Final_project

February 23, 2021

1 Final task: Dimensionality reduction via principal component analysis

```
[1]: %matplotlib inline
     import matplotlib.pvplot as plt
     from matplotlib import animation, rc
     from mpl_toolkits.mplot3d import Axes3D
     from matplotlib import cm
     from matplotlib.ticker import LinearLocator, FormatStrFormatter
     from IPython.display import HTML
     import random
     import copy
     import numpy as np
     import scipy as scy
     from tqdm.notebook import trange, tqdm
     import time
     from scipy.stats import maxwell
     #used for the video
     import subprocess
     import glob
     import os
```

1.1 Introduction:

The analysis of highly dimentional systems described by a set of correlated coordinate can be very tedous, we therefore aim to simplify the system in such a way that that the analysis becomes computationally affordable and that the simplified model still describes the system in a revelent way. In this simulation we are here looking at dimentionality reduction using Principal componant analysis (PCA). The initial correlated set of corrdinates are modified by calculating the covarience matrix, diagonalising it, and obtaining it's eingenvectors and eingenvalues. With this transformation we order the coordinated by oder of magnitude of their eingen values. The eigenvectors give the new coordinates and the ones with higher eingenvalues are the ones along which the varience of the system is highter, and therefore the ones that describe the pricipal motion of the system. With a system with n sets of coordinates we can now choose the m (with m << n) most important ones

and analyse the system in lower dimentions with this new set of coordinates.

This simulation is runed with a relatively simple system (2D) and aims to show how PCA technic can help reduce dimentions by at the same time preserving the main characteristics of the system. We are here studying a 2D potential and obtaining out data of the energy surface by implementing a Monte Carlo method with Metropolis algorithm. We will study the difference of the data in the inital coordinates (x and y) and in the new set calculated from the covarience martix σ .

1.2 Task I: Implementation

Monte Carlo (MC) method with Metropolis algorithm that samples a 2D energy surface with:

$$U(x,y) = k_b T(0.28(0.25(a \cdot x + b \cdot y)^4 + 0.1(a \cdot x + b \cdot y)^3 - 3.24(a \cdot x + b \cdot y)^2 + 6.856(a \cdot y - b \cdot x)^2) + 3.5)$$

with a = 0.809 and b = 0.588.

- Calculates the potential energy given values of x and y using the above equation.
- \rightarrow Takes : x, y
- \rightarrow Returns : U(x,y)

```
[2]: def Potential_energy (x,y):
    a = 0.809
    b = 0.588

ax_plus_by = a*x + Dimensionalityb*y
    ay_minus_bx =a*y - b*x

U = k_b * Temp * (0.28*( 0.25*ax_plus_by**4 + 0.1*ax_plus_by**3 - 3.

→24*ax_plus_by**2 + 6.856*ay_minus_bx**2)+3.5)
    return U
```

• Metropolis algorithm:

From our previous position (x,y) of sampling with choose a other position at a set distance dR but with random orientation.

- If the potential is lower at this new location we sample there.
- If it is higher we either discard the move or accept it with a certain probability depending of the difference of energy between the point positions and a ranom number.

This results in a sampling trajectory that will both tend to go towards the minimum points of energy but htat can also go over barriers of potential, with a probability proportional to the gradient of the barrier. This allow us to find sevral minimas, as long as the sampling time is large engough.

- -> Takes: (x,y) last sampling position
- -> Returns: True/False (if move if accepted of not), new/same position of sampling, potential at this position

```
[3]: def Move(x,y):
         #Calculate U1
         U_1 = Potential_energy(x,y)
         #Move by dR = 0.01 nm in a random direction
         angle = np.random.uniform(0,2*np.pi)
         dx = np.sin(angle)*dR
         dy = np.cos(angle)*dR
         #Calculate U2
         U_2 = Potential_energy(x+dx, y+dy)
         if U_1 > U_2:
             #accept move
             return True, x+dx , y+dy, U_2
         else :
             P = np.exp(-(U_2 - U_1)/(k_b*Temp))
             q = np.random.uniform(0,1)
             if q < P:
                 #accept move
                 return True, x+dx , y+dy, U_2
             else :
                 #Discard move
                 return False, x, y, U_1
```

1.3 Task II: Simulation

```
[4]: Temp = 300 #K

#Step size
dR = 0.01 #nm

#Initial position of the first sampling
x_0 = 2
y_0 = 2

# Number of accepted mooves
Nbr_MC = 2000000

k_b = 1.380649e-23 #J.K-1
Na = 6.02214086e23 #mol-1
```

- Monte carlo simulation (with Metropolis algorithm included in the funcion 'Move')
- -> Returns : x_list, and y_list of the position of the sampling over time and U_list which correspong to the value of the potential at those positions.

```
[]: #Lists used to strore the trajectory and the potential along it.
     x_list = [x_0] + [0]*Nbr_MC
     y_list = [y_0] + [0]*Nbr_MC
     U_list = [Potential_energy(x_0,y_0)] + [0]*Nbr_MC
     #Number of accepted moves
     move_Nbr = 0
     while move_Nbr < Nbr_MC :</pre>
         #Prints the percentage of mooves reasiled out of Nbr MC
         if (100*move Nbr/Nbr MC)\%5 == 0:
             print(str(100 * move_Nbr/Nbr_MC)[:3])
         # Metropolis algorithm
         check_value, x, y, U = Move( x_list[move_Nbr], y_list[move_Nbr])
         #If check value = True the move has been accepted and we perform the sample.
         # we store the data of thetrajectory in the lists. Otherwise we pass, \Box
      \hookrightarrowstaying
         # at the same position.
         if check_value == True :
             move_Nbr += 1
             x_{int}[move_Nbr] = x
             y_list[move_Nbr] = y
             U_list[move_Nbr] = U
     print('Done')
```

0.0

5.0

10.

15.

20.

• Saves the data from the simulation or retreive data from the simulation run earlier

```
[]: #np.savez('save2.npz', x_list, y_list, U_list)

[5]: npzfile = np.load('save.npz')
    x_list = npzfile['arr_0']
    y_list = npzfile['arr_1']
    U_list = npzfile['arr_2']
```

Calculating the minimum

Seeing the plot of the 2D energy surface with can distinguish 2 minimas. In the following cell we calculate their exact position and depth.

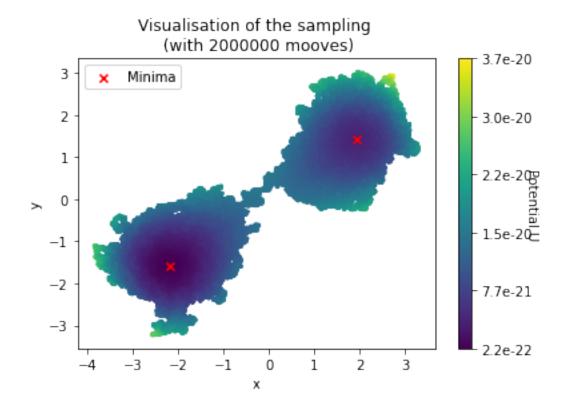
HBox(children=(FloatProgress(value=0.0, max=2000001.0), HTML(value='')))

Plot of the data

```
[7]: plt.scatter(x_list, y_list, c = U_list, s=3)
    plt.scatter([x_list[min_ind_1],x_list[min_ind_2]],__
     plt.xlabel('x')
    plt.ylabel('y')
    t = 'Visualisation of the sampling \n (with ' + str(Nbr MC) + ' mooves)'
    plt.title(t)
    plt.legend()
    cbar = plt.colorbar()
    U_ticks = np.linspace(min(U_list), max(U_list), 6)
    U_ticks = [str(U_ticks[i])[0:3] + str(U_ticks[i])[-4:] for i in_
     →range(len(U_ticks))]
    cbar.ax.set_yticklabels(U_ticks)
    cbar.set_label('Potential U', rotation=270)
    print('Minimum 1 at (x,y) = (', x_list[min_ind_1],y_list[min_ind_1], ') with U_
     \rightarrow= ', U_min_1, 'J')
    print()
    print('Minimum 2 at (x,y) = (', x_list[min_ind_2],y_list[min_ind_2], ') with U_\( \)
     \Rightarrow= ', U_min_2, 'J')
```

Minimum 1 at: (-2.183668404027096 -1.5881000022882406) with U = 2.2978051441872487e-22

Minimum 2 at: (1.9408301256758511 1.4107920871090323) with U = 4.075809248912603e-21



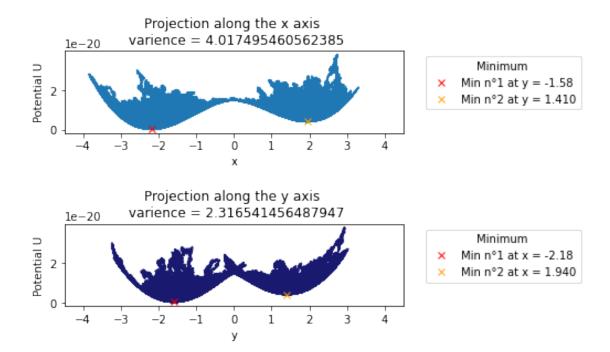
```
[9]: # Projection of the data along the x axisDimensionality
plt.subplot(211)

#Trajectory :
plt.plot(x_list, U_list)

#Indicates the mininmas:
MIN1_X = 'Min n°1 at y = ' + str(y_list[min_ind_1])[0:5]
MIN2_X = 'Min n°2 at y = ' + str(y_list[min_ind_2])[0:5]
plt.plot([x_list[min_ind_1]], [U_list[min_ind_1]], c = 'red', marker='x', u \linestyle='None', label= MIN1_X)
plt.plot([x_list[min_ind_2]], [U_list[min_ind_2]], c = 'orange', marker='x', u \linestyle='None', label= MIN2_X)

#Titles and labels for the plot :
t = 'Projection along the x axis \n varience = ' + str(np.var(x_list))
plt.title(t)
plt.xlim(-4.5, 4.5)
```

```
plt.xlabel('x')
plt.ylabel('Potential U')
plt.legend(title='Minimum', bbox_to_anchor=(1.05, 1), loc='upper left')
# Projection of the data along the y axis
plt.subplot(212)
#Trajectory :
plt.plot(y_list, U_list, c = 'midnightblue')
#Indicates the mininmas:
MIN1_Y = 'Min n°1 at x = ' + str(x_list[min_ind_1])[0:5]
MIN2_Y = 'Min n^2 at x = ' + str(x_list[min_ind_2])[0:5]
plt.plot([y_list[min_ind_1]], [U_list[min_ind_1]],c = 'red', marker='x',__
→linestyle='None',label= MIN1_Y )
plt.plot([y_list[min_ind_2]], [U_list[min_ind_2]],c = 'orange', marker='x',__
→linestyle='None',label= MIN2_Y )
#Titles and labels for the plot :
t = 'Projection along the y axis \n varience = ' + str(np.var(y_list))
plt.title(t)
plt.xlim(-4.5, 4.5)
plt.xlabel('y')
plt.ylabel('Potential U')
plt.legend(title='Minimum', bbox_to_anchor=(1.05, 1), loc='upper left')
#General adjustments :
plt.tight_layout()
plt.subplots_adjust(bottom=0.1, right=0.8, top=0.9)
plt.show()
```



We can clearly see the two minimas in the two projections, both corrdinates hold information about the main characteristic of the energy surface. Alough the varience along x is 1.7 times larger then the one along y

[]: it is describing only describing jjj % of the tatal varience of the system.

1.4 Task III: Sampling

Free energy:

$$\Delta G(x,y) = -k_b T \ln(P(x,y))$$
$$= (U2 - U1) = \Delta U$$

With:

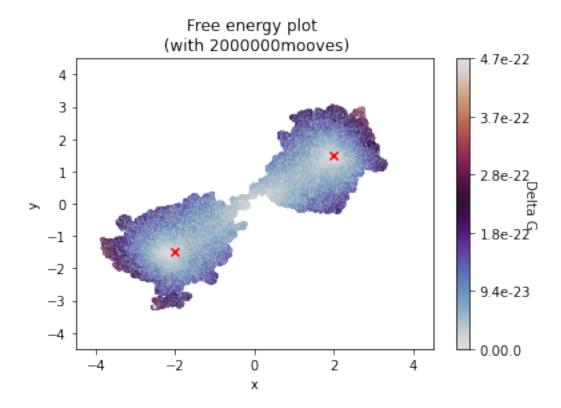
$$ln(P) = -\frac{(U2 - U1)}{k_b T}$$

The plot will then show us the gradien of the energy surface. In the cell below we calculate the free energy ΔG for our trajectory.

```
[10]: lnP_list = [0]* Nbr_MC
for i in trange (1, Nbr_MC):
    U_1 = Potential_energy( x_list[i-1], y_list[i-1])
    U_2 = Potential_energy( x_list[i], y_list[i])
    lnP = abs((U_2 - U_1))
    lnP_list[i-1] = lnP
```

```
HBox(children=(FloatProgress(value=0.0, max=1999999.0), HTML(value='')))
```

Plot



The lighter areas are ones with the lowest free energy and correspondingly with the smallest potential gradient. We can see that the free energy is minimised around the area of minimum potential but also along the conecting pathay between them. Because the simulation was run using the metropolis algorithm, the trajectory is more likely to follow a path with low gradient.

1.5 Task IV: PCA

Means

Variances

Covariances and matrix σ_{ij}

$$\sigma_{ij} = \begin{pmatrix} var(x) & cov(x,y) \\ cov(y,x) & var(y) \end{pmatrix}$$

```
_ij = [[4.01749747 2.81880472] [2.81880472 2.31654261]]
```

Eigenvalues λ_k and eigenvectors $\vec{e}_{\lambda k}$

```
Lambda1 = 6.1113318329016835 with eigenvector e1: [0.80276223 \ 0.59629925]
Lambda2 = 0.22270825116709414 with eigenvector e2: [-0.59629925 \ 0.80276223]
```

Unit vector connecting our two minimas : [0.8088054040909834, 0.58807637115703]

The eigenvectors match the data as expected, as we can see that our first vector has the same direction as the connecting axis between the two minimas and that our second vector is perpendicular to the first our as expected.

1.6 Task V: Interpretation

Angle θ between the eigenvector \vec{e}_1

```
[16]: Theta = np.arctan(eig_vec_1[0]/eig_vec_1[1])
print('Theta ~', str(round(Theta/np.pi, 3)),' ')
```

Theta ~ 0.297

The angle θ found seems correct compared to the plot of the potential displayed in Task II.

Clockwise rotation matrix R

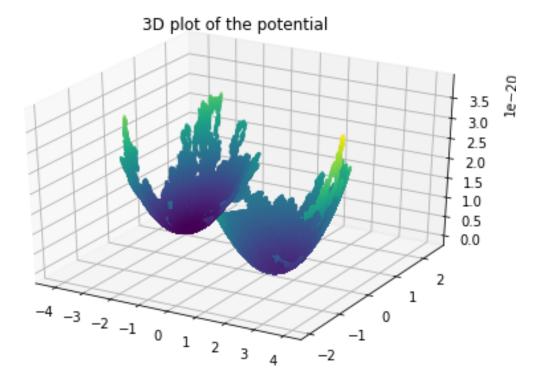
$$R(\theta) = \begin{pmatrix} Cos(\theta) & Sin(\theta) \\ -Sin(\theta) & Cos(\theta) \end{pmatrix}$$

$$R\begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} x' \\ y' \end{pmatrix}$$

For x' and y' being the data in the rotated frame and x,y our initial data.

plt.tight_layout()

plt.show()

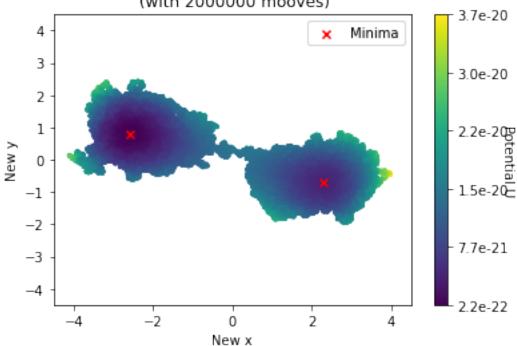


```
[20]: plt.scatter(x_rot_list, y_rot_list, c = U_list, s=3)
     plt.scatter([x_rot_list[min_ind_1],x_rot_list[min_ind_2]],__
      →marker='x',label='Minima')
     plt.xlabel('New x')
     plt.ylabel('New y')
     plt.xlim(-4.5,4.5)
     plt.ylim(-4.5,4.5)
     t = 'Visualisation of the sampling after \n clockwise rotation of ' +_{\sqcup}

→str(round(Theta/np.pi, 3)) + 'pi\n (with ' + str(Nbr_MC) + ' mooves)'
     plt.title(t)
     plt.legend()
     cbar = plt.colorbar()
     U_ticks = np.linspace(min(U_list), max(U_list), 6)
     U_ticks = [str(U_ticks[i])[0:3] + str(U_ticks[i])[-4:] for i in_
      →range(len(U_ticks))]
     cbar.ax.set_yticklabels(U_ticks)
     cbar.set_label('Potential U', rotation=270)
```

```
Minimum 1 at: (x,y) = (-2.57 \ 0.805) with U = 2.2978051441872487e-22
Minimum 2 at: (x,y) = (2.289 \ -0.71) with U = 4.075809248912603e-21
```

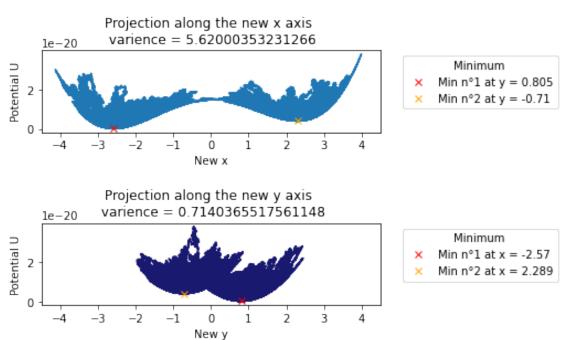
Visualisation of the sampling after clockwise rotation of 0.297pi (with 2000000 mooves)

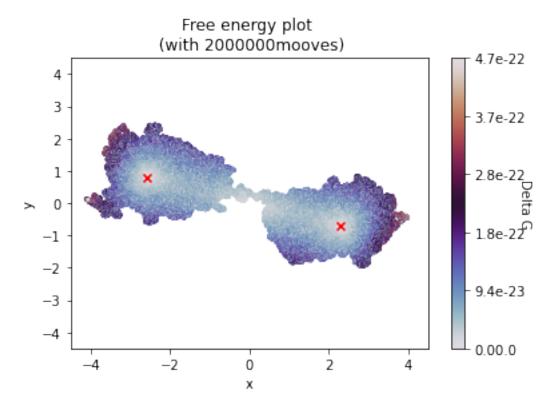


```
print('\n',str(sigma new[0,0]/v2*100)[0:4],'% of the varience of the total__
       ⇔system is')
      print('contained in the new x coordinate and ', str(sigma_new[1,1]/v2*100)[0:
       \rightarrow 4],'% in the new y')
     Previous covarience matrix :
       ij =
      [[4.01749747 2.81880472]
      [2.81880472 2.31654261]]
       63.4 % of the varience of the total system is
     contained in x and 36.5 % in y
     New covarience matrix :
       _ij =
      [[ 5.62000353 -1.62844832]
      [-1.62844832 0.71403655]]
       88.7 % of the varience of the total system is
     contained in the new x coordinate and 11.2 % in the new y
[22]: plt.subplot(211)
     plt.plot(x_rot_list, U_list)
     MIN1_X = 'Min n°1 at y = ' + str(y_rot_list[min_ind_1])[0:5]
      MIN2_X = 'Min n^2 at y = ' + str(y_rot_list[min_ind_2])[0:5]
      plt.plot([x_rot_list[min_ind_1]], [U_list[min_ind_1]],c = 'red', marker='x',u
      →linestyle='None',label= MIN1_X )
      plt.plot([x_rot_list[min_ind_2]], [U_list[min_ind_2]],c = 'orange', marker='x',u
      →linestyle='None',label= MIN2_X )
      t = 'Projection along the new x axis \n varience = ' + str(sigma_new[0,0])
      plt.title(t)
      plt.xlim(-4.5, 4.5)
      plt.xlabel('New x')
      plt.ylabel('Potential U')
      plt.legend(title='Minimum', bbox_to_anchor=(1.05, 1), loc='upper left')
      plt.subplot(212)
      plt.plot(y_rot_list, U_list, c = 'midnightblue')
     MIN1_Y = 'Min n^1 at x = ' + str(x_rot_list[min_ind_1])[0:5]
      MIN2_Y = 'Min n^2 at x = ' + str(x_rot_list[min_ind_2])[0:5]
      plt.plot([y_rot_list[min_ind_1]], [U_list[min_ind_1]],c = 'red', marker='x',_
      →linestyle='None',label= MIN1 Y )
      plt.plot([y_rot_list[min_ind_2]], [U_list[min_ind_2]],c = 'orange', marker='x',_
      →linestyle='None',label= MIN2_Y )
```

```
t = 'Projection along the new y axis \n varience = ' + str(sigma_new[1,1])
plt.title(t)
plt.xlim(-4.5, 4.5)
plt.xlabel('New y')
plt.ylabel('Potential U')
plt.legend(title='Minimum ', bbox_to_anchor=(1.05, 1), loc='upper left')

plt.tight_layout()
plt.subplots_adjust(bottom=0.1, right=0.8, top=0.9)
plt.show()
```





```
[24]: plt.subplot(211)
    plt.plot(x_rot_list[0:-1], lnP_list)

'''

MIN1_X = 'Min n°1 at y = ' + str(y_rot_list[min_ind_1])[0:5]

MIN2_X = 'Min n°2 at y = ' + str(y_rot_list[min_ind_2])[0:5]

plt.plot([x_rot_list[min_ind_1]], [U_list[min_ind_1]], c = 'red', \( \to \)

\to marker='x', label= MIN1_X )

plt.plot([x_rot_list[min_ind_2]], [U_list[min_ind_2]], c = 'orange', \( \to \)

\to marker='x', label= MIN2_X )

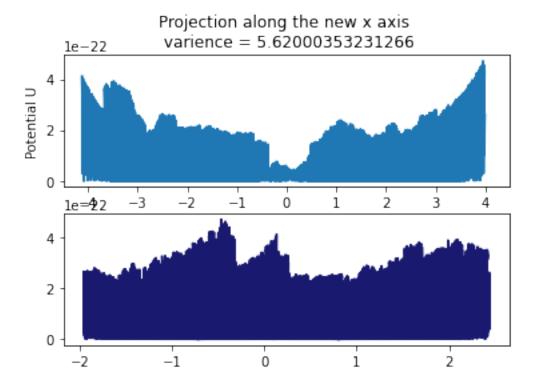
'''

t = 'Projection along the new x axis \n varience = ' + str(sigma_new[0,0])
```

```
plt.title(t)
plt.xlim(-4.5, 4.5)
plt.xlabel('New x')
plt.ylabel('Potential U')
#plt.legend(title='Minimum ', bbox_to_anchor=(1.05, 1), loc='upper left')

plt.subplot(212)
plt.plot(y_rot_list[0:-1], lnP_list, c = 'midnightblue')
```

[24]: [<matplotlib.lines.Line2D at 0x7fa4e444e760>]



From the covarience matrix and from the above plot of the data along our new x and y axis we can clearly see that most of the varience is not contained in the first componant x (88% of the varience in contrast to 63% previously).

- "the histogram of Task III" is it not a plot that we are expecting?
- ΔG equivalent to U???
- add comments on the percentages
- V?
- interpretation
- \bullet Intro

Expected percentages of varience in the intro

[]:

[]:	
[]:	