

## CLASS CS-6635, Assignment - 1

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### Part 1: Generate your own data and visualize it [30 pts]

```
In [490... ## Import all required libraries
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from math import pi
from pandas.plotting import parallel_coordinates
np.random.seed(21)
import plotly.graph_objs as go
from matplotlib.colors import ListedColormap
```

Create an array of 100 random numbers uniformly distributed between 0 and 1. Create a second array of 200 random numbers with a Gaussian distribution between 1 and 100. Note: You will need to find an appropriate mean and standard deviation for the Gaussian. It is okay if few of the numbers generated are outside the [1,100] range.

Answer: Created an array named "arr\_uniform" having 100 random numbers uniformly distributed between 0-1 and an array named "arr\_normal" having 200 random numbers followed Gaussian distribution between 1 and 100

```
In [511... ## 100 random number uniformly distributed between 0 and 1
arr_uniform = np.random.uniform(0,1,100)

## 200 random number Gaussian distribution between 1 and 100
# mean = 50, std = 17
arr_normal = np.random.normal(50,17,200)
```

1. [4 pts] Create a box plot for visualization of both arrays.

Answer: Created the box plot visualization for both the arrays represented in Fig 1. The box plot has the information of min, max, mean (diamond shape), median, lower quartile and upper quartile values. As we are sampling very few numbers from the population slight deviation in the values observed from the actual distribution. The box represents the inter quartile range bounded between lower quartile Q1 (25%) and upper quartile Q3 (75%) with the whiskers represent min and max value. In

uniform distribution the box is quite spread out because the probability of getting the numbers between 0-1 is nearly the same. However, in normal distribution the probability of getting numbers near to the mean are high as compared to the the numbers far away from mean. In the box plot of normal distribution an outlier (circle) is also noticed.

```
In [539... plt.figure(figsize=(12,5))
plt.subplot(1,2,1)

# box plot
plt.boxplot(arr_uniform, showmeans=True);
plt.grid(linestyle = "--")
plt.title('Box plot of 100 random numbers uniformly distributed between 0-1')
plt.ylabel('Distribution of Values')

median = np.median(arr_uniform)
mean = np.mean(arr_uniform)
q1 = np.percentile(arr_uniform, 25)
q3 = np.percentile(arr_uniform, 75)

plt.text(1.1, median, f'Median: {median:.2f}', verticalalignment='center')
plt.text(1.1, mean+0.06, f'Mean: {mean:.2f}', verticalalignment='center')
plt.text(1.1, q1, f'Q1: {q1:.2f}', verticalalignment='center')
plt.text(1.1, q3, f'Q3: {q3:.2f}', verticalalignment='center')
plt.subplot(1,2,2)

# box plot
plt.boxplot(arr_normal, showmeans=True);
plt.grid(linestyle = "--")
plt.title('Box plot of 200 random numbers normally distributed between 1-100')
plt.ylabel('Distribution of Values')
median = np.median(arr_normal)
mean = np.mean(arr_normal)
q1 = np.percentile(arr_normal, 25)
q3 = np.percentile(arr_normal, 75)

plt.text(1.1, median, f'Median: {median:.2f}', verticalalignment='center')
plt.text(1.1, mean+5, f'Mean: {mean:.2f}', verticalalignment='center')
plt.text(1.1, q1, f'Q1: {q1:.2f}', verticalalignment='center')
plt.text(1.1, q3, f'Q3: {q3:.2f}', verticalalignment='center')
plt.subplot(1,2,2)
plt.show()
```

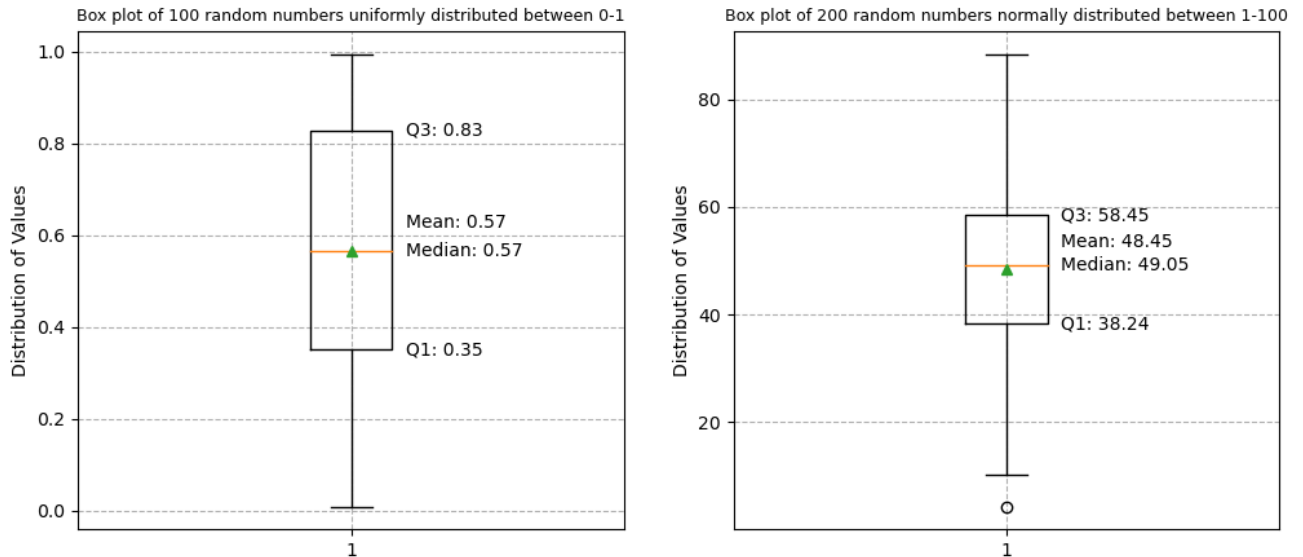


Fig.1 Left: Box plot of uniform distribution Right: Box plot of gaussian distribution

2. [4 pts] Partition the data into 20 bins and create a histogram of both arrays using the 20 bins with a bar chart. You may not use a histogram function from a plotting library, however, you may use a bar chart function.

Answer: Partitioned the data into 20 bins and created bar graph for both the arrays using 20 bins shown in fig 2. The uniform distribution of bar graph shows roughly similar frequencies between the entire range 0-1. However, the bar graph of Gaussian distribution follows a bell shape where the frequency of the bins near to the mean is higher as compared to the bins away from mean.

```
In [521.. plt.figure(figsize=(12,5))
plt.subplot(1,2,1)
# create bins
hist, bins = np.histogram(arr_uniform, bins=20)
plt.bar(bins[0:-1], hist, width=bins[1]-bins[0], edgecolor='black', align='e
plt.title('Histogram of 100 random numbers uniformly distributed between 0-1
plt.xlabel('Data between 0 and 1', fontsize = 14)
plt.ylabel('Frequency', fontsize = 14)

plt.subplot(1,2,2)
# create bins
hist, bins = np.histogram(arr_normal, bins=20)
plt.bar(bins[0:-1], hist, width=bins[1]-bins[0], edgecolor='black', align='e
plt.title('Histogram of 200 random numbers with Gaussian distribution between
plt.xlabel('Data between 1 and 100', fontsize = 14)
plt.ylabel('Frequency', fontsize = 14)
plt.show()
```

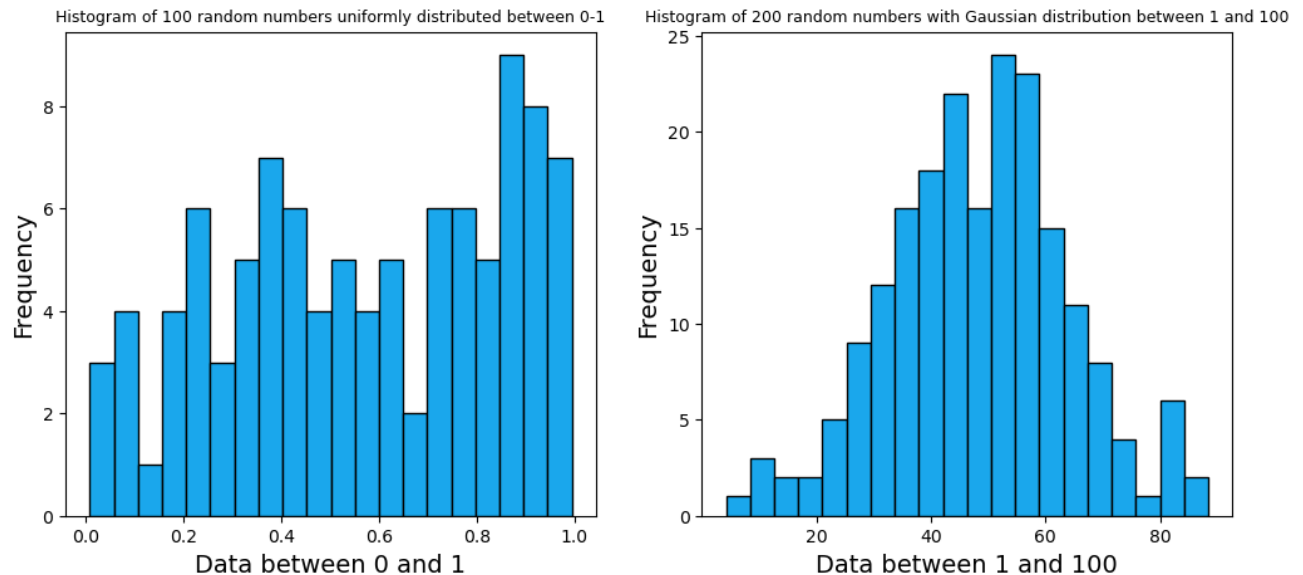


Fig.2 Left: Bar graph of uniform distribution with 20 bins Right: Bar graph of gaussian distribution with 20 bins

3. [6 pts] Write the arrays into a binary file. Read it back into an array. Visualize the arrays that were read in by plotting the cumulative distribution function as a line graph.

Answer: Wrote both the arrays in two different bin files and read the arrays back from the files. Fig 3. represents the cumulative distribution plot where the y axis represents the probability 0-1 and x axis represents the data. The CDF curve is linear for uniform distribution shows equal probability of the data. The CDF of gaussian distribution is a s shaped curve represents high probability near mean.

```
In [538]: file_path = 'uniform.bin'
          ## save the uniform array into binary file
          with open(file_path, 'wb') as file:
              np.save(file, arr_uniform)

          ## read from the file
          with open(file_path, 'rb') as file:
              arr_uniform_file = np.load(file)

          file_path = 'normal.bin'

          ## save the gaussian array into binary file
          with open(file_path, 'wb') as file:
              np.save(file, arr_normal)

          ## read from the file
          with open(file_path, 'rb') as file:
```

```

arr_normal_file = np.load(file)

# Sort the array
arr_uniform_sort = np.sort(arr_uniform_file)

# cumulative probabilities
cdf = np.arange(1, len(arr_uniform_sort) + 1) / len(arr_uniform_sort)

plt.figure(figsize=(16,6))
plt.subplot(1,2,1)
plt.plot(arr_uniform_sort, cdf, c= "#1AA7EC")
plt.grid(linestyle = '--')
plt.title('Cumulative Distribution Function for Uniform Distribution')
plt.xlabel('Data between 0 and 1', fontsize = 14)
plt.ylabel('Probability', fontsize = 14)

# Sort the array
arr_normal_sort = np.sort(arr_normal_file)

# cumulative probabilities
cdf = np.arange(1, len(arr_normal_sort) + 1) / len(arr_normal_sort)

plt.subplot(1,2,2)
# Plotting the CDF
plt.plot(arr_normal_sort, cdf,c= "#1AA7EC")
plt.grid(linestyle = '--')
plt.title('Cumulative Distribution Function for Gaussian Distribution')
plt.xlabel('Data between 1 and 100', fontsize = 14)
plt.ylabel('Probability', fontsize = 14)

```

Out[538... Text(0, 0.5, 'Probability')

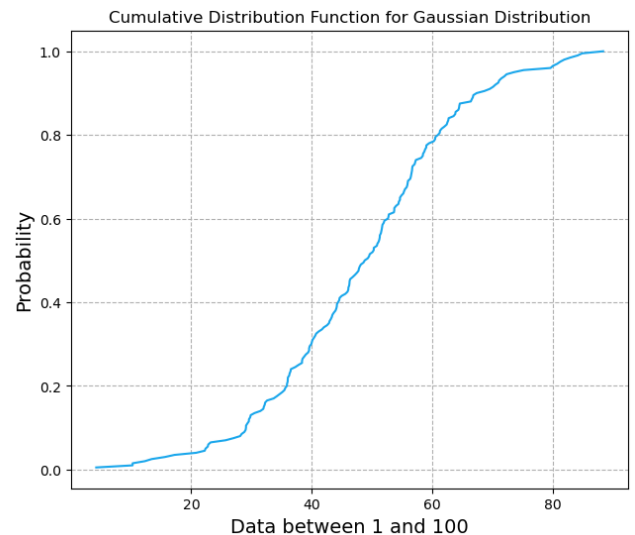
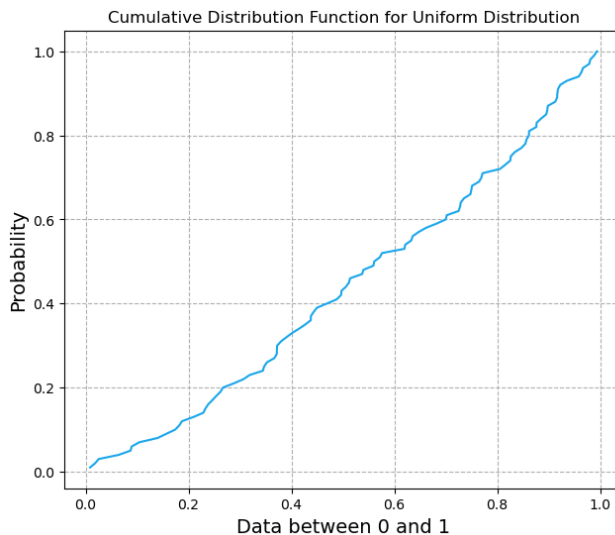


Fig.3 Left: CDF of uniform distribution Right: CDF of gaussian distribution

4. [4 pts] Create 2D arrays using uniform random sampling and gaussian random sampling with 5,000 points on  $[0,1] \times [0,1]$ . Plot the arrays with a scatter plot and compare. Note: You can use Python's built in random.sample package.

Answer Created 2D arrays using uniform random sampling and gaussian random sampling with 5,000 points on  $[0,1] \times [0,1]$ . Fig 4 shows scatter plot of 5000 x 5000 points from both the distribution. It can be referred the data points are quite concentrated near to the centre  $[0.5, 0.5]$  in case of gaussian sampling and data points are quite spread out in the entire range of  $[0,1]$  in case of uniform sampling.

```
In [530... np.random.seed(31)
arr_uniform_2d = np.random.sample((2,5000))
plt.scatter(arr_uniform_2d[0],arr_uniform_2d[1], s = 12, c= "#1AA7EC", alpha = 0.5)

arr_normal_2d = np.random.normal(0.5,0.13,(2,5000))

plt.scatter(arr_normal_2d[0],arr_normal_2d[1], c = 'black', s = 12, alpha = 0.5)
plt.xlabel(" X, values between [0,1]")
plt.ylabel(" Y, values between [0,1]")
plt.legend()
plt.title('Scatter plot of uniform and gaussian sampling data')
plt.show()
```

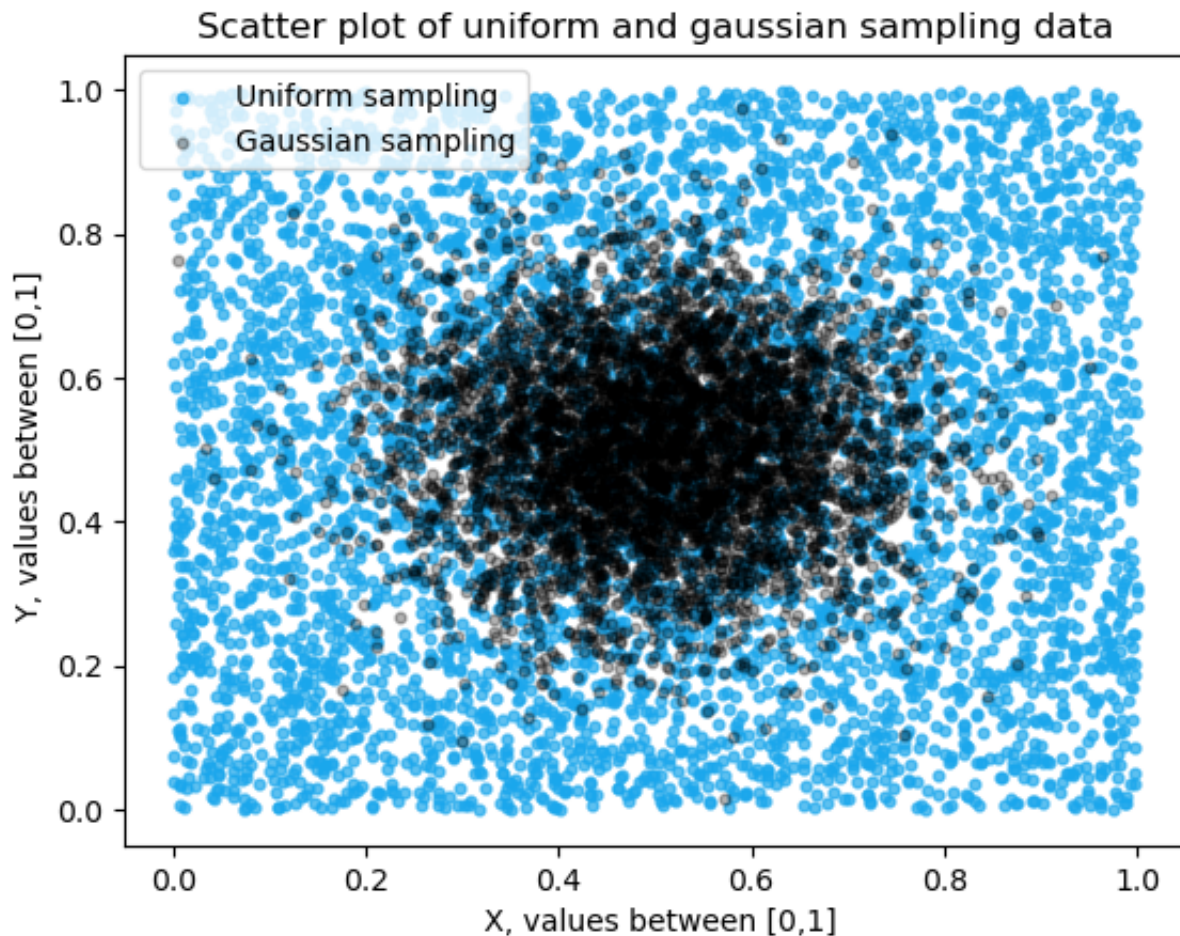


Fig.4 Scatter Plot of uniform and gaussian sampling data

4b. [8 pts] For both sampling arrays, generate 100 bins along both dimensions (think of counting the number of points in each grid cell). The output will be a 2D array of size 100x100. Show these arrays as images.

Answer: Fig 5 represents 2D histogram of uniform and gaussian sampling. The bins frequencies are spread out in uniform sampling and frequencies are concentrated near to mean in gaussian sampling.

```
In [531... plt.figure(figsize=(16,6))

plt.subplot(1,2,1)
hist,xbins, ybins, image = plt.hist2d(arr_uniform_2d[0], arr_uniform_2d[1],
plt.colorbar()
plt.xlabel(" X data")
plt.ylabel(" Y data")
plt.title('2D uniform sampling histogram')

plt.subplot(1,2,2)
hist,xbins, ybins, image = plt.hist2d(arr_normal_2d[0], arr_normal_2d[1], bi
```



```
plt.colorbar()
plt.xlabel(" X data")
plt.ylabel(" Y data")
plt.title('2D normal sampling histogram')
plt.show()
```

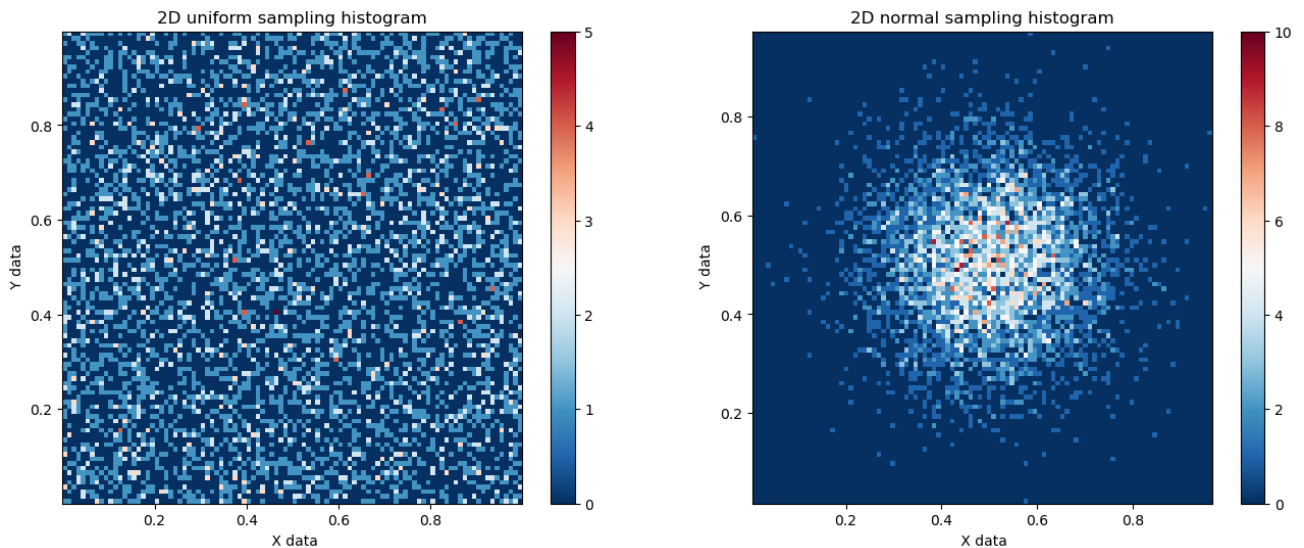


Fig.5 Left: 2D histogram of uniform distribution Right: 2D histogram of gaussian distribution

4c. [4 pts] Now plot both uniform and gaussian sampled arrays as contour plots with 10 levels. Hint: Use `tricontourf` in matplotlib as it is unstructured data.

Answer: Fig 6 represents contour plot of uniform and gaussian sampling histograms. The bins having equal values are shown as contours. Mostly similar contour values observed in entire range  $[0,1]$  in uniform sampling. In gaussian distribution near the extreme ends the values are almost 0 and high at the centre.

```
In [439... plt.figure(figsize=(16,6))
plt.subplot(1,2,1)
hist,xbins, ybins, image = plt.hist2d(arr_uniform_2d[0], arr_uniform_2d[1],
plt.contourf(hist, extent = [0,1,0,1], levels = np.linspace(hist.min(),hist.
plt.colorbar()
plt.xlabel(" X Bins data")
plt.ylabel(" Y Bins data")
plt.title('2D uniform histogram contour')

plt.subplot(1,2,2)
hist,xbins, ybins, image = plt.hist2d(arr_normal_2d[0], arr_normal_2d[1], bi
plt.contourf(hist, extent = [0,1,0,1], levels = np.linspace(hist.min(),hist.
plt.colorbar()
```



```
plt.xlabel(" X Bins data")
plt.ylabel(" Y Bins data")
plt.title('2D normal histogram contour')
plt.show()
```

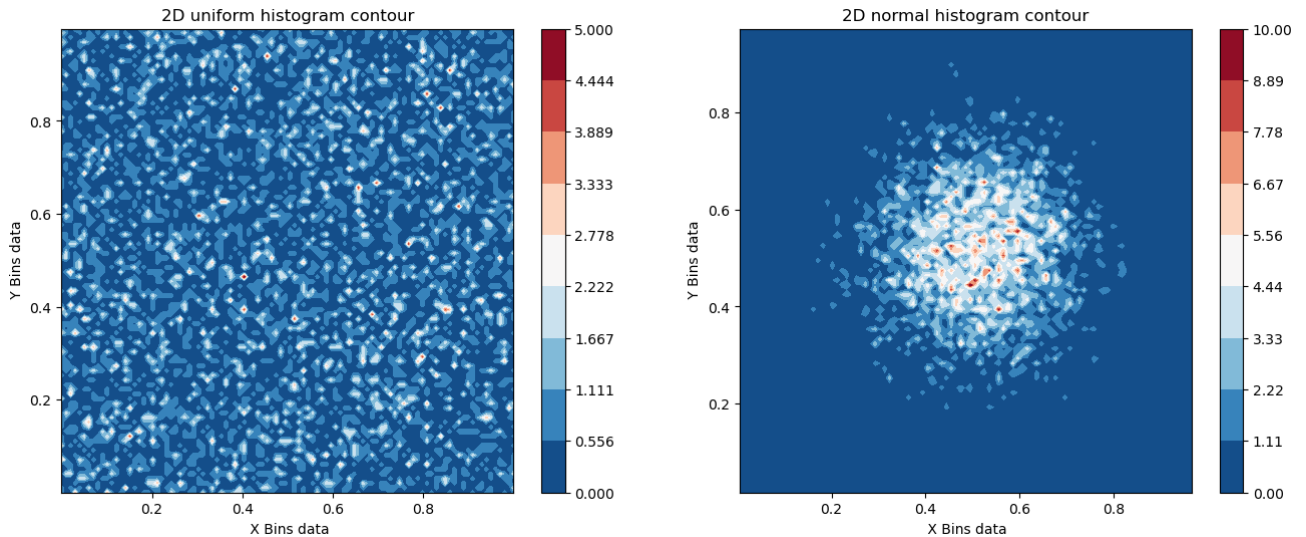


Fig 6 represents contour plot of uniform and gaussian sampling histograms

## Part 2: Intersting data sets for visualization [20 points]

1. [5 pts] Download the NOAA Land Ocean Temperature Anomalies Data Set: <https://my.eng.utah.edu/~cs6635/NOAA-Temperatures.csv>. Create a bar plot of the data. Include a label called "Year" along the x-axis and a label called Degrees F +/- From Average along the y-axis. Color each bar with either red/blue based on whether there is a positive/negative change in temperature. Describe trends in the data.

Answer: Figure 7 represents the temperature anomalies from the year 1880 to 2017 as per NOAA data set. It can be seen that from 1880 to till around 1939 the temprature difference is negative and beyond 1940 the trends are towards positive temperature difference. There are few years between 1945 to 1975 which shows a negative temperature difference. From the year 1975 to 2017 it shows increase in tempreature i.e getting hotter as the year passes. In the year 2016 it reaches a maximum value of 0.9.

```
In [449... # download the data from the given URL
df = pd.read_csv('https://my.eng.utah.edu/~cs6635/NOAA-Temperatures.csv', head=
# bar graph
plt.bar(np.asarray(df['Year'],int),df['Value'], color = ['blue' if f<0 else
plt.xlabel('Year')
plt.ylabel('Degress F +/- From Average')
plt.title('Bar graph showing land ocean temperature anomalies of NOAA dataset')
```

```
plt.show()
```

Bar graph showing land ocean temperature anomalies of NOAA dataset

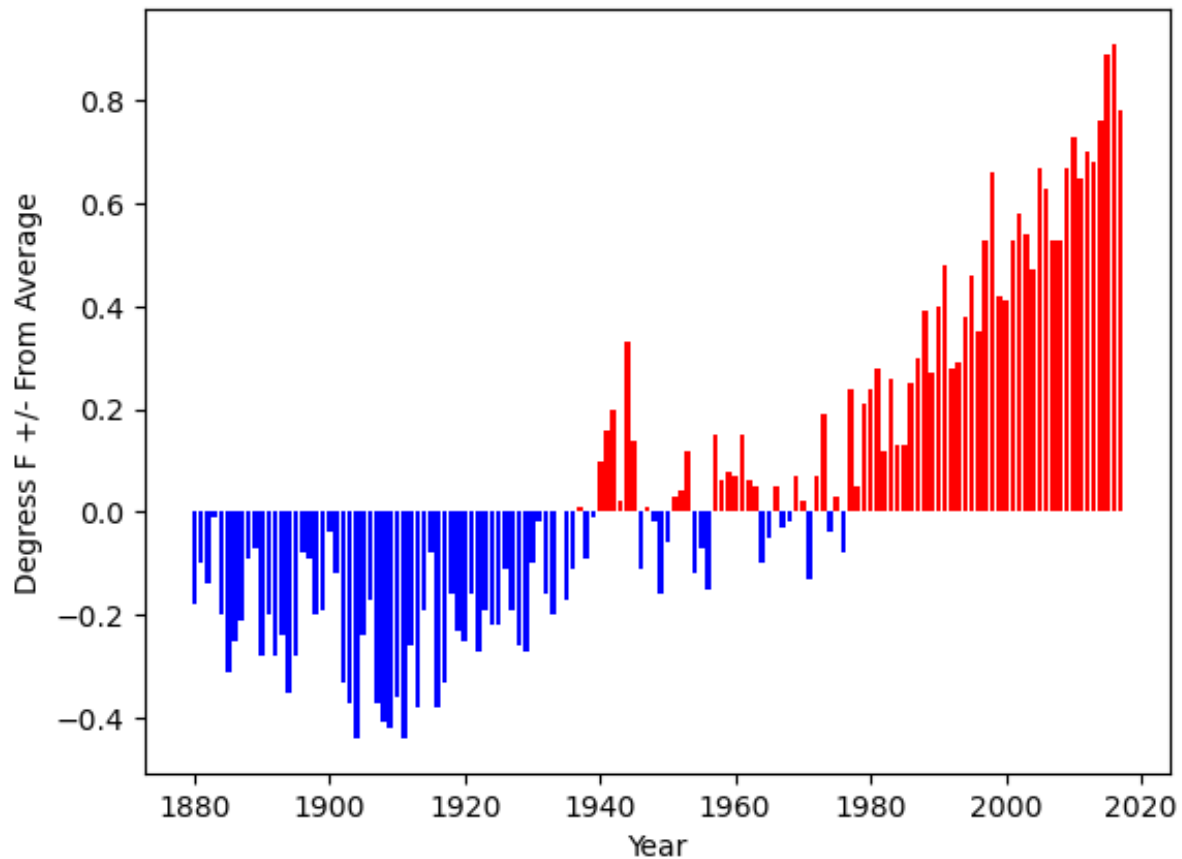


Figure 7 represents the temperature anomalies from the year 1880 to 2020 as per NOAA data set

2. [5 pts] Download the dataset

<https://my.eng.utah.edu/~cs6635/Breakfast-Cereals.xls> and generate a radar/star chart with 8 nutritional statistics for 3 cereals.

Answer: I have chosen Apple Cinnamon Cheerios, Basic 4 and Cheerios as three different Cereals and the 8 nutritional categories as 'Calories', 'Protein', 'Fat', 'Fiber', 'Carbohydrates', 'Sugars', 'Vitamins', 'Potassium'. The figure 8 shows the different nutritional values of 3 cereals. It can be inferred that Basic 4 has higher value of calories and almost similar value of potassium when compared with cheerios. There are quite overlapping areas observed in this plot which makes difficult to infer all the information from this radar plot. For example the information of Apple Cinnamon Cheerios is completely hidden.

```
In [465... # read the nutrition dataset
data_nutrition = pd.read_excel('Breakfast-Cereals.xls')
```



## Nutritional Comparison of 3 different Breakfast Cereals



Figure 8 radar plot for 3 cereals nutrition values.

```

## select the required columns
data_cereal = data_nutrition[['Cereal', 'Calories', 'Protein', 'Fat', 'Fiber',
                              'Sugars', 'Vitamins', 'Potassium']]

## select 3 cereals
selected_cereals = data_cereal.head(3)

## define the categories
columns_for_radar = ['Calories', 'Protein', 'Fat', 'Fiber', 'Carbohydrates',
                     'Sugars', 'Vitamins', 'Potassium']

## defining the data for the radar plot for each cereal r and theta
data_for_radar = []

for i, row in selected_cereals.iterrows():
    trace = go.Scatterpolar(
        r=row[columns_for_radar].values,
        theta=columns_for_radar,
        fill='toself',
        name=row['Cereal']
    )
    data_for_radar.append(trace)

layout = go.Layout(
    title='Nutritional Comparison of 3 different Breakfast Cereals',
    showlegend=True,
)
fig = go.Figure(data=data_for_radar, layout=layout)
fig.show()

```

Fig 8. Radar plot of 3 cereals having different nutritional values.

[10 pts] Five Thirty Eight maintains a server with many interesting datasets: <https://github.com/fivethirtyeight/data>. Choose two different data sets to visualize. Visualize each data set using what you think is the most appropriate plot for the given data. Choose between Parallel Coordinates and Scatter Plot for each data set and use each plotting type only once. It is also helpful to color lines or points according to certain features in the data set to get more insight. Describe the trends you can find in the data by experimenting with these plots.

Answer. The Dataset I choose is the "early senate polls". "<https://github.com/fivethirtyeight/data/blob/master/early-senate-polls/early-senate-polls.csv>". The dataset has year, election\_result, presidential\_approval and poll\_average. I represented the data in a scatter plot between election result and poll average with the colormap represents the year varying from 2006 to 2012. I convert the presidential

approval as categorical variable to represent this information as scatter plot markers shape i.e if presidential approval is > 30 % then the marker shape is "o" otherwise "x". In this way, i could able to represent the entire dataset in a single scatter plot. From the figure it can be seen that there exists a linear relationship between election result and poll average irrespective of years. If the poll average shows -ve margin the election result also follows the negative margin and vice versa. In most of the cases the presidential approval is > 30%. Only is one case in the year 2008, the presidential average is below 30% but the election result and poll average shows positive margin. In rest of the cases if the presidential average is below 30% then the poll average and election result have -ve margin.

```
In [486... data_poll = pd.read_csv('early-senate-polls.csv')
data_poll['presidential_approval'].describe()
data_poll['cat_Pa'] = ["o" if p < 30 else "x" for p in data_poll['presidenti

subset_o = data_poll[data_poll['cat_Pa'] == "o"]
subset_x = data_poll[data_poll['cat_Pa'] == "x"]

plt.figure(figsize=(8, 6))

plt.scatter(subset_o['poll_average'], subset_o['election_result'],
            c=subset_o['year'], marker='x', alpha=0.7, s = 80, label='Preside
plt.scatter(subset_x['poll_average'], subset_x['election_result'],
            c=subset_x['year'], marker='o', alpha=0.7, s = 80, label='Preside

plt.colorbar(label='Year')
plt.title('Election Result vs Poll Average in Years 2006 - 2012')
plt.xlabel('Poll Average', fontsize=14)
plt.ylabel('Election Result', fontsize=14)
plt.legend()
plt.show()
```

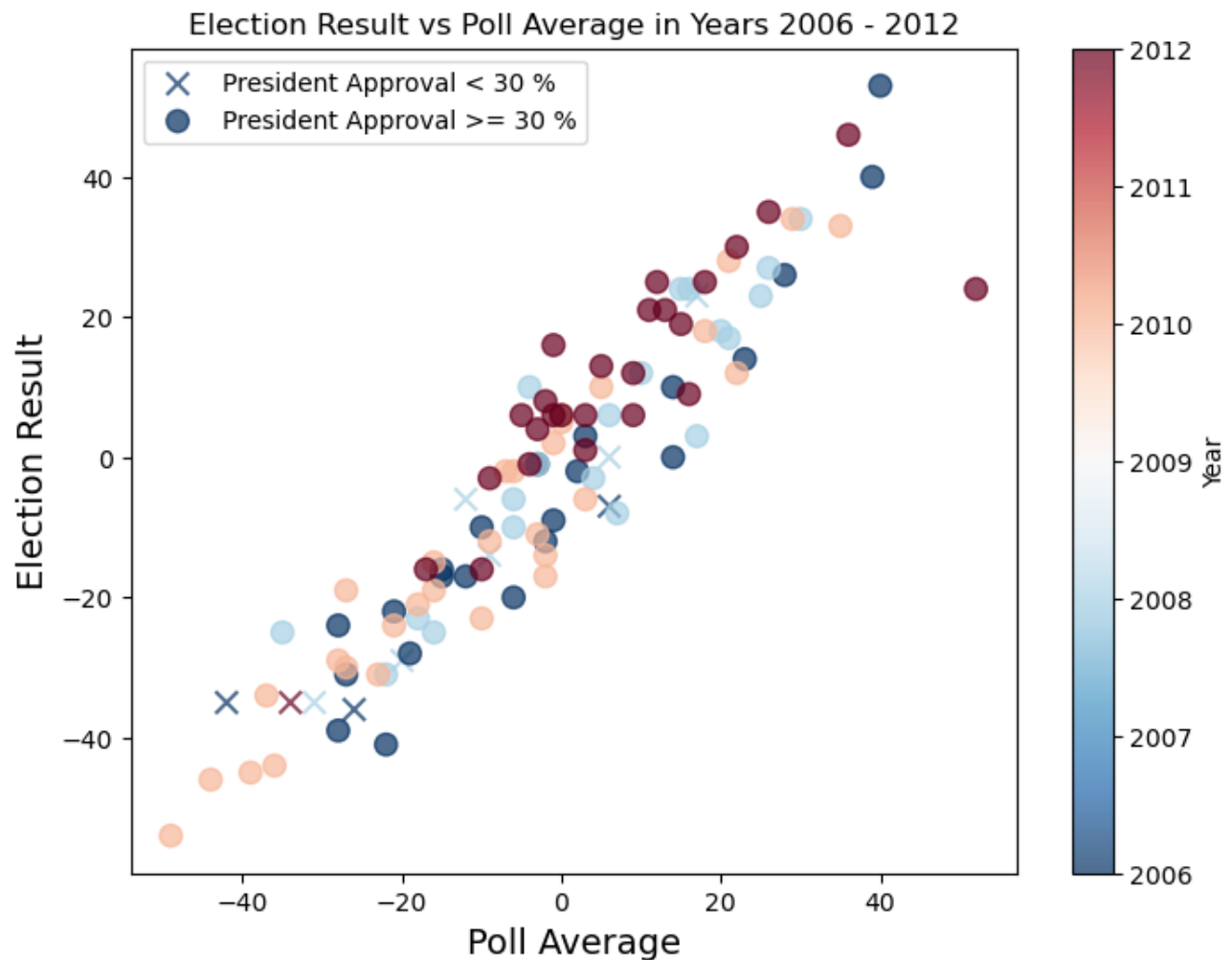


Fig 9. Scatter plot of early senate polls

Answer. The Dataset I choose is the divorce data set." <https://github.com/fivethirtyeight/data/blob/master/marriage/divorce.csv>  
 The dataset has 18 different demographic categories and their corresponding divorce values from the year 1960 to till 2012. I used the paralld coordinate plot to represent this data, where each verical bar shows the demography catgores and the line repreents the divroce values for each categoired for each year. It can be seen that the divroce values is quite high around 0.4 for the catgory poor\_4554 i.e poor in the age range of 45-54. Similarly poor in the age range of 35-44 the divorce value is around 0.35. Lowest divorce value around 0.05 has been observed for the rich in the age range of 35-44 and rich in the age range of 45-54. Furthermore as year passes by the divorce value also increases. Year 1960 shows lowest divorce values followed by 1970 and 1980.

Dataset "<https://github.com/fivethirtyeight/data/blob/master/marriage/divorce.csv>"

```
In [493... data_divorce = pd.read_csv('divorce.csv')
```

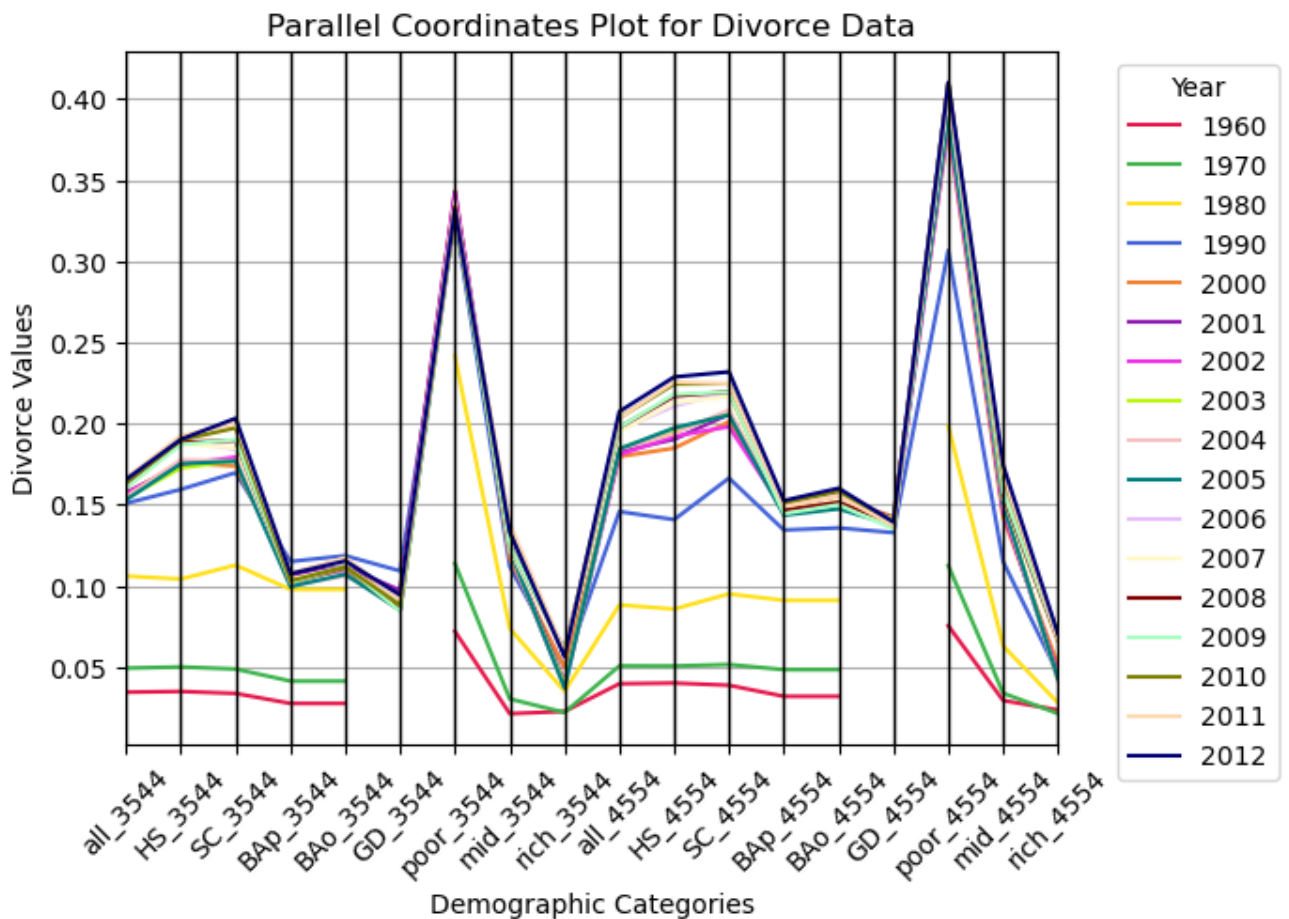
```

data_divorce = data_divorce.drop(['Unnamed: 0', 'date'], axis=1)
color_list = ['#e6194b', '#3cb44b', '#ffe119', '#4363d8', '#f58231',
              '#911eb4', '#46f0f0', '#f032e6', '#bcf60c', '#fabebe',
              '#008080', '#e6beff', '#9a6324', '#fffac8', '#800000',
              '#aaffc3', '#808000', '#ffd8b1', '#000075']

parallel_coordinates(data_divorce, class_column='year', colormap=ListedColorMap)

plt.title('Parallel Coordinates Plot for Divorce Data')
plt.xlabel('Demographic Categories')
plt.ylabel('Divorce Values')
plt.xticks(rotation=45)
plt.legend(title='Year', bbox_to_anchor=(1.05, 1), loc='upper left')
plt.show()

```



### Part 3: Questions on The Value of Visualization Paper [20 pts]

(<https://my.eng.utah.edu/~cs6635/Value-of-Visualization.pdf>)

"The Value of Visualization" by Jarke J. Van Wijk



**1 [5 pts] Why is assessing value of visualizations important? What are the two measures for deciding the value of visualizations?**

Assesing the the value of visualizations is important because it helps to evaluate how efficiently and effectively visulization guides in decision making, understanding and representing information. Two key measures which decides the value of visualizations are effectivness and efficiency. An effective visualization tool should be able to guide on decision making, providing insight into the large dataset and improves the existing knowledge of the user. Efficiency is related to the usage of resources such as time, effort and hardware while developing an efficient visualization tool should use less resources.

**2 [5 pts] Briefly describe a mathematical model for the visualization block shown in Fig. 1.**

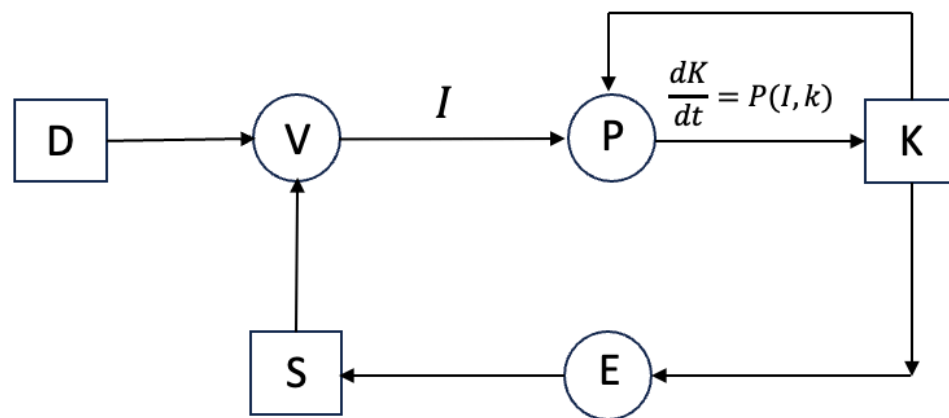


Figure 1 of the paper represents the general workflow how visualization works. In the figure the visualization function  $V$  transforms the input data  $D$  into time varying image  $I(t)$  guided by certain specification  $S$ . This specification  $S$  represents the specification of hardware, algorithm and specific parameters. The output  $I$  can be an image/animation/ auditory or haptic feedback. This output is perceived by the user resulting in increase in knowledge  $K$ , which is evaluate as rate of change of knowledge with time. This rate of change depends on the current knowledge of the user, the cognitive and perceptual abiltiy of the user and the characteristic of the output image. Furthermore, the user adapts the specifications based on the updated knowledge and explores the data further. This represents an interactive way of learning and understanding the data

through visualization.

### 3 [5 pts] State four parameters that describe the costs associated with any visualization technique.

There are four different cost that describe the total costs associated with any visualization technique.

1. Initial development cost : Cost associated with development and implementation of the visualization technique for example hardware cost.
2. Initial cost per user : Time spend by user to select and acquire of Visualization technique and understanding its usage and changing it as per requirement.
3. Initial cost per session : Data conversion cost.
4. Perception and exploration cost: User time in understanding the visualization and exploring the data.

Number of users ( $n$ ), how many times ( $m$ ) each user visualize the data, how many exploration steps ( $k$ ) in each session and time  $T$  also affects the total cost. For a great visualization tool  $n, m$  should be on higher side i.e number of users using the visualization tool frequently with lower value of exploration ( $k$ ) and time ( $T$ ).

### 4 [5 pts] What are the pros and cons of interactivity of visualizations?##

Pros:

1. Enhances the understanding of the data
2. Exploration of the data and get improved insights
3. Developing need basis interactive tool to understand the data better

Cons:

1. Freely modifying the visualization results into subjectivity user can modify the visualization to emphasize the result he/she wants to show.
2. Difficulty in comparison when higher level of customization is done in visualization.
3. Higher cost because of the interaction.

## PART 4: 3D Scalar Volume Dataset

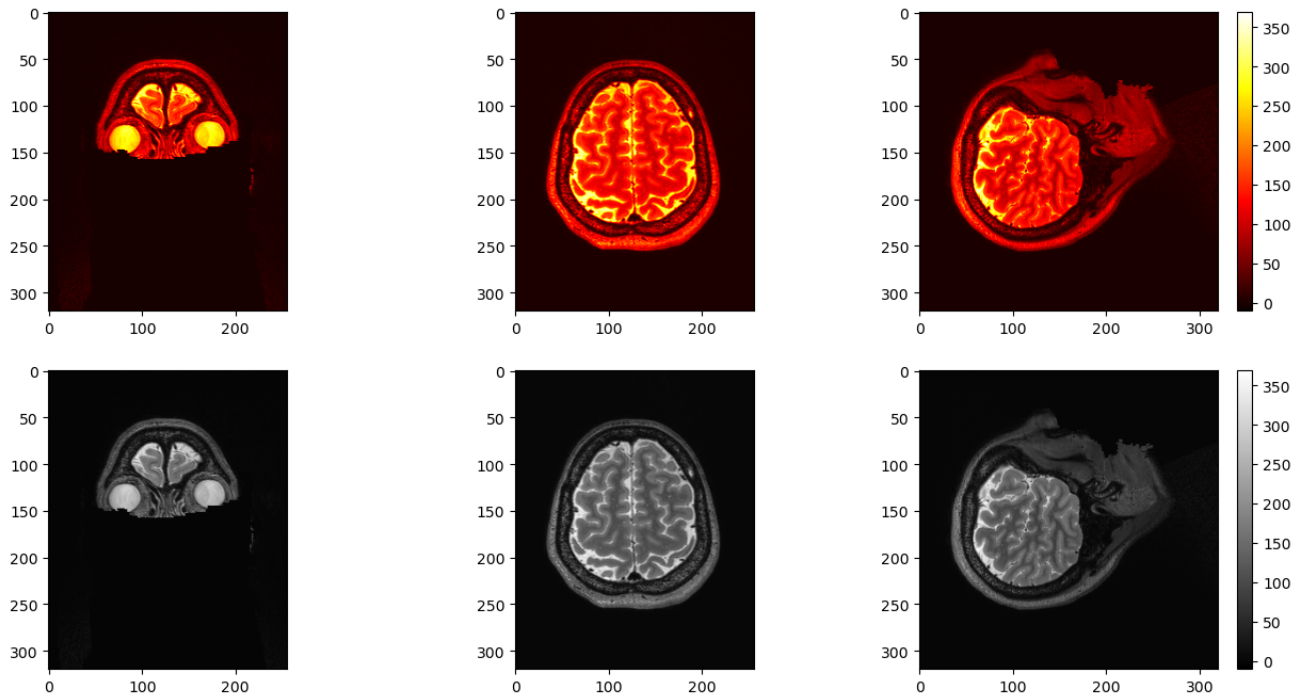
MATLAB/Python also can be used for analysis and visualization of 3D volume data sets, such as brain MRI images. Download the brain MRI data set from <https://my.eng.utah.edu/~cs6635/T2.nii.gz> . The data format is .nii with 320 x 320 x 256 dimensions. Load data in MATLAB/Python.

Extract one slice for each axis (three slices total) from the volume and save them as images. Use at least two colormaps to show the three image set and describe the difference this choice makes.

Answer: I took 3 different slices from 3 different axis. I have used two different color map "hot" and "gray" to represent the data. The magma colormap shows a clear gradient of colors depending on the values. Higher values have yellowish color with 0 values have black color. In case of gray color map its the gradient form black to gray with higher values have gray color. I can see the difference in both the colormap however the hot colormap is very prominent and distictive.

```
In [537... import nibabel as nib
mri = nib.load("T2.nii")
mri.shape
plt.figure(figsize=(16,8))
mri_data = mri.get_fdata()
plt.subplot(2,3,1)
plt.imshow(mri_data[56,:,:], cmap='hot')
## save image
plt.savefig('x_mri.jpg', format='jpg')
plt.subplot(2,3,2)
plt.imshow(mri_data[:,56,:], cmap='hot')
# save image
plt.savefig('y_mri.jpg', format='jpg')
plt.subplot(2,3,3)
plt.imshow(mri_data[:, :, 56], cmap='hot')
# save image
plt.savefig('z_mri.jpg', format='jpg')
plt.colorbar()
plt.subplot(2,3,4)
plt.imshow(mri_data[56,:,:], cmap='gray')
plt.subplot(2,3,5)
plt.imshow(mri_data[:,56,:], cmap='gray')
plt.subplot(2,3,6)
plt.imshow(mri_data[:, :, 56], cmap='gray')
plt.colorbar()
```

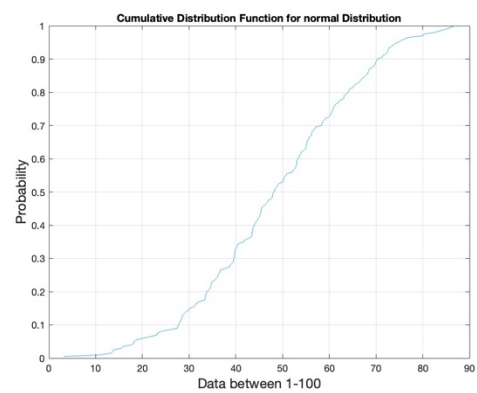
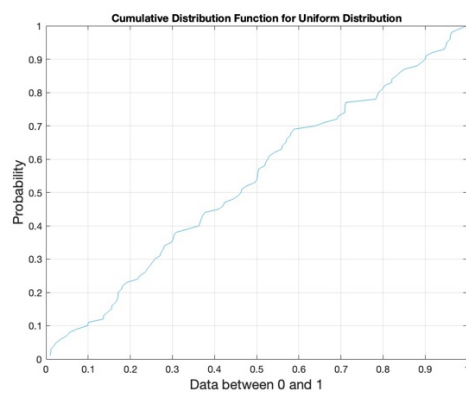
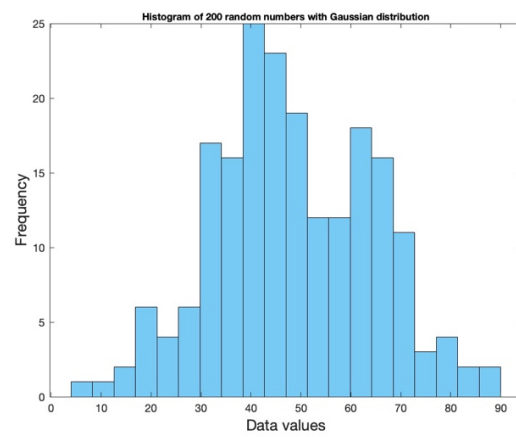
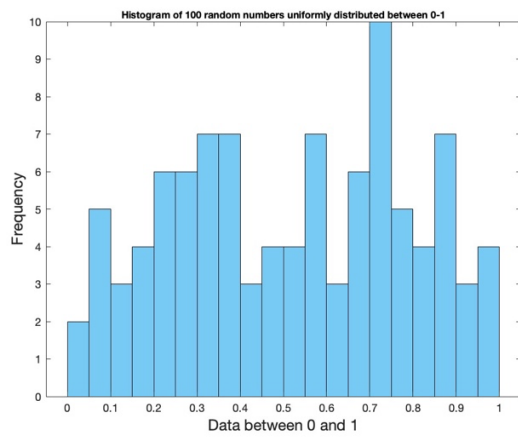
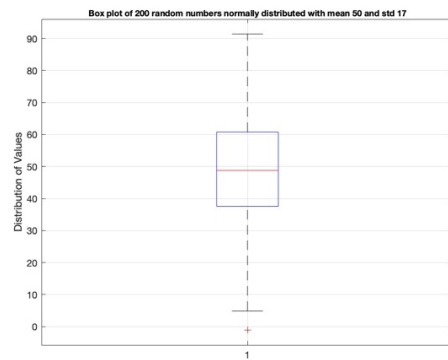
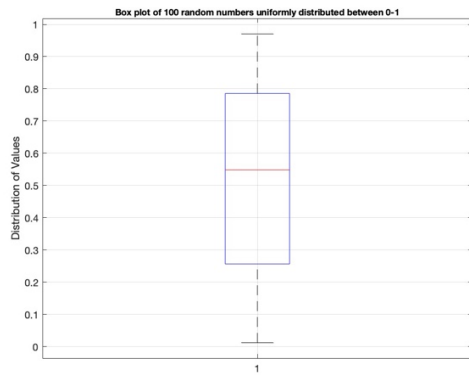
Out[537... <matplotlib.colorbar.Colorbar at 0x2bffcbbd30>

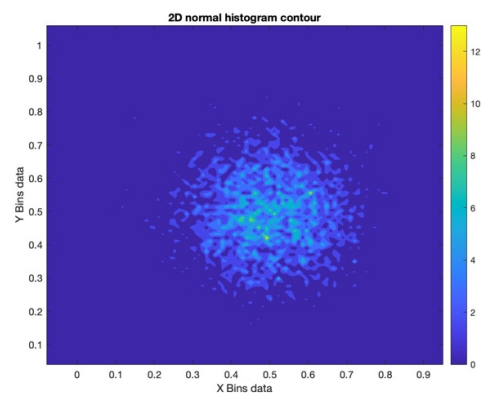
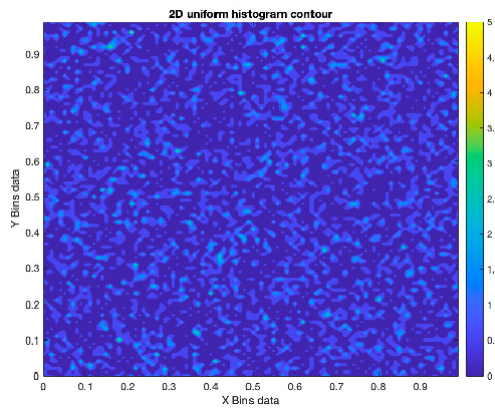
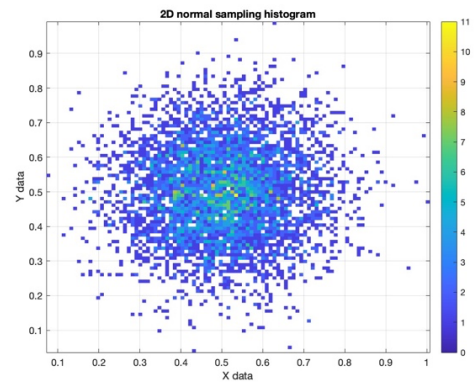
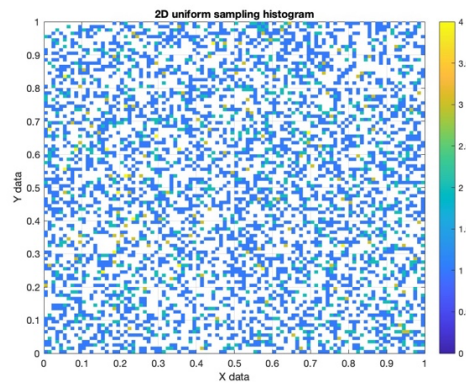
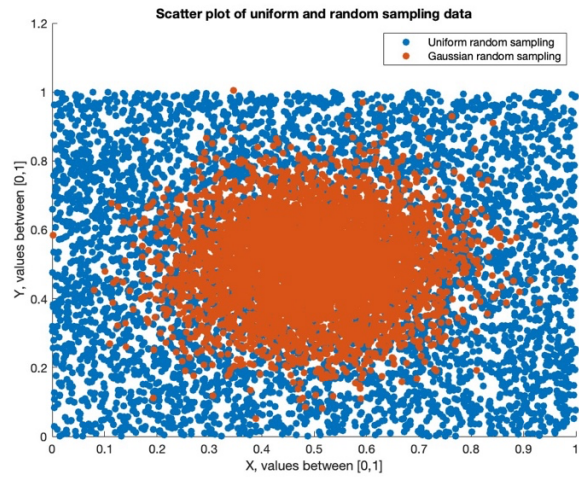


#### References:

1. [https://matplotlib.org/stable/api/\\_as\\_gen/matplotlib.pyplot.contourf.html](https://matplotlib.org/stable/api/_as_gen/matplotlib.pyplot.contourf.html)
2. <https://sashamaps.net/docs/resources/20-colors/>
3. <https://nipy.org/nibabel/>
4. [https://matplotlib.org/stable/api/\\_as\\_gen/matplotlib.pyplot.hist2d.html](https://matplotlib.org/stable/api/_as_gen/matplotlib.pyplot.hist2d.html)

## Part 1 matlab:





## CODE

```
% Uniform Distribution
arr_uniform = rand(1, 100);

% Gaussian Distribution
arr_normal = 17 * randn(1, 200) + 50;

figure;
boxplot(arr_uniform);
grid on;
title('Box plot of 100 random numbers uniformly distributed between 0-1', 'FontSize', 9);
ylabel('Distribution of Values');

figure;
boxplot(arr_normal);
grid on;
title('Box plot of 200 random numbers normally distributed with mean 50 and std 17', 'FontSize', 9);
ylabel('Distribution of Values');

figure;
histogram(arr_uniform, 20, 'EdgeColor', 'black', 'FaceColor', '#1AA7EC');
title('Histogram of 100 random numbers uniformly distributed between 0-1', 'FontSize', 9);
xlabel('Data between 0 and 1', 'FontSize', 14);
ylabel('Frequency', 'FontSize', 14);

figure;
histogram(arr_normal, 20, 'EdgeColor', 'black', 'FaceColor', '#1AA7EC');
title('Histogram of 200 random numbers with Gaussian distribution', 'FontSize', 9);
xlabel('Data values', 'FontSize', 14);
ylabel('Frequency', 'FontSize', 14);

% write file
file_path_uniform = 'uniform.bin';
fi = fopen(file_path_uniform, 'w');
fwrite(fi, arr_uniform, 'double');
fclose(fi);

% Read from the binary file
fi = fopen(file_path_uniform, 'r');
arr_uniform_file = fread(fi, 'double');
fclose(fi);

% Save the Gaussian array to a binary file
file_path_normal = 'normal.bin';
fi = fopen(file_path_normal, 'w');
fwrite(fi, arr_normal, 'double');
fclose(fi);

% Read from the binary file
fi = fopen(file_path_normal, 'r');
arr_normal_file = fread(fi, 'double');
fclose(fi);

% Sort the arrays
arr_uniform_sorted = sort(arr_uniform_file);
arr_normal_sorted = sort(arr_normal_file);

% Calculate the cumulative probabilities
cdf_uniform = (1:length(arr_uniform_sorted)) / length(arr_uniform_sorted);
```



```

cdf_normal = (1:length(arr_normal_sorted)) / length(arr_normal_sorted);

figure;
plot(arr_uniform_sorted, cdf_uniform, 'Color', '#1AA7EC');
grid on;
title('Cumulative Distribution Function for Uniform Distribution');
xlabel('Data between 0 and 1', 'FontSize', 14);
ylabel('Probability', 'FontSize', 14);

figure;
plot(arr_normal_sorted, cdf_normal, 'Color', '#1AA7EC');
grid on;
title('Cumulative Distribution Function for normal Distribution');
xlabel('Data between 1-100', 'FontSize', 14);
ylabel('Probability', 'FontSize', 14);

% Generate a 2D array for uniform distribution
arr_uniform_2d = rand(2, 5000);

% Generate a 2D array for normal distribution
arr_normal_2d = 0.5 + 0.13 * randn(2, 5000);

figure;
scatter(arr_uniform_2d(1,:), arr_uniform_2d(2,:), 'filled', 'Color', '#1AA7EC');
hold on;

scatter(arr_normal_2d(1,:), arr_normal_2d(2,:), "filled", 'Color', 'black');

xlabel('X, values between [0,1]');
ylabel('Y, values between [0,1]');
legend('Uniform random sampling', 'Gaussian random sampling');
title('Scatter plot of uniform and random sampling data');

hold off;

figure ;
histogram2(arr_uniform_2d(1,:), arr_uniform_2d(2,:), 100, 'FaceColor', 'flat', 'DisplayStyle', 'tile');
colorbar;
xlabel('X data');
ylabel('Y data');
title('2D uniform sampling histogram');

figure ;
histogram2(arr_normal_2d(1,:), arr_normal_2d(2,:), 100, 'FaceColor', 'flat', 'DisplayStyle', 'tile');
colorbar;
xlabel('X data');
ylabel('Y data');
title('2D normal sampling histogram');

figure;
h = histogram2(arr_uniform_2d(1,:), arr_uniform_2d(2,:), 100, 'DisplayStyle', 'tile', 'Visible', 'off');
contourf(h.XBinEdges(1:end-1), h.YBinEdges(1:end-1), h.Values, linspace(min(h.Values(:)), max(h.Values(:)), 10), 'LineStyle',
'none');
colorbar;
xlabel('X Bins data');
ylabel('Y Bins data');
title('2D uniform histogram contour');

figure;

```

```
h = histogram2(arr_normal_2d(1,:), arr_normal_2d(2,:), 100, 'DisplayStyle', 'tile', 'Visible', 'off');
contourf(h.XBinEdges(1:end-1), h.YBinEdges(1:end-1), h.Values', linspace(min(h.Values(:)), max(h.Values(:)), 10), 'LineStyle',
'none');
colorbar;
xlabel('X Bins data');
ylabel('Y Bins data');
title('2D normal histogram contour');
```