v}) = 0$$

We have shown

$$(\lambda - \bar{\lambda})(\bar{\mathbf{v}} \bullet \mathbf{v}) = 0$$

Let

$$\mathbf{v} = \begin{pmatrix} a_1 + ib_1 \\ \vdots \\ a_n + ib_n \end{pmatrix}$$

Then

$$\vec{\mathbf{v}} = \begin{pmatrix} a_1 - ib_1 \\ \vdots \\ a_n - ib_n \end{pmatrix}$$

So

$$ar{m{v}} ullet m{v} = (a_1^2 + b_1^2) + \dots + (a_n^2 + b_n^2)$$

But \mathbf{v} eigenvector is $\neq \mathbf{0}$, so $\bar{\mathbf{v}} \bullet \mathbf{v} \neq \mathbf{0}$, so

$$(\lambda - \bar{\lambda})(\bar{\mathbf{v}} \bullet \mathbf{v}) = 0 \iff \lambda - \bar{\lambda} = 0 \iff \lambda = \bar{\lambda} \iff \lambda \in \mathbb{R}$$

Claim 2. For $A \in \mathcal{M}_{mn}(\mathbb{R})$, the eigenvalues of A^TA are real and nonnegative

Proof. We know that for $A \in \mathcal{M}_{mn}$, A^TA symmetric and from previous claim, if $A \in \mathcal{M}_{mn}(\mathbb{R})$, then A^TA is symmetric and real and with real eigenvalues

Let (λ, \mathbf{v}) be an eigenpair of A^TA , with \mathbf{v} chosen so that $\|\mathbf{v}\| = 1$

Norms are functions $V \to \mathbb{R}_+$, so $\|A \mathbf{v}\|$ and $\|A \mathbf{v}\|^2$ are ≥ 0 and thus

$$0 \le ||A\mathbf{v}||^2 = (A\mathbf{v}) \bullet (A\mathbf{v}) = (A\mathbf{v})^T (A\mathbf{v})$$

$$= \mathbf{v}^T A^T A \mathbf{v} = \mathbf{v}^T (A^T A \mathbf{v}) = \mathbf{v}^T (\lambda \mathbf{v})$$

$$= \lambda (\mathbf{v}^T \mathbf{v}) = \lambda (\mathbf{v} \bullet \mathbf{v}) = \lambda ||\mathbf{v}||^2$$

$$= \lambda \quad \square$$

Claim 3. For $A \in \mathcal{M}_{mn}(\mathbb{R})$, the nonzero eigenvalues of A^TA and AA^T are the same

Proof. Let (λ, \mathbf{v}) be an eigenpair of $A^T A$ with $\lambda \neq 0$. Then $\mathbf{v} \neq \mathbf{0}$ and

$$A^T A \mathbf{v} = \lambda \mathbf{v} \neq \mathbf{0}$$

Left multiply by A

$$AA^TA\mathbf{v} = \lambda A\mathbf{v}$$

Let $\mathbf{w} = A\mathbf{v}$, we thus have $AA^T\mathbf{w} = \lambda\mathbf{w}$; in other words, $A\mathbf{v}$ is an eigenvector of AA^T corresponding to the (nonzero) eigenvalue λ

The reverse works the same way..

The singular value decomposition (SVD)

Theorem 23 (SVD)

$$A \in \mathcal{M}_{mn}$$
 with singular values $\sigma_1 \geq \cdots \geq \sigma_r > 0$ and $\sigma_{r+1} = \cdots = \sigma_n = 0$

Then there exists $U \in \mathcal{M}_m$ orthogonal, $V \in \mathcal{M}_n$ orthogonal and a block matrix $\Sigma \in \mathcal{M}_{mn}$ taking the form

$$\Sigma = \begin{pmatrix} D & 0_{r,n-r} \\ 0_{m-r,r} & 0_{m-r,n-r} \end{pmatrix}$$

where

$$D = \mathsf{diag}(\sigma_1, \ldots, \sigma_r) \in \mathcal{M}_r$$

such that

$$A = U\Sigma V^T$$

Definition 24

We call a factorisation as in Theorem 23 the singular value decomposition of A. The columns of U and V are, respectively, the **left** and **right singular vectors** of A

U and V^T are rotation or reflection matrices, Σ is a scaling matrix

 $U \in \mathcal{M}_m$ orthogonal matrix with columns the eigenvectors of AA^T

 $V \in \mathcal{M}_n$ orthogonal matrix with columns the eigenvectors of $A^T A$

Outer product form of the SVD

Theorem 25 (Outer product form of the SVD)

 $A \in \mathcal{M}_{mn}$ with singular values $\sigma_1 \ge \cdots \ge \sigma_r > 0$ and $\sigma_{r+1} = \cdots = \sigma_n = 0$, $\boldsymbol{u}_1, \ldots, \boldsymbol{u}_r$ and $\boldsymbol{v}_1, \ldots, \boldsymbol{v}_r$, respectively, left and right singular vectors of A corresponding to these singular values

Then

$$A = \sigma_1 \mathbf{u}_1 \mathbf{v}_1^T + \dots + \sigma_r \mathbf{u}_r \mathbf{v}_r^T$$

Computing the SVD (case of \neq eigenvalues)

To compute the SVD, we use the following result

Theorem 26

Let $A \in \mathcal{M}_n$ symmetric, $(\lambda_1, \mathbf{u}_1)$ and $(\lambda_2, \mathbf{u}_2)$ be eigenpairs, $\lambda_1 \neq \lambda_2$. Then $\mathbf{u}_1 \bullet \mathbf{u}_2 = 0$

Singular values decomposition (SVD)

Proof of Theorem 26

$$A \in \mathcal{M}_n$$
 symmetric, $(\lambda_1, \boldsymbol{u}_1)$ and $(\lambda_2, \boldsymbol{u}_2)$ eigenpairs with $\lambda_1 \neq \lambda_2$

$$\lambda_{1}(\mathbf{v}_{1} \bullet \mathbf{v}_{2}) = (\lambda_{1}\mathbf{v}_{1}) \bullet \mathbf{v}_{2}$$

$$= A\mathbf{v}_{1} \bullet \mathbf{v}_{2}$$

$$= (A\mathbf{v}_{1})^{T}\mathbf{v}_{2}$$

$$= \mathbf{v}_{1}^{T}A^{T}\mathbf{v}_{2}$$

$$= \mathbf{v}_{1}^{T}(A\mathbf{v}_{2}) \qquad [A \text{ symmetric so } A^{T} = A]$$

$$= \mathbf{v}_{1}^{T}(\lambda_{2}\mathbf{v}_{2})$$

$$= \lambda_{2}(\mathbf{v}_{1}^{T}\mathbf{v}_{2})$$

$$= \lambda_{2}(\mathbf{v}_{1} \bullet \mathbf{v}_{2})$$

So
$$(\lambda_1 - \lambda_2)(\mathbf{v}_1 \bullet \mathbf{v}_2) = 0$$
. But $\lambda_1 \neq \lambda_2$, so $\mathbf{v}_1 \bullet \mathbf{v}_2 = 0$

Computing the SVD (case of \neq eigenvalues)

If all eigenvalues of A^TA (or AA^T) are distinct, we can use Theorem 26

- 1. Compute $A^TA \in \mathcal{M}_n$
- 2. Compute eigenvalues $\lambda_1, \ldots, \lambda_n$ of $A^T A$; order them as $\lambda_1 > \cdots > \lambda_n \geq 0$ (> not \geq since \neq)
- 3. Compute singular values $\sigma_1 = \sqrt{\lambda_1}, \dots, \sigma_n = \sqrt{\lambda_n}$
- 4. Diagonal matrix D in Σ is either in \mathcal{M}_n (if $\sigma_n > 0$) or in \mathcal{M}_{n-1} (if $\sigma_n = 0$)

- 5. Since eigenvalues are distinct, Theorem 26 ⇒ eigenvectors are orthogonal set. Compute these eigenvectors in the same order as the eigenvalues
- 6. Normalise them and use them to make the matrix V, i.e., $V = [\mathbf{v}_1 \cdots \mathbf{v}_n]$
- 7. To find the u_i , compute, for i = 1, ..., r,

$$oldsymbol{u}_i = rac{1}{\sigma_i} A oldsymbol{v}_i$$

and ensure that $\|\boldsymbol{u}_i\|=1$

Computing the SVD (case where some eigenvalues are =)

- 1. Compute $A^T A \in \mathcal{M}_n$
- 2. Compute eigenvalues $\lambda_1, \ldots, \lambda_n$ of $A^T A$; order them as $\lambda_1 \geq \cdots \geq \lambda_n \geq 0$
- 3. Compute singular values $\sigma_1 = \sqrt{\lambda_1}, \dots, \sigma_n = \sqrt{\lambda_n}$, with $r \leq n$ the index of the last positive singular value
- 4. For eigenvalues that are distinct, proceed as before
- 5. For eigenvalues with multiplicity > 1, we need to ensure that the resulting eigenvectors are LI and orthogonal

Dealing with eigenvalues with multiplicity > 1

When an eigenvalue has (algebraic) multiplicity > 1, e.g., characteristic polynomial contains a factor like $(\lambda - 2)^2$, things can become a little bit more complicated

The proper way to deal with this involves the so-called Jordan Normal Form (another matrix decomposition)

In short: not all square matrices are diagonalisable, but all square matrices admit a JNF

Sometimes, we can find several LI eigenvectors associated to the same eigenvalue. Check this. If not, need to use the following

Definition 27 (Generalised eigenvectors)

 $extbf{ extit{x}}
eq extbf{ extit{0}}$ generalized eigenvector of rank m of $A \in \mathcal{M}_n$ corresponding to eigenvalue λ if

$$(A - \lambda \mathbb{I})^m \mathbf{x} = \mathbf{0}$$

but

$$(A-\lambda \mathbb{I})^{m-1} \mathbf{x} \neq \mathbf{0}$$

Procedure for generalised eigenvectors

 $A \in \mathcal{M}_n$ and assume λ eigenvalue with algebraic multiplicity k

Find \mathbf{v}_1 , "classic" eigenvector, i.e., $\mathbf{v}_1 \neq \mathbf{0}$ s.t. $(A - \lambda \mathbb{I})\mathbf{v}_1 = \mathbf{0}$

Find generalised eigenvector \mathbf{v}_2 of rank 2 by solving for $\mathbf{v}_2 \neq \mathbf{0}$,

$$(A-\lambda\mathbb{I})\mathbf{v}_2=\mathbf{v}_1$$

. . .

Find generalised eigenvector \mathbf{v}_k of rank k by solving for $\mathbf{v}_k
eq \mathbf{0}$,

$$(A-\lambda \mathbb{I})\mathbf{v}_k = \mathbf{v}_{k-1}$$

Then $\{\mathbf{v}_1, \dots, \mathbf{v}_k\}$ LI

Back to the normal procedure

With the LI eigenvectors $\{v_1, \ldots, v_k\}$ corresponding to λ

Apply Gram-Schmidt to get orthogonal set

For all eigenvalues with multiplicity > 1, check that you either have LI eigenvectors or do what we just did

When you are done, be back on your merry way to step 6 in the case where eigenvalues are all \neq

I am caricaturing a little here: there can be cases that do not work exactly like this, but this is general enough..

Applications of the SVD

Many applications of the SVD, both theoretical and practical..

- 1. Obtaining a unique solutions to least squares when A^TA singular
- 2. Image compression

Least squares revisited

Theorem 28

Let $A \in \mathcal{M}_{mn}$, $\mathbf{x} \in \mathbb{R}^n$ and $\mathbf{b} \in \mathbb{R}^m$. The least squares problem $A\mathbf{x} = \mathbf{b}$ has a unique least squares solution $\tilde{\mathbf{x}}$ of minimal length (closest to the origin) given by

$$\tilde{\mathbf{x}} = A^+ \mathbf{b}$$

where A^+ is the pseudoinverse of A

Definition 29 (Pseudoinverse)

 $A = U\Sigma V^T$ an SVD for $A \in \mathcal{M}_{mn}$, where

$$\Sigma = \begin{pmatrix} D & 0 \\ 0 & 0 \end{pmatrix}$$
, with $D = \mathsf{diag}(\sigma_1, \ldots, \sigma_r)$

(D contains the nonzero singular values of A ordered as usual)

The **pseudoinverse** (or **Moore-Penrose inverse**) of A is $A^+ \in \mathcal{M}_{nm}$ given by

$$A^+ = V \Sigma^+ U^T$$

with

$$\Sigma^+ = egin{pmatrix} D^{-1} & 0 \ 0 & 0 \end{pmatrix} \in \mathcal{M}_{\it nm}$$

Compressing images

Consider an image (for simplicity, assume in shades of grey). This can be stored in a matrix $A \in \mathcal{M}_{mn}$

Take the SVD of A. Then the small singular values carry information about the regions with little variation and can perhaps be omitted, whereas the large singular values carry information about more "dynamic" regions of the image

Suppose A has r nonzero singular values. For $k \le r$, let

$$A_k = \sigma_1 \mathbf{u}_1 \mathbf{v}_1^T + \dots + \sigma_k \mathbf{u}_k \mathbf{v}_k^T$$

(so for k = r we get the usual outer product form)

Least squares problems **QR** factorisation Singular values decomposition (SVD) Principal component analysis (PCA) Support vector machines

Dimensionality reduction

One of the reasons the SVD is used is for dimensionality reduction. However, SVD has many many other uses

Now we look at another dimensionality reduction technique. PCA

PCA is often used as a blackbox technique, here we take a look at the math behind it

What is PCA?

Linear algebraic technique

Helps reduce a complex dataset to a lower dimensional one

Non-parametric method: does not assume anything about data distribution (distribution from the statistical point of view)

Principal component analysis (PCA)

A crash course on probability

A running example: fingerprints

Change of basis

Back to PCA

A 2D example to begin: hockey players

Brief "review" of some probability concepts

Proper definition of *probability* requires to use *measure theory*.. will not get into details here

A **random variable** X is a *measurable* function $X: \Omega \to E$, where Ω is a set of outcomes (*sample space*) and E is a measurable space

$$\mathbb{P}(X \in S \subseteq E) = \mathbb{P}(\omega \in \Omega | X(\omega) \in S)$$

Distribution function of a r.v., $F(x) = \mathbb{P}(X \le x)$, describes the distribution of a r.v.

R.v. can be discrete or continuous or .. other things.

Definition 30 (Variance)

Let X be a random variable. The **variance** of X is given by

$$Var X = E\left[\left(X - E(X)\right)^2\right]$$

where E is the expected value

Definition 31 (Covariance)

Let X, Y be jointly distributed random variables. The **covariance** of X and Y is given by

$$cov(X, Y) = E[(X - E(X))(Y - E(Y))]$$

Note that $cov(X,X) = E\left[(X - E(X))^2\right] = Var X$

In practice: "true law" versus "observation"

In statistics: we reason on the *true law* of distributions, but we usually have only access to a sample

We then use **estimators** to .. estimate the value of a parameter, e.g., the mean, variance and covariance

Definition 32 (Unbiased estimators of the mean and variance)

Let x_1, \ldots, x_n be data points (the *sample*) and

$$\bar{x} = \frac{1}{n} \sum_{i=1}^{n} x_i$$

be the mean of the data. An unbiased estimator of the variance of the sample is

$$\sigma^2 = \frac{1}{n-1} \sum_{i=1}^n (x_i - \bar{x})^2$$

Definition 33 (Unbiased estimator of the covariance)

Let $(x_1, y_1), \ldots, (x_n, y_n)$ be data points,

$$\bar{x} = \frac{1}{n} \sum_{i=1}^{n} x_i \text{ and } \bar{y} = \frac{1}{n} \sum_{i=1}^{n} y_i$$

be the means of the data. An estimator of the covariance of the sample is

$$cov(x,y) = \frac{1}{n} \sum_{i=1}^{n} (x_i - \bar{x})(y_i - \bar{y})$$

What does covariance do?

Variance explains how data disperses around the mean, in a 1-D case

Covariance measures the relationship between two dimensions. E.g., height and weight

More than the exact value, the sign is important:

- \triangleright cov(X,Y) > 0: both dimensions change in the same "direction"; e.g., larger height usually means higher weight
- \triangleright cov(X, Y) < 0: both dimensions change in reverse directions; e.g., time spent on social media and performance in this class
- \triangleright cov(X,Y)=0: the dimensions are independent from one another; e.g., sex/gender and "intelligence"

The covariance matrix

Typically, we consider more than 2 variables..

Definition 34

Suppose p random variables X_1, \ldots, X_p . Then the covariance matrix is the symmetric matrix

$$\begin{pmatrix} \operatorname{cov}(X_1, X_1) & \operatorname{cov}(X_1, X_2) & \cdots & \operatorname{cov}(X_1, X_p) \\ \operatorname{cov}(X_2, X_1) & \operatorname{cov}(X_2, X_2) & \cdots & \operatorname{cov}(X_2, X_p) \\ \vdots & & \vdots & & \vdots \\ \operatorname{cov}(X_p, X_1) & \operatorname{cov}(X_p, X_2) & \cdots & \operatorname{cov}(X_p, X_p) \end{pmatrix}$$

i.e., using the properties of covariance,

$$\begin{pmatrix} \operatorname{Var} X_1 & \operatorname{cov}(X_1, X_2) & \cdots & \operatorname{cov}(X_1, X_p) \\ \operatorname{cov}(X_1, X_2) & \operatorname{Var} X_2 & \cdots & \operatorname{cov}(X_2, X_p) \\ \vdots & \vdots & & \vdots \\ \operatorname{cov}(X_1, X_p) & \operatorname{cov}(X_2, X_p) & \cdots & \operatorname{Var} X_p \end{pmatrix}$$

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Principal component analysis (PCA)

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A 2D example to begin: hockey players

Example of a PCA problem

We collect a bunch of information about a bunch of people.. for instance this data from Loughborough University

This dataset contains the height, weight and 4 fingerprint measurements (length, width, area and circumference), collected from 200 participants.

What best describes a participant?

The variables

Each participant is associated to 11 variables

- "Participant Number"
- "Gender"
- ➤ "Age"
- "Dominant Hand"
- "Height (cm) (average of 3 measurements)"
- "Weight (kg) (average of 3 measurements)"
- "Fingertip Temperature (°C)"
- "Fingerprint Height (mm)"
- "Fingerprint Width (mm)"
- "Fingerprint Area (mm2)"
- "Fingerprint Circumference (mm)"

Nature of variables

Variables have different natures

- ightharpoonup "Participant Number": $\in \mathbb{N}$ (not interesting)
- "Gender": categorical
- ightharpoonup "Age": $\in \mathbb{N}$
- "Dominant Hand": categorical
- ▶ "Height (cm) (average of 3 measurements)": $\in \mathbb{R}$
- ▶ "Weight (kg) (average of 3 measurements)": $\in \mathbb{R}$
- ▶ "Fingertip Temperature (°C)": $\in \mathbb{R}$
- ▶ "Fingerprint Height (mm)": $\in \mathbb{R}$
- ightharpoonup "Fingerprint Width (mm)": $\in \mathbb{R}$
- ▶ "Fingerprint Area (mm2)": $\in \mathbb{R}$
- ▶ "Fingerprint Circumference (mm)": $\in \mathbb{R}$

Setting things up

Each participant is a row in the matrix (an observation)

Each variable is a column

So we have an 200×10 matrix (we discard the "Participant number" column)

We want to find what carries the most information

For this, we are going to project the information in a new basis in which the first "dimension" will carry most variance, the second dimension will carry a little less, etc.

In order to do so, we need to learn how to change bases

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In the following slide,

 $[x]_{\mathcal{B}}$

denotes the coordinates of \boldsymbol{x} in the basis \mathcal{B}

The aim of a change of basis is to express vectors in another coordinate system (another basis)

We do so by finding a matrix allowing to move from one basis to another

Change of basis

Definition 35 (Change of basis matrix)

$$\mathcal{B} = \{ extbf{\emph{u}}_1, \dots, extbf{\emph{u}}_n \}$$
 and $\mathcal{C} = \{ extbf{\emph{v}}_1, \dots, extbf{\emph{v}}_n \}$ bases of vector space V

The change of basis matrix $P_{\mathcal{C}\leftarrow\mathcal{B}}\in\mathcal{M}_n$.

$$P_{\mathcal{C}\leftarrow\mathcal{B}}=[[\boldsymbol{u}_1]_{\mathcal{C}}\cdots[\boldsymbol{u}_n]_{\mathcal{C}}]$$

has columns the coordinate vectors $[u_1]_{\mathcal{C}}, \ldots, [u_n]_{\mathcal{C}}$ of vectors in \mathcal{B} with respect to \mathcal{C}

Theorem 36

 $\mathcal{B} = \{ \mathbf{u}_1, \dots, \mathbf{u}_n \}$ and $\mathcal{C} = \{ \mathbf{v}_1, \dots, \mathbf{v}_n \}$ bases of vector space V and $P_{\mathcal{C} \leftarrow \mathcal{B}}$ a change of basis matrix from B to C

- 1. $\forall \mathbf{x} \in V$, $P_{\mathcal{C} \leftarrow \mathcal{B}}[\mathbf{x}]_{\mathcal{B}} = [\mathbf{x}]_{\mathcal{C}}$
- 2. $P_{\mathcal{C} \leftarrow \mathcal{B}}$ s.t. $\forall \mathbf{x} \in V$, $P_{\mathcal{C} \leftarrow \mathcal{B}}[\mathbf{x}]_{\mathcal{B}} = [\mathbf{x}]_{\mathcal{C}}$ is unique
- 3. $P_{\mathcal{C} \leftarrow \mathcal{B}}$ invertible and $P_{\mathcal{C} \leftarrow \mathcal{B}}^{-1} = P_{\mathcal{B} \leftarrow \mathcal{C}}$

Row-reduction method for changing bases

Theorem 37

 $\mathcal{B} = \{ \mathbf{u}_1, \dots, \mathbf{u}_n \}$ and $\mathcal{C} = \{ \mathbf{v}_1, \dots, \mathbf{v}_n \}$ bases of vector space V. Let \mathcal{E} be any basis for V,

$$B = [[\textbf{\textit{u}}_1]_{\mathcal{E}}, \ldots, [\textbf{\textit{u}}_n]_{\mathcal{E}}] \ \textit{and} \ C = [[\textbf{\textit{v}}_1]_{\mathcal{E}}, \ldots, [\textbf{\textit{v}}_n]_{\mathcal{E}}]$$

and let [C|B] be the augmented matrix constructed using C and B. Then

$$RREF([C|B]) = [I|P_{C \leftarrow B}]$$

If working in \mathbb{R}^n , this is quite useful with \mathcal{E} the standard basis of \mathbb{R}^n (it does not matter if $\mathcal{B} = \mathcal{E}$)

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So the question now becomes

How do we find what new basis to look at our data in?

(Changing the basis does not change the data, just the view you have of it)

(Think of what happens when you do a headstand.. your up becomes down, your right and left switch, but the world does not change, just your view of it)

(Changes of bases are fundamental operations in Science)

Principal component analysis (PCA)

A crash course on probability

A running example: fingerprints

Change of basis

Back to PCA

A 2D example to begin: hockey players

Setting things up

I will use notation (mostly) as in Joliffe's *Principal Component Analysis* (PDF of older version available for free from UofM Libraries)

$$\mathbf{x} = (x_1, \dots, x_p)$$
 vector of p random variables

We seek a linear function $\alpha_1^T \mathbf{x}$ with maximum variance, where $\alpha_1 = (\alpha_{11}, \dots, \alpha_{1p})$, i.e.,

$$\boldsymbol{\alpha}_1^T \boldsymbol{x} = \sum_{j=1}^p \alpha_{1j} x_j$$

Then we seek a linear function $\alpha_2^T x$ with maximum variance, uncorrelated to $\alpha_1^T x$

And we continue...

At kth stage, we find a linear function $\alpha_k^T \mathbf{x}$ with maximum variance, uncorrelated to $\alpha_1^T \mathbf{x}, \dots, \alpha_{k-1}^T \mathbf{x}$

 $\alpha_i^T x$ is the *i*th **principal component** (PC)

Case of known covariance matrix

Suppose we know Σ , covariance matrix of x (i.e., typically: we know x)

Then the kth PC is

$$z_k = \alpha_k^T \mathbf{x}$$

where $lpha_k$ is an eigenvector of Σ corresponding to the kth largest eigenvalue λ_k

If, additionally, $\|\alpha_k\| = \alpha_k^T \alpha = 1$, then $\lambda_k = \text{Var } z_k$

Why is that?

Let us start with

$$lpha_1^{ extit{T}}$$
x

We want maximum variance, where $\alpha_1 = (\alpha_{11}, \dots, \alpha_{1p})$, i.e.,

$$oldsymbol{lpha}_1^T oldsymbol{x} = \sum_{j=1}^p lpha_{1j} x_j$$

with the constraint that $\| {m lpha}_1 \| = 1$

We have

Var
$$\alpha_1^T \mathbf{x} = \alpha_1^T \Sigma \alpha_1$$

Objective

We want to maximise Var $\alpha_1^T x$, i.e.,

$$lpha_1^T \Sigma lpha_1$$

under the constraint that $\|lpha_1\|=1$

⇒ use **Lagrange multipliers**

Maximisation using Lagrange multipliers

(A.k.a. super-brief intro to multivariable calculus)

We want the max of $f(x_1, \ldots, x_n)$ under the constraint $g(x_1, \ldots, x_n) = k$

1. Solve

$$\nabla f(x_1,\ldots,x_n) = \lambda \nabla g(x_1,\ldots,x_n)$$
$$g(x_1,\ldots,x_n) = k$$

where $\nabla = (\frac{\partial}{\partial x_1}, \dots, \frac{\partial}{\partial x_n})$ is the **gradient operator**

2. Plug all solutions into $f(x_1, ..., x_n)$ and find maximum values (provided values exist and $\nabla g \neq \mathbf{0}$ there)

 λ is the **Lagrange multiplier**

The gradient

(Continuing our super-brief intro to multivariable calculus)

 $f: \mathbb{R}^n \to \mathbb{R}$ function of several variables, $\nabla = \left(\frac{\partial}{\partial x_1}, \dots, \frac{\partial}{\partial x_n}\right)$ the gradient operator

Then

$$\nabla f = \left(\frac{\partial}{\partial x_1} f, \dots, \frac{\partial}{\partial x_n} f\right)$$

So ∇f is a *vector-valued* function, $\nabla f: \mathbb{R}^n \to \mathbb{R}^n$; also written as

$$\nabla f = f_{x_1}(x_1,\ldots,x_n)\boldsymbol{e}_1 + \cdots f_{x_n}(x_1,\ldots,x_n)\boldsymbol{e}_n$$

where f_{x_i} is the partial derivative of f with respect to x_i and $\{e_1, \dots, e_n\}$ is the standard basis of \mathbb{R}^n

Bear with me...

(You may experience a brief period of discomfort)

$$\alpha_1^T \Sigma \alpha_1$$
 and $\|\alpha_1\|^2 = \alpha_1^T \alpha_1$ are functions of $\alpha_1 = (\alpha_{11}, \dots, \alpha_{1p})$

In the notation of the previous slide, we want the max of

$$f(\alpha_{11},\ldots,\alpha_{1p}):=\boldsymbol{\alpha}_1^T\boldsymbol{\Sigma}\boldsymbol{\alpha}_1$$

under the constraint that

$$g(\alpha_{11},\ldots,\alpha_{1p}):=\boldsymbol{\alpha}_1^T\boldsymbol{\alpha}_1=1$$

and with gradient operator

$$\nabla = \left(\frac{\partial}{\partial \alpha_{11}}, \dots, \frac{\partial}{\partial \alpha_{1p}}\right)$$

Effect of ∇ on g

g is easiest to see:

$$\nabla g(\alpha_{11}, \dots, \alpha_{1p}) = \left(\frac{\partial}{\partial \alpha_{11}}, \dots, \frac{\partial}{\partial \alpha_{1p}}\right) (\alpha_{11}, \dots, \alpha_{1p}) \begin{pmatrix} \alpha_{11} \\ \vdots \\ \alpha_{1p} \end{pmatrix}$$

$$= \left(\frac{\partial}{\partial \alpha_{11}}, \dots, \frac{\partial}{\partial \alpha_{1p}}\right) (\alpha_{11}^2 + \dots + \alpha_{1p}^2)$$

$$= (2\alpha_{11}, \dots, 2\alpha_{1p})$$

$$= 2\alpha_1$$

(And that's a general result: $\nabla \|\mathbf{x}\|_2^2 = 2\mathbf{x}$ with $\|\cdot\|_2$ the Euclidean norm)

Effect of ∇ on f

Expand (write $\Sigma = [s_{ii}]$ and do not exploit symmetry)

$$\alpha_{1}^{T} \Sigma \alpha_{1} = (\alpha_{11}, \dots, \alpha_{1p}) \begin{pmatrix} s_{11} & s_{12} & \dots & s_{1p} \\ s_{21} & s_{22} & \dots & s_{2p} \\ \vdots & \vdots & & \vdots \\ s_{p1} & s_{p2} & s_{pp} \end{pmatrix} \begin{pmatrix} \alpha_{11} \\ \alpha_{12} \\ \vdots \\ \alpha_{1p} \end{pmatrix}$$

$$= (\alpha_{11}, \dots, \alpha_{1p}) \begin{pmatrix} s_{11}\alpha_{11} + s_{12}\alpha_{12} + \dots + s_{1p}\alpha_{1p} \\ s_{21}\alpha_{11} + s_{22}\alpha_{12} + \dots + s_{2p}\alpha_{1p} \\ \vdots \\ s_{p1}\alpha_{11} + s_{p2}\alpha_{12} + \dots + s_{pp}\alpha_{1p} \end{pmatrix}$$

$$= (s_{11}\alpha_{11} + s_{12}\alpha_{12} + \dots + s_{1p}\alpha_{1p})\alpha_{11} + (s_{21}\alpha_{11} + s_{22}\alpha_{12} + \dots + s_{2p}\alpha_{1p})\alpha_{12}$$

$$\vdots \\ + (s_{p1}\alpha_{11} + s_{p2}\alpha_{12} + \dots + s_{pp}\alpha_{1p})\alpha_{1p}$$

p. 95 - Principal component analysis (PCA)

We have

e have
$$\alpha_1^T \Sigma \alpha_1 = (s_{11}\alpha_{11} + s_{12}\alpha_{12} + \dots + s_{1p}\alpha_{1p})\alpha_{11} \\ + (s_{21}\alpha_{11} + s_{22}\alpha_{12} + \dots + s_{2p}\alpha_{1p})\alpha_{12} \\ \vdots \\ + (s_{p1}\alpha_{11} + s_{p2}\alpha_{12} + \dots + s_{pp}\alpha_{1p})\alpha_{1p}$$

$$\Longrightarrow \frac{\partial}{\partial \alpha_{11}} \alpha_1^T \Sigma \alpha_1 = (s_{11}\alpha_{11} + s_{12}\alpha_{12} + \dots + s_{1p}\alpha_{1p}) + s_{11}\alpha_{11} \\ + s_{21}\alpha_{12} \\ \vdots \\ + s_{p1}\alpha_{1p} \\ = s_{11}\alpha_{11} + s_{12}\alpha_{12} + \dots + s_{1p}\alpha_{1p} \\ + s_{11}\alpha_{11} + s_{21}\alpha_{12} + \dots + s_{p1}\alpha_{1p} \\ = 2(s_{11}\alpha_{11} + s_{12}\alpha_{12} + \dots + s_{1p}\alpha_{1p})$$

 $_{p.~96}$ (last equality stems from symmetry of Σ)

In general, for $i = 1, \ldots, p$,

$$\frac{\partial}{\partial \alpha_{1i}} \boldsymbol{\alpha}_{1}^{T} \boldsymbol{\Sigma} \boldsymbol{\alpha}_{1} = s_{i1} \alpha_{11} + s_{i2} \alpha_{12} + \dots + s_{ip} \alpha_{1p}$$
$$+ s_{i1} \alpha_{11} + s_{2i} \alpha_{12} + \dots + s_{pi} \alpha_{1p}$$
$$= 2(s_{i1} \alpha_{11} + s_{i2} \alpha_{12} + \dots + s_{ip} \alpha_{1p})$$

(because of symmetry of Σ)

As a consequence,

$$\nabla \alpha_1^T \Sigma \alpha_1 = 2 \Sigma \alpha_1$$

So solving

$$\nabla f(x_1,\ldots,x_n) = \lambda \nabla g(x_1,\ldots,x_n)$$

means solving

$$2\Sigma\alpha_1 = \lambda 2\alpha_1$$

i.e.,

$$\Sigma \alpha_1 = \lambda \alpha_1$$

 \implies (λ, α_1) eigenpair of Σ , with α_1 having unit length

Picking the right eigenvalue

 (λ, α_1) eigenpair of Σ , with α_1 having unit length

But which λ to choose?

Recall that we want $\operatorname{Var} \alpha_1^T \mathbf{x} = \alpha_1^T \Sigma \alpha_1$ maximal

We have

$$\text{Var } \alpha_1^T \mathbf{x} = \alpha_1^T \Sigma \alpha_1 = \alpha_1^T (\Sigma \alpha_1) = \alpha_1^T (\lambda \alpha_1) = \lambda (\alpha_1^T \alpha_1) = \lambda$$

 \implies we pick $\lambda = \lambda_1$, the largest eigenvalue (covariance matrix symmetric so eigenvalues real)

What we have this far..

The first principal component is $\alpha_1^T \mathbf{x}$ and has variance λ_1 , where λ_1 the largest eigenvalue of Σ and α_1 an associated eigenvector with $\|\alpha_1\| = 1$

We want the second principal component to be *uncorrelated* with $\alpha_1^T \mathbf{x}$ and to have maximum variance $\text{Var } \alpha_2^T \mathbf{x} = \alpha_2^T \Sigma \alpha_2$, under the constraint that $\|\alpha_2\| = 1$

 $\alpha_2^T \mathbf{x}$ uncorrelated to $\alpha_1^T \mathbf{x}$ if $cov(\alpha_1^T \mathbf{x}, \alpha_2^T \mathbf{x}) = 0$

We have

$$cov(\alpha_1^T \mathbf{x}, \alpha_2^T \mathbf{x}) = \alpha_1^T \Sigma \alpha_2$$

$$= \alpha_2^T \Sigma^T \alpha_1$$

$$= \alpha_2^T \Sigma \alpha_1 \quad [\Sigma \text{ symmetric}]$$

$$= \alpha_2^T (\lambda_1 \alpha_1)$$

$$= \lambda \alpha_2^T \alpha_1$$

So $\alpha_2^T x$ uncorrelated to $\alpha_1^T x$ if $\alpha_1 \perp \alpha_2$

This is beginning to sound a lot like Gram-Schmidt, no?

In short

Take whatever covariance matrix is available to you (known Σ or sample S_X) – assume sample from now on for simplicity

For i = 1, ..., p, the *i*th principal component is

$$z_i = \mathbf{v}_i^T \mathbf{x}$$

where v_i eigenvector of S_X associated to the *i*th largest eigenvalue λ_i

If \mathbf{v}_i is normalised, then $\lambda_i = \text{Var } z_k$

Covariance matrix

 Σ the covariance matrix of the random variable, S_X the sample covariance matrix

 $X \in \mathcal{M}_{mp}$ the data, then the (sample) covariance matrix S_X takes the form

$$S_X = \frac{1}{n-1} X^T X$$

where the data is centred!

Sometimes you will see $S_X = 1/(n-1)XX^T$. This is for matrices with observations in columns and variables in rows. Just remember that you want the covariance matrix to have size the number of variables, not observations, this will give you the order in which to take the product

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A 2D example

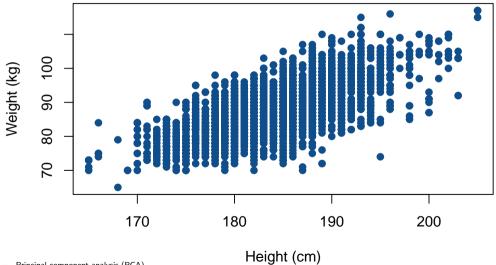
See a dataset on this page for a dataset of height and weight of some hockey players

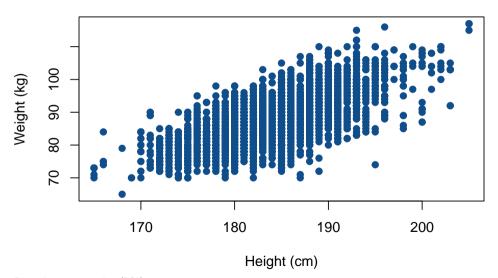
```
data = read.csv("https://figshare.com/ndownloader/files/5303173")
head(data, n=3)
                                 name position side height weight
##
    year country no
             RUS 10 tverdovsky oleg
## 1 2001
                                                      185
                                                              84 1976-
## 2 2001 RUS 2 vichnevsky vitali
                                                    188
                                                              86 1980-
## 3 2001 RUS 26 petrochinin evgeni
                                                      182
                                                              95 1976-
##
                      club
                              age cohort
                                               bmi
    anaheim mighty ducks 24.95277 1976 24.54346
## 1
## 2 anaheim mighty ducks 21.11978 1980 24.33228
## 3 severstal cherepovetal 25.22930 1976 28.68011
dim(data)
## [1] 6292
             13
```

In case you are wondering, this is a database of ice hockey players at IIHF world championships, 2001-2016, assembled by the dataset's author

See some comments here

As usual, it is a good idea to plot this to get a sense of the lay of the land





The author of the study is interested in the evolution of weights, so it is likely that the same person will be in the dataset several times. Let us check this. First check will be 'FALSE' if the number of unique names does not match the number of rows in the dataset

```
length(unique(data$name)) == dim(data)[1]
## [1] FALSE
length(unique(data$name))
## [1] 3278
```

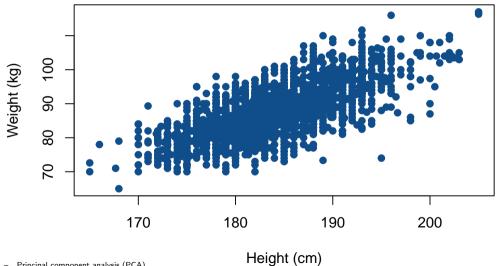
We are not interested in the evolution of weights, so far, so let us simplify things: if we have more than one record for someone, let us take their weight as the average of the

```
weights recorded for them. To be extra careful, we could check as well that there are
no major variations on player height, as this could indicate homonymies.
# First, we make a data frame with just the names
data_simplified = data.frame(name = unique(data$name))
W = C()
h = c()
for (n in data_simplified$name) {
    tmp = data[which(data$name == n),]
    h = c(h, mean(tmp\$height))
    w = c(w, mean(tmp\$weight))
```

data_simplified\$weight = w data_simplified\$height = h

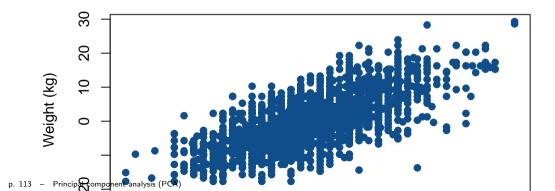
head(data_simplified)

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Centre the data and plot the result

```
data$weight.c = data$weight-mean(data$weight)
data$height.c = data$height-mean(data$height)
```



Centre the data and plot the result

```
mean(data$weight)

## [1] 87.71555

mean(data$height)

## [1] 183.8596
```

Covariance

The function 'cov' returns the covariance of two samples. Note that the functions deals equally well with data that is not centred as with data that is centred.

```
cov(data$height, data$weight)
## [1] 26.63506
cov(data$height.c, data$weight.c)
## [1] 26.63506
```

Covariance matrix

As we could see from plotting the data, there is a positive linear relationship between the two variables. Let us compute the sample covariance matrix.

```
# Select X as the matrix we want to process
X = as.matrix(data[,c("height.c", "weight.c")])
S = 1/(dim(X)[1]-1)*t(X) %*% X
S
## height.c weight.c
## height.c 29.66176 26.63506
## weight.c 26.63506 47.81112
```

Covariance matrix

The off-diagonal entries do match the computed covariance. Let us check that the variances are indeed a match too.

```
var(X[,1])
## [1] 29.66176
var(X[,2])
## [1] 47.81112
```

Hey, that works. Is math not cool? ;)

Okay, joke aside, let us now compute the principal components. For this, we need eigenvalues and eigenvectors.

```
ev = eigen(S)
ev
## eigen() decomposition
## $values
## [1] 66.87496 10.59793
##
## $vectors
##
             [.1] \qquad [.2]
## [1,] 0.5820222 -0.8131729
## [2,] 0.8131729 0.5820222
```

The result of a call to 'eigen' will not necessarily be well ordered. (Here, we are in 2D, so chances are good. When we increase the size, things might become more iffy.) So we order the eigenvalues in decreasing order and apply the same ordering to the eigenvectors. (As the eigenvectors are the columns of 'ev\$vectors', we order the columns.

```
idx_order = order(ev$values, decreasing = TRUE)
    ev$values = ev$values[idx order]
    ev$vectors = ev$vectors[, idx order]
    ev
    ## eigen() decomposition
    ## $values
    ## [1] 66.87496 10.59793
    ##
    ## $vectors
                   [.1] \qquad [,2]
p. 119 ##Principal-component analysis (RCA) 0 9131700
```

Let us normalise the first eigenvector. This way, we now that the variance of the first principal component will be the corresponding eigenvalue.

```
ev$vectors[,1] = ev$vectors[,1] / sqrt(sum(ev$vectors[,1]^2))
ev$vectors[,1]
## [1] 0.5820222 0.8131729
sum(ev$vectors[,1]^2)
## [1] 1
```

Let us plot this first eigenvector (well, the line carrying this first eigenvector). To use the function 'abline', we need to give the coefficients of the line in the form of (intercept,slope). Intercept is easy, as the line goes through the origin (by construction and because we have centred the data). The slope is also quite simple..

```
plot(data$height.c, data$weight.c,
    pch = 19, col = "dodgerblue4",
    xlab = "Height (cm)", ylab = "Weight (kg)")
abline(a = 0, b = ev$vectors[2,1]/ev$vectors[1,1],
    col = "red", lwd = 3)
```

Let us rotate the data so that the red line becomes the "x-axis". To do that, we use a rotation matrix.

$$R_{\theta} = egin{pmatrix} \cos heta & -\sin heta \ \sin heta & \cos heta \end{pmatrix}.$$

To find the angle θ , recall that tan θ is equal to opposite length over adjacent length, i.e.,

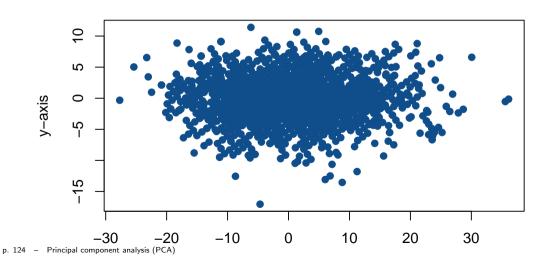
$$\tan \theta = \frac{\text{ev} \{ \text{vectors}[2, 1] \}}{\text{ev} \{ \text{vectors}[1, 1] \}}$$

So we just use the arctan of this. (Note that angles are in radians.)

```
theta = atan(ev$vectors[2,1]/ev$vectors[1,1])
     theta
     ## [1] 0.949583
    R_{\text{theta}} = \text{matrix}(c(\cos(\text{theta}), -\sin(\text{theta}),
                             sin(theta), cos(theta)).
p. 122 - Principal component analysis (PGA) = 2, byrow = TRUE)
```

And now we rotate the points. (In this case, we think of the points as vectors, of course.)

```
tmp_in = matrix(c(data$weight.c, data$height.c),
                nc = 2
tmp_out = c()
for (i in 1:dim(tmp_in)[1]) {
    tmp_out = rbind(tmp_out,
                    t(R_theta %*% tmp_in[i,]))
data$weight.c_r = tmp_out[,1]
data$height.c_r = tmp_out[,2]
```



Note that the axes have changed quite a lot, hence the very different aspect. Let us plot with the same range as for the non-rotated data for the y-axis.

```
plot(data$height.c_r, data$weight.c_r,
    pch = 19, col = "dodgerblue4",
    xlab = "x-axis", ylab = "y-axis",
    ylim = range(data$weight.c))
abline(h = 0, col = "red", lwd = 2)
```



Now do things "properly".

GS = pracma::gramSchmidt(A = ev\$vectors)

```
GS
## $Q
            [,1] \qquad [,2]
##
## [1,] 0.5820222 -0.8131729
## [2,] 0.8131729 0.5820222
##
## $R.
## [,1] [,2]
```

[1,] 1 -3.510848e-17 ## [2,] 0 1.000000e+00

Now recall we saw a theorem that told us how to construct a new basis..

```
A=matrix(c(GS\$0,1,0,0,1), nr = 2)
Α
           [,1] [,2] [,3] [,4]
##
## [1,] 0.5820222 -0.8131729 1 0
## [2,] 0.8131729 0.5820222 0 1
pracma::rref(A)
## [,1] [,2] [,3] [,4]
## [1.] 1 0 0.5820222 0.8131729
## [2,] 0 1 -0.8131729 0.5820222
P = pracma::rref(A)[,c(3,4)]
P
```

Least squares problems **QR** factorisation Singular values decomposition (SVD) Principal component analysis (PCA) Support vector machines



Clustering vs classification

Clustering is partitioning an unlabelled dataset into groups of similar objects

Classification sorts data into specific categories using a labelled dataset

Clustering

From Wikipedia

Cluster analysis or clustering is the task of grouping a set of objects in such a way that objects in the same group (called a cluster) are more similar (in some sense) to each other than to those in other groups (clusters).

There are a myriad of ways to do clustering, this is an extremely active field of research and application. See the Wikipedia page for leads

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Classification

From Wikipedia

In statistics, classification is the problem of identifying which of a set of categories (sub-populations) an observation (or observations) belongs to. Examples are assigning a given email to the "spam" or "non-spam" class, and assigning a diagnosis to a given patient based on observed characteristics of the patient (sex, blood pressure, presence or absence of certain symptoms, etc.).



Support vector machines (SVM)

We are given a training dataset of n points of the form

$$(\mathbf{x}_1, y_1), \ldots, (\mathbf{x}_n, y_n)$$

where $x_i \in \mathbb{R}^p$ and $y_i = \{-1, 1\}$. The value of y_i indicates the class to which the point x_i belongs

We want to find a surface S in \mathbb{R}^p that divides the group of points into two subgroups

Once we have this surface \mathcal{S} , any additional point that is added to the set can then be classified as belonging to either one of the sets depending on where it is with respect to the surface \mathcal{S}

Linear SVM

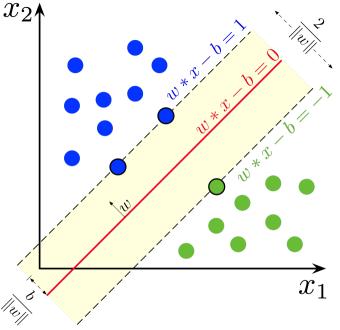
We are given a training dataset of n points of the form

$$(\mathbf{x}_1, y_1), \ldots, (\mathbf{x}_n, y_n)$$

where $\mathbf{x}_i \in \mathbb{R}^p$ and $y_i = \{-1, 1\}$. The value of y_i indicates the class to which the point \mathbf{x}_i belongs

Linear SVM – Find the "maximum-margin hyperplane" that divides the group of points \mathbf{x}_i for which $y_i = 1$ from the group of points for which $y_i = -1$, which is such that the distance between the hyperplane and the nearest point \mathbf{x}_i from either group is maximized.

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Maximum-margin hyperplane and margins for an SVM trained with samples from two classes. Samples on the margin are the **support vectors**

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Any hyperplane can be written as the set of points x satisfying

$$\mathbf{w}^\mathsf{T}\mathbf{x} - b = 0$$

where \boldsymbol{w} is the (not necessarily normalized) **normal vector** to the hyperplane (if the hyperplane has equation $a_1z_1+\cdots+a_pz_p=c$, then (a_1,\ldots,a_n) is normal to the hyperplane)

The parameter $b/\|\mathbf{w}\|$ determines the offset of the hyperplane from the origin along the normal vector \mathbf{w}

Remark: a hyperplane defined thusly is not a subspace of \mathbb{R}^p unless b=0. We can of course transform the data so that it is...

Linearly separable points

Let X_1 and X_2 be two sets of points in \mathbb{R}^p

Then X_1 and X_2 are **linearly separable** if there exist $w_1, w_2, ..., w_p, k \in \mathbb{R}$ such that

- every point $x \in X_1$ satisfies $\sum_{i=1}^p w_i x_i > k$
- every point $x \in X_2$ satisfies $\sum_{i=1}^p w_i x_i < k$

where x_i is the *i*th component of x

Hard-margin SVM

If the training data is linearly separable, we can select two parallel hyperplanes that separate the two classes of data, so that the distance between them is as large as possible

The region bounded by these two hyperplanes is called the "margin", and the maximum-margin hyperplane is the hyperplane that lies halfway between them

With a normalized or standardized dataset, these hyperplanes can be described by the equations

- $\mathbf{w}^\mathsf{T}\mathbf{x} b = 1$ (anything on or above this boundary is of one class, with label 1)
- $\mathbf{w}^\mathsf{T}\mathbf{x} b = -1$ (anything on or below this boundary is of the other class, with label -1)

Distance between these two hyperplanes is $2/\|\boldsymbol{w}\|$

 \Rightarrow to maximize the distance between the planes we want to minimize $\|\boldsymbol{w}\|$

The distance is computed using the distance from a point to a plane equation

We must also prevent data points from falling into the margin, so we add the following constraint: for each i either

$$\mathbf{w}^\mathsf{T}\mathbf{x}_i - b \ge 1$$
, if $y_i = 1$

or

$$\mathbf{w}^{\mathsf{T}}\mathbf{x}_{i} - b < -1$$
, if $y_{i} = -1$

(Each data point must lie on the correct side of the margin)

This can be rewritten as

$$y_i(\mathbf{w}^\mathsf{T}\mathbf{x}_i - b) \ge 1$$
, for all $1 \le i \le n$

or

$$y_i(\mathbf{w}^\mathsf{T}\mathbf{x}_i - b) - 1 \ge 0$$
, for all $1 \le i \le n$

We get the optimization problem:

Minimize
$$\|\mathbf{w}\|$$
 subject to $y_i(\mathbf{w}^\mathsf{T}\mathbf{x}_i - b) - 1 > 0$ for $i = 1, \ldots, n$

The **w** and *b* that solve this problem determine the classifier, $\mathbf{x} \mapsto \operatorname{sgn}(\mathbf{w}^\mathsf{T}\mathbf{x} - b)$ where $\operatorname{sgn}(\cdot)$ is the **sign function**.

The maximum-margin hyperplane is completely determined by those x_i that lie nearest to it

These x_i are the support vectors

Writing the goal in terms of Lagrange multipliers

Recall that our goal is to minimize $\|\mathbf{w}\|$ subject to $y_i(\mathbf{w}^\mathsf{T}\mathbf{x}_i - b) - 1 \ge 0$ for i = 1, ..., n

Using Lagrange multipliers $\lambda_1, \ldots, \lambda_n$, we have the function

$$L_P := F(\boldsymbol{w}, b\lambda_1, \dots, \lambda_n) = \frac{1}{2} \|\boldsymbol{w}\|^2 - \sum_{i=1}^n \lambda_i y_i(\boldsymbol{x}_i \boldsymbol{w} + b) + \sum_{i=1}^n \lambda_i$$

Note that we have as many Lagrange multipliers as there are data points. Indeed, there are that many inequalities that must be satisfied

The aim is to minimise L_p with respect to \boldsymbol{w} and b while the derivatives of L_p w.r.t. λ_i vanish and the $\lambda_i > 0$, $i = 1, \ldots, n$

Lagrange multipliers

We have already seen Lagrange multipliers, when we were studying PCA

Maximisation using Lagrange multipliers (V1.0)

We want the max of $f(x_1, \ldots, x_n)$ under the constraint $g(x_1, \ldots, x_n) = k$

1. Solve

$$\nabla f(x_1,\ldots,x_n) = \lambda \nabla g(x_1,\ldots,x_n)$$
$$g(x_1,\ldots,x_n) = k$$

where $\nabla = (\frac{\partial}{\partial x_1}, \dots, \frac{\partial}{\partial x_n})$ is the **gradient operator**

2. Plug all solutions into $f(x_1, ..., x_n)$ and find maximum values (provided values exist and $\nabla g \neq \mathbf{0}$ there)

 λ is the **Lagrange multiplier**

The gradient

 $f:\mathbb{R}^n o\mathbb{R}$ function of several variables, $\nabla=\left(rac{\partial}{\partial x_1},\ldots,rac{\partial}{\partial x_n}
ight)$ the gradient operator

Then

$$\nabla f = \left(\frac{\partial}{\partial x_1} f, \dots, \frac{\partial}{\partial x_n} f\right)$$

So ∇f is a *vector-valued* function, $\nabla f : \mathbb{R}^n \to \mathbb{R}^n$; also written as

$$\nabla f = f_{x_1}(x_1,\ldots,x_n)\boldsymbol{e}_1 + \cdots + f_{x_n}(x_1,\ldots,x_n)\boldsymbol{e}_n$$

where f_{x_i} is the partial derivative of f with respect to x_i and $\{e_1, \dots, e_n\}$ is the standard basis of \mathbb{R}^n

Lagrange multipliers (V2.0)

However, the problem we were considering then involved a single multiplier λ

Here we want $\lambda_1, \ldots, \lambda_n$

Lagrange multiplier theorem

Theorem 38

Let $f: \mathbb{R}^n \to \mathbb{R}$ be the objective function, $g: \mathbb{R}^n \to \mathbb{R}^c$ be the constraints function, both being C^1 . Consider the optimisation problem

maximize
$$f(x)$$

subject to $g(x) = 0$

Let x^* be an optimal solution to the optimization problem, such that $\operatorname{rank}(Dg(x^*)) = c < n$, where $Dg(x^*)$ denotes the matrix of partial derivatives

$$[\partial g_j/\partial x_k]$$

Then there exists a unique Lagrange multiplier $\lambda^* \in \mathbb{R}^c$ such that

$$Df(x^*) = \lambda^{*T} Dg(x^*)$$

Lagrange multipliers (V3.0)

Here we want $\lambda_1, \ldots, \lambda_n$

But we also are looking for $\lambda_i \geq 0$

So we need to consider the so-called Karush-Kuhn-Tucker (KKT) conditions

Karush-Kuhn-Tucker (KKT) conditions

Consider the optimisation problem

maximize
$$f(x)$$
 subject to $g_i(x) \leq 0$ $h_i(x) = 0$

Form the Lagrangian

$$L(\mathbf{x}, \mu, \lambda) = f(\mathbf{x}) + \mu^T \mathbf{g}(\mathbf{x}) + \lambda^T \mathbf{h}(\mathbf{x})$$

Theorem 39

If (\mathbf{x}^*, μ^*) is a saddle point of $L(\mathbf{x}, \mu)$ in $\mathbf{x} \in \mathbf{X}$, $\mu \geq \mathbf{0}$, then \mathbf{x}^* is an optimal vector for the above optimization problem. Suppose that $f(\mathbf{x})$ and $g_i(\mathbf{x})$, $i=1,\ldots,m$, are convex in \mathbf{x} and that there exists $\mathbf{x}_0 \in \mathbf{X}$ such that $\mathbf{g}(\mathbf{x}_0) < 0$. Then with an optimal vector \mathbf{x}^* for the above optimization problem there is associated a non-negative vector μ^* such that $L(\mathbf{x}^*, \mu^*)$ is a saddle point of $L(\mathbf{x}, \mu)$

KKT conditions

$$\frac{\partial}{\partial w_{\nu}} L_{P} = w_{\nu} - \sum_{i}^{n} \lambda_{i} y_{i} x_{i\nu} = 0 \qquad \nu = 1, \dots, p$$

$$\frac{\partial}{\partial b} L_{P} = -\sum_{i=1}^{n} \lambda_{i} y_{i} = 0$$

$$y_{i}(\mathbf{x}_{i}^{T} \mathbf{w} + b) - 1 \ge 0 \qquad i = 1, \dots, n$$

$$\lambda_{i} \ge 0 \qquad i = 1, \dots, n$$

$$\lambda_{i}(y_{i}(\mathbf{x}_{i}^{T} \mathbf{w} + b) - 1) = 0 \qquad i = 1, \dots, n$$

Support vector machines

Soft-margin SVM

To extend SVM to cases in which the data are not linearly separable, the **hinge loss** function is helpful

$$\max\left(0,1-y_i(\mathbf{w}^\mathsf{T}\mathbf{x}_i-b)\right)$$

 y_i is the *i*th target (i.e., in this case, 1 or -1), and $\mathbf{w}^\mathsf{T}\mathbf{x}_i - b$ is the *i*-th output

This function is zero if the constraint is satisfied, in other words, if x_i lies on the correct side of the margin

For data on the wrong side of the margin, the function's value is proportional to the distance from the margin

The goal of the optimization then is to minimize

$$\lambda \|\mathbf{w}\|^2 + \left[\frac{1}{n} \sum_{i=1}^n \max\left(0, 1 - y_i(\mathbf{w}^\mathsf{T} \mathbf{x}_i - b)\right)\right]$$

where the parameter $\lambda>0$ determines the trade-off between increasing the margin size and ensuring that the \mathbf{x}_i lie on the correct side of the margin

Thus, for sufficiently small values of λ , it will behave similar to the hard-margin SVM, if the input data are linearly classifiable, but will still learn if a classification rule is viable or not