```
In [1]: import numpy as np
    import pandas as pd
    import matplotlib.pyplot as plt
    import seaborn as sns
    from math import sqrt
    from tqdm import tqdm
    from timeit import default_timer as timer
```

Intro to ML

Ex0

1. Algebra, Random variables

 Ω = The sample space (the set of outcomes)

 $\omega =$ The outcome

 $P(\omega) > 0$ = The probability of an outcome

 $\sum_{\omega \in \Omega} P(\omega) = 1$ So sum of probabilities is 1

X = a real-valued random variable (satunnaismuuttuja)

 $X(\omega)$ creates a mapping between real numbers and the possible outcomes

a)

An operator L is linear if it obeys:

$$L(x + y) = Lx + Ly$$

where x, y are eg. functions

and

$$L(\lambda x) = \lambda L x$$

where $\lambda \in \Re$

Show that the operator E is linear:

$$E[X(\omega)] = \sum_{\omega} P(\omega)X(\omega)$$

$$E[F(\omega) + G(\omega)] = \sum_{\omega} P(\omega) (F(\omega) + G(\omega))$$

$$E[F(\omega) + G(\omega)] = \sum_{\omega} P(\omega)F(\omega) + \sum_{\omega} P(\omega)G(\omega)$$

$$E[F(\omega) + G(\omega)] = E[F(\omega)] + E[G(\omega)]$$

$$E[F + G] = E[F] + E[G]$$

and:

$$E[cF(\omega)] = \sum_{\omega} P(\omega)cF(\omega)$$
$$E[cF(\omega)] = c\sum_{\omega} P(\omega)F(\omega)$$
$$E[cF(\omega)] = cE[F(\omega)]$$

$$Var[X(\omega)] = E[(X(\omega) - E[X(\omega)])^{2}]$$

b)
$$\text{Var}[X] = E[(X - E[X])^2]$$

$$\text{Var}[X] = E[X^2 - 2XE[X] + E[X]^2]$$

$$\text{Var}[X] = E[X^2] - E[2XE[X]] + E[E[X]^2]$$

$$\text{Var}[X] = E[X^2] - 2E[X]E[X] + E[X]^2$$

$$\text{Var}[X] = E[X^2] - 2E[X]^2 + E[X]^2$$

$$\text{Var}[X] = E[X^2] - E[X]^2$$

Here E[X] is the expectation value ie. a number so eg. E[2XE[X]] = 2E[X]E[X] works

2. Bayes Rule

a) Derivation

The probability of event A given B is:

$$P(A|B) = \frac{P(A \cap B)}{P(B)}$$

The probability of event B given A is:

$$P(B|A) = \frac{P(B \cap A)}{P(A)}$$

Noting that:

$$P(B \cap A) = P(A \cap B)$$

We get:

$$P(A|B)P(B) = P(B|A)P(A)$$

Rearranging gives Bayes:

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

b) Medical test

A = allergy

 $\neg A = \text{no allergy}$

T = positive test

 $\neg T$ = negative test

false positive in 1 % of the healthy:

$$P(T|\neg A) = 0.01$$

true negative in 99 % of the healthy:

$$P(\neg T | \neg A) = 0.99$$

false negative in 10 % of the allergic:

$$P(\neg T|A) = 0.1$$

true positive in 90 % of the allergic:

$$P(T|A) = 0.9$$

15 % of the population in Finland have allergy:

$$P(A) = 0.15$$

 $P(\neg A) = 0.85$

the probability that a person is allergic, if the test is positive:

$$P(A|T) = \frac{P(A \cap T)}{P(T)}$$

$$P(A|T) = \frac{P(T|A)P(A)}{P(T)}$$

Write:

$$P(T) = P(A \cap T) + P(\neg A \cap T)$$

$$P(T) = P(T|A)P(A) + P(T|\neg A)P(\neg A)$$

Bayes with the marginal:

$$P(A|T) = \frac{P(T|A)P(A)}{P(T|A)P(A) + P(T|\neg A)P(\neg A)}$$

$$\begin{array}{c|cccc} & A & \neg A & \textbf{SUM} \\ \hline T & & 0.01 & \\ \hline \neg T & 0.1 & \\ \hline \text{SUM} & 1 & 1 & 2 \\ \hline \end{array}$$

Bayes with the marginal:

$$P(A|T) = \frac{0.9 \times 0.15}{0.9 \times 0.15 + 0.01 \times 0.85}$$

$$P(A|T) = 0.94076655$$

$$P(A|T) \approx 0.94$$

3. Matrix calculus

Matrix $A \in \mathbb{R}^{n \times n}$ Eigenvalue $\lambda \in \mathbb{R}$ of AEigenvector $x \in \mathbb{R}^n$ if Eigenfunction $Ax = \lambda x$ A has n orthonormal eigenvectors $x_i \in \mathbb{R}^n$ and corresponding eigenvalue $\lambda_i \in \mathbb{R}$ where $i \in [1, n]$

Orthonormality means:

$$\begin{aligned} \boldsymbol{x}_i^\top \boldsymbol{x}_i &= 1 \text{ and } \\ \boldsymbol{x}_i^\top \boldsymbol{x}_j &= 0 \text{ if } i \neq j \end{aligned}$$

A new matrix:

$$\mathbf{B} = \sum_{j=1}^n \lambda_j x_j x_j^{\top}$$
 "spectral decomposition"

A matrix A can be expressed in the basis of it's eigenvectors $X = [x_1 + x_2 + ... + x_n]$ and X^{\top}

$$\mathbf{A} = X \Lambda X^{\top}$$

where Λ is a diagonal matrix with $\mathrm{Tr}(\Lambda) = \sum_{i=1}^n \lambda_i$

$$A = \begin{pmatrix} \uparrow & \uparrow & & \uparrow \\ x_1 & x_2 & \dots & x_n \\ \downarrow & \downarrow & & \downarrow \end{pmatrix} \begin{pmatrix} \lambda_1 & & & 0 \\ & \lambda_2 & & \\ & & \ddots & \\ 0 & & & \lambda_n \end{pmatrix} \begin{pmatrix} \leftarrow & x_1 & \rightarrow \\ \leftarrow & x_2 & \rightarrow \\ \leftarrow & \vdots & \rightarrow \\ \leftarrow & x_n & \rightarrow \end{pmatrix}$$

Multiplying the matrices yields $A = \sum_{i=1}^{n} \lambda_i x_i x_i^{\top}$ Hence A = B and the eigenvectors and -values are the same for B as well.

4. Optimization

Constants:

 $a \in \mathbb{R}$

 $b \in \mathbb{R}$

 $c \in \mathbb{R}$

A function:

$$f(x) = ax^4 + bx + c$$
$$f'(x) = 4ax^3 + b$$

a)

The extreme values are found when f'(x) = 0

$$4ax^{3} + b = 0$$

$$4ax^{3} = -b$$

$$x^{3} = \frac{-b}{4a}$$

$$x = \sqrt[3]{\frac{-b}{4a}}$$

b)

Conditions: a > 0 and none of them can go to the infinity (and beyond)

5. Algorithms

The fibonacci numbers F(i) are defined for $i \in \mathbb{N}$ as:

$$F(i + 2) = F(i + 1) + F(i)$$
 when $F(1) = F(2) = 1$

I know that no run time experiments are needed but I was curious so here they are anyway. Also those python formulations are neat so I'm going to keep them. Required O-notations are written under the plots and in the function descriptions.

```
In [1]: def fibol(n):
    """Function to calculate values from fibonacci sequence using r
    ecursion. Scales O(2^n)"""
    if n<= 1: return n
    else: return fibol(n-1)+fibol(n-2)</pre>
```

```
In [2]:
         def fibo2(n):
              """Function to calculate valuesfrom fibonacci sequence. Scales
              return int(((1+sqrt(5))**n-(1-sqrt(5))**n)/(2**n*sqrt(5)))
In [14]:
         length = 50
In [15]:
         idx = np.arange(length)
         fibo1_df = pd.DataFrame(index=idx,columns=["Time","F"])
In [32]:
          fibo1 list = []
          fibol times = []
In [33]: for n in tqdm(range(length)):
              start = timer()
              fibo1_list.append(fibo1(n))
              end = timer()
              time = end - start
              fibo1_times.append(time)
         100% | 50/50 [3:11:21<00:00, 2436.57s/it]
In [34]: | fibo1_df["F"]=fibo1_list
          fibol df["Time"]=fibol times
In [44]: | sns.scatterplot(x=idx,y=fibo1 df.Time)
         plt.title("Fibonacci with recursion")
         plt.xlabel("n")
         plt.ylabel("Time")
Out[44]: Text(0, 0.5, 'Time')
                            Fibonacci with recursion
            5000
            4000
            3000
            2000
            1000
                  0
                         10
                                 20
                                        30
                                                        50
```

So it almost doubles the time every time: Scales $O(2^n)$ Well the big-O is the worst case scenario here (not like one usually thinks)

```
fibo2 df = pd.DataFrame(index=idx,columns=["Time","F"])
In [23]:
          fibo2 list = []
          fibo2 times = []
In [24]:
         for n in tqdm(range(length)):
              start = timer()
              fibo2_list.append(fibo2(n))
              end = timer()
              time = end - start
              fibo2_times.append(time)
         100% | 50/50 [00:00<00:00, 9364.79it/s]
         fibo2_df["F"]=fibo2_list
In [25]:
          fibo2_df["Time"]=fibo2_times
In [27]: sns.scatterplot(x=idx, y=fibo2_df.Time)
         plt.title("Fibonacci with sqrt formula")
         plt.xlabel("n")
         plt.ylabel("Time")
Out[27]: Text(0, 0.5, '')
                          Fibonacci with sqrt formula
           0.015
           0.010
           0.005
           0.000
          -0.005
          -0.010
          -0.015
```

This follows basically O(1) scaling

10

20

30

50

6. Data analysis

 V1
 V2
 V3
 V4
 V5
 V6
 V7
 V8

 53
 -4.941701
 -4.229189
 7.417093
 -2.490971
 5.551929
 0.107738
 8.076430
 -8.317607
 1

 809
 -3.022010
 -2.411519
 7.275234
 -1.656121
 4.202106
 1.116153
 7.700970
 -6.539154
 1

 981
 -3.573994
 -3.071186
 6.382967
 -1.323073
 7.302999
 1.176597
 8.920917
 -6.716963
 1

 892
 -5.807970
 -5.219509
 6.553538
 -3.051890
 6.672151
 -0.283036
 9.336143
 -9.051607
 1

4 rows × 32 columns

In [3]: x_data.shape

Out[3]: (1000, 32)

In [4]: x_data.describe()

Out[4]:

	V1	V2	V 3	V4	V 5	V 6	
count	1000.000000	1000.000000	1000.000000	1000.000000	1000.000000	1000.000000	100
mean	-5.493603	-4.848361	6.748369	-3.592207	5.432545	-1.009983	٠
std	1.860624	1.868029	1.002521	1.798006	1.561889	1.801655	
min	-12.051620	-11.449646	3.402454	-9.883042	0.650691	-7.255290	•
25%	-6.720468	-6.091409	6.098733	-4.771207	4.390488	-2.164160	1
50%	-5.483653	-4.871369	6.761059	-3.571100	5.406136	-0.987092	٠
75%	-4.240198	-3.555943	7.438199	-2.368798	6.432138	0.173440	1
max	0.765007	1.443165	10.269498	1.833573	10.671700	4.451196	1

8 rows × 32 columns

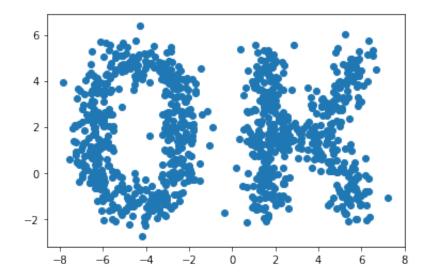
Here I do what is asked: obtain the data indices without referring to them...

```
In [5]: indices = np.where(x_data.describe().loc["std",:] > 2)[0]
In [6]: use = x_data.iloc[:,indices]
```

And here they are just for plotting but okay I use some other syntax if that pleases..

```
In [7]: plt.scatter(use.iloc[:,0],use.iloc[:,1])
```

Out[7]: <matplotlib.collections.PathCollection at 0x1a176f4668>



This should be ok 🧐

```
In [ ]:
```