

#FOOTBALL PLAYER CLASSIFICATION ANALYSIS

#ABSTRACT

Understanding the foundations of data simulation and the various applications of data across industries is the aim of this study. The 19238 rows and 110 columns of the FIFA football players list of the year 2022 are used in various data analysis processes, such as simulating discrete and continuous random variables, which involve statistical analysis, visualisation, verification of the central limit theorem, outlier detection, and probability calculations. Furthermore, we simulate Markov Chains and Real Data Analysis, encompassing variance reduction techniques, factor analysis, recurrent events, ergodicity, joint distribution analysis, sensitivity analysis, and visualisation. We also compare different approaches to simulation.

Data on FIFA football players list is included with the target variable in the dataset, the overall score of the different players. It contains the following data: Attacking, Skills, Defense, Mentality, GK Skills, age, height, weight, calculation scores such as move, dribble, goalkeeping, club name, etc. The project's goal is to comprehend the philosophy behind simulating various data kinds under various circumstances in many disciplines.

#CHAPTER 1

#INTRODUCTION

The Data Set contains details on video game of FIFA football players for the year 2022. The target variable includes the age of the various players which could help to give the insights of how to choose the teams. The other variables like player attributes with statistics as Attacking, Skills, Defense, Mentality, GK Skills, player personal data like Nationality, Club, DateOfBirth, Wage, Salary, etc. seem to be potential predictive indicators that may correlate or relate to create the ideal team in the game. The data could be used to analyze the key drivers to select the team and build a model to predict the best team out of it. This dataset contains details on 19,238 football players including information on their demographics, player status, individual details, and a target variable indicating the age of the player so that it could be analysed how to select the player based on their overall score of all the given attributes. There are 110 feature variables on each player that could relate to selecting the team along with the binary target overall score. Performing some exploratory analysis on this data can help better understand the factors at play for player selecting and identify patterns that differentiate other players from creating a good team. These insights can then inform the development of a Good FIFA team in the videogame. This analysis aims to:

- Familiarize the reader with the data through distribution plots and summary statistics
- Identify correlations between the predictor variables
- Determine which variables have the strongest relationship with overall stats
- Outline relevant football players based on the available attributes.

#CHAPTER 2

#DATA DESCRIPTION

This dataset contains: Number of rows: 19238 Number of columns: 110 The columns include:

- RowNumber - likely just an index
- playerURL - unique ID for each player
- Short name - name of customer
- Long name - Complete name of the player
- Playerpositions - positions that the player can be able to play as an individual
- Overall - overall score of the players based upon the stats
- Potential - the potential of the player
- Age - age of player
- value - the overall value of the

player • wage - net amount that need to be given to the player in euros • height - heights of the different players • clubname - the name of the club that a player plays for • jersey number - this indicates the jersey number of the players • clubcountry - the details of the club that the player is belongs to which country • nationalteam - the national team the player plays for. This dataset contains details on 19,238 football players including information on their demographics, overall status, financial details, and a target variable indicating the overall score of the different variables included. 110 feature variables on each player could relate to selecting a team along with the binary target overall score. Performing some exploratory analysis on this data can help better understand the factors at play for selecting the team of players and identify patterns that differentiate best players for a team and the remaining ones. These insights can then inform the development of a predictive player selecting system for the video game.

The following sections will cover the key findings, including various plots and quantitative summaries of the data.

#DISCRIPTIVE ANALYSIS

Looking at the distribution of the target variable the average overall score of the players is 65. The highest overall score is 93. Next distributions of some key numeric variables:

AGE

Player age follows a somewhat normal distribution centered around 25 years old. Younger players appear more likely to be selected in the team and have more skills.

Passing

This describes about the passing ability scores of the various players which can be able to classify the players to make a better team.

Dribbling

This helps the reader to classify players based on the dribbling capacity of the individual player.

#CHAPTER 3

Mean:

The mean, which is the average of a collection of values, is a measure of central tendency in statistics. It is computed by taking the total number of values in a dataset and dividing the result by the sum of the individual values. The one representative number that represents the centre of the data is provided by the arithmetic mean, or \bar{x} . It is a basic idea in statistics that is used to explain the average or usual value within a set. In many fields, such as mathematics, economics, and data analysis, the mean is used extensively as a crucial central tendency indicator. By combining several values into a single representative measure, the mean can shed light on the general behaviour of a dataset.

Variance

A statistical metric called variance is used to quantify how widely or dispersed a group of values within a dataset are. It is a crucial metric for determining how different individual data points are from the mean. Each data point's divergence from the mean is squared in order to calculate the variance. The mean is then determined by averaging these squared deviations. (An increased

variance implies a higher level of dispersion within the sample, implying that the values deviate more from the average. On the other hand, a smaller variance denotes less variability and a closer proximity of the values to the mean. In order to evaluate the consistency and dependability of data, variance is a fundamental idea in statistics and an important

Standard Deviation

A statistical measure called standard deviation is used in conjunction with variance to provide a more comprehensible depiction of the distribution or dispersion of a collection of values within a dataset. Standard deviation is only the variance squared, whereas variance indicates the average of the squared deviations from the mean. Essentially, the standard deviation gives an indication of the average separation between each data point and the mean. Greater dispersion is implied by a bigger standard deviation, whereas a smaller standard deviation indicates that the values in the dataset are closely concentrated around the mean, indicating less variability. In statistics, standard deviation is a commonly used and straightforward metric that is often used to evaluate the risk, volatility, or variability of data in domains including finance, science, and

Mode

A dataset's mode is a statistical measure that indicates the value or values that appear the most frequently. The mode draws attention to the values that occur most frequently, as opposed to the mean and median, which concentrate on central tendency. A dataset may contain one mode (unimodal), multiple modes (multimodal), or none at all if every value occurs exactly once. When defining the essential characteristics of discrete or category data, like the most prevalent category within a set, the mode is especially helpful. It can be used to find ranges or intervals with the highest frequency in continuous data. The mode is frequently employed in several disciplines, including as statistics, sociology, and

Markov's Chain

A mathematical representation of a system or process that alternates between states in accordance with specific probabilistic criteria is called a Markov chain. The Markov property, which asserts that the system's future state depends solely on its current state and not on how it got there, is the essential component of a Markov chain. Stated differently, the system just remembers its current state and not its previous ones.

A set of states and a set of transition probabilities between states create a Markov chain. The probabilities are frequently grouped in a transition matrix, where the likelihood of changing from state I to state J is represented by the entry (i, j) . The concept that the system must be in one of the states following a transition is reflected in the matrix's rows, which add up to 1.

Markov chains are widely used in modelling physical processes such as particle motion, economic and biological systems, and several elements of machine learning and artificial intelligence, including reinforcement learning and natural language processing. They are very helpful for modelling random and uncertain systems because they make it possible to analyse and forecast how the system will behave in the future.

Continuous Random Variables

A basic idea in probability theory, continuous random variables are variables that can have an uncountably large number of values within a given range. X is the random variable; unlike their discrete counterparts, continuous random variables are described by probability density

functions (PDFs). The chance that the variable will fall within a given range is calculated by integrating the probability distribution function (PDF) over the given interval. The PDF indicates the probability that the variable will assume a specific value within the period. Notably, there is technically no chance that a continuous random variable will take exactly one value.

Temperature, weight, and other physical parameters are common instances of continuous random variables. Continuous distributions, like the normal distribution, are widely used in statistical analyses to model behaviour of the variables.

Discrete Random Variables

A key idea in probability theory are discrete random variables, which are variables that can only have discrete, distinct values with particular probabilities. Discrete random variables usually entail countable outcomes, as opposed to continuous random variables, which might assume an uncountably infinite number of values within a range. The chance of a discrete random variable taking any value is represented by the probability mass function (PMF), and the total of these probabilities over all possible values is 1. The results of a fair six-sided die or the number of heads in a sequence of coin flips are two examples of discrete random variables. A discrete random variable's cumulative distribution function (or CDF) gives the likelihood that the variable will be less than or equal to a given value.

Factor Analysis

In order to reduce the dimensionality of the data and reveal the underlying structure, factor analysis is a statistical approach used to identify latent factors that underlie a set of observed variables. This approach makes the assumption that fewer, so-called latent, unobservable factors have an impact on the measured variables. component loadings show the type and intensity of the association between each latent component and the observed variables. Eigenvalues, which show how much of the variance in the observed variables is explained by each factor, are the outcome of the analysis. It is possible to use rotation techniques to improve the results' interpretability. Factor analysis is especially useful in the social sciences and psychology, where researchers aim to find and understand the latent constructs that contributes to the observed correlations.

#CHAPTER 4

4.1 Simulation Data Analysis

```
import scipy.stats as st
import matplotlib.pyplot as plt
import numpy as np
import pandas as pd

mu, sigma, n = 0, 0.1, 5000
file_path = 'players_22.csv'

footballplayers_dataset = pd.read_csv(file_path, low_memory=False)

footballplayers_dataset =
footballplayers_dataset[footballplayers_dataset['overall'] != 0]
```

```
mean_value = np.mean(footballplayers_dataset['overall'])
```

```
print("Mean : ", mean_value)
```

```
footballplayers_dataset =
```

```
footballplayers_dataset.reset_index(drop=True)
```

```
Mean : 65.77218150631529
```

```
footballplayers_dataset
```

	sofifa_id	player_url \
0	158023	https://sofifa.com/player/158023/lionel-messi/...
1	188545	https://sofifa.com/player/188545/robert-lewand...
2	20801	https://sofifa.com/player/20801/c-ronaldo-dos-...
3	190871	https://sofifa.com/player/190871/neymar-da-sil...
4	192985	https://sofifa.com/player/192985/kevin-de-bruy...
...
19234	261962	https://sofifa.com/player/261962/defu-song/220002
19235	262040	https://sofifa.com/player/262040/caoimhin-port...
19236	262760	https://sofifa.com/player/262760/nathan-logue/...
19237	262820	https://sofifa.com/player/262820/luke-rudden/2...
19238	264540	https://sofifa.com/player/264540/emanuel-lalch...

	short_name	long_name \
0	L. Messi	Lionel Andrés Messi Cuccittini
1	R. Lewandowski	Robert Lewandowski
2	Cristiano Ronaldo	Cristiano Ronaldo dos Santos Aveiro
3	Neymar Jr	Neymar da Silva Santos Júnior
4	K. De Bruyne	Kevin De Bruyne
...
19234	Song Defu	宋德福
19235	C. Porter	Caoimhin Porter
19236	N. Logue	Nathan Logue-Cunningham
19237	L. Rudden	Luke Rudden
19238	E. Lalchhanchhuaha	Emanuel Lalchhanchhuaha

	player_positions	overall	potential	value_eur	wage_eur	age
...	\					
0	RW, ST, CF	93	93	78000000.0	320000.0	34
...						
1	ST	92	92	119500000.0	270000.0	32
...						
2	ST, LW	91	91	45000000.0	270000.0	36
...						
3	LW, CAM	91	91	129000000.0	270000.0	29
...						
4	CM, CAM	91	91	125500000.0	350000.0	30
...						
...

...						
19234	CDM	47	52	70000.0	1000.0	22
...						
19235	CM	47	59	110000.0	500.0	19
...						
19236	CM	47	55	100000.0	500.0	21
...						
19237	ST	47	60	110000.0	500.0	19
...						
19238	CAM	47	60	110000.0	500.0	19
...						

	lcb	cb	rcb	rb	gk	\
0	50+3	50+3	50+3	61+3	19+3	
1	60+3	60+3	60+3	61+3	19+3	
2	53+3	53+3	53+3	60+3	20+3	
3	50+3	50+3	50+3	62+3	20+3	
4	69+3	69+3	69+3	75+3	21+3	
...	
19234	46+2	46+2	46+2	48+2	15+2	
19235	44+2	44+2	44+2	48+2	14+2	
19236	45+2	45+2	45+2	47+2	12+2	
19237	26+2	26+2	26+2	32+2	15+2	
19238	41+2	41+2	41+2	45+2	16+2	

	player_face_url	\
0	https://cdn.sofifa.net/players/158/023/22_120.png	
1	https://cdn.sofifa.net/players/188/545/22_120.png	
2	https://cdn.sofifa.net/players/020/801/22_120.png	
3	https://cdn.sofifa.net/players/190/871/22_120.png	
4	https://cdn.sofifa.net/players/192/985/22_120.png	
...	...	
19234	https://cdn.sofifa.net/players/261/962/22_120.png	
19235	https://cdn.sofifa.net/players/262/040/22_120.png	
19236	https://cdn.sofifa.net/players/262/760/22_120.png	
19237	https://cdn.sofifa.net/players/262/820/22_120.png	
19238	https://cdn.sofifa.net/players/264/540/22_120.png	

	club_logo_url	\
0	https://cdn.sofifa.net/teams/73/60.png	
1	https://cdn.sofifa.net/teams/21/60.png	
2	https://cdn.sofifa.net/teams/11/60.png	
3	https://cdn.sofifa.net/teams/73/60.png	
4	https://cdn.sofifa.net/teams/10/60.png	
...	...	
19234	https://cdn.sofifa.net/teams/112541/60.png	
19235	https://cdn.sofifa.net/teams/445/60.png	
19236	https://cdn.sofifa.net/teams/111131/60.png	
19237	https://cdn.sofifa.net/teams/111131/60.png	
19238	https://cdn.sofifa.net/teams/113040/60.png	

```

club_flag_url \
0      https://cdn.sofifa.net/flags/fr.png
1      https://cdn.sofifa.net/flags/de.png
2      https://cdn.sofifa.net/flags/gb-eng.png
3      https://cdn.sofifa.net/flags/fr.png
4      https://cdn.sofifa.net/flags/gb-eng.png
...
19234  https://cdn.sofifa.net/flags/cn.png
19235  https://cdn.sofifa.net/flags/ie.png
19236  https://cdn.sofifa.net/flags/ie.png
19237  https://cdn.sofifa.net/flags/ie.png
19238  https://cdn.sofifa.net/flags/in.png

```

```

nation_logo_url \
0      https://cdn.sofifa.net/teams/1369/60.png
1      https://cdn.sofifa.net/teams/1353/60.png
2      https://cdn.sofifa.net/teams/1354/60.png
3      NaN
4      https://cdn.sofifa.net/teams/1325/60.png
...
19234  NaN
19235  NaN
19236  NaN
19237  NaN
19238  NaN

```

```

nation_flag_url
0      https://cdn.sofifa.net/flags/ar.png
1      https://cdn.sofifa.net/flags/pl.png
2      https://cdn.sofifa.net/flags/pt.png
3      https://cdn.sofifa.net/flags/br.png
4      https://cdn.sofifa.net/flags/be.png
...
19234  https://cdn.sofifa.net/flags/cn.png
19235  https://cdn.sofifa.net/flags/ie.png
19236  https://cdn.sofifa.net/flags/ie.png
19237  https://cdn.sofifa.net/flags/ie.png
19238  https://cdn.sofifa.net/flags/in.png

```

```
[19239 rows x 110 columns]
```

```
overall_score_data = footballplayers_dataset['overall']
```

```
overall_score_data
```

```

0      93
1      92
2      91
3      91

```

```

4          91
19234      ..
19235      47
19236      47
19237      47
19238      47
Name: overall, Length: 19239, dtype: int64

```

Determination of my Dataset

```

import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns

file_path = 'players_22.csv'

footballplayers_dataset = pd.read_csv(file_path, low_memory=False)

print(footballplayers_dataset.info())
print(footballplayers_dataset.describe())

plt.figure(figsize=(8, 6))
sns.histplot(footballplayers_dataset['overall'], kde=True, bins=30,
color='skyblue')
plt.title('Distribution of Overall Score')
plt.xlabel('Overall Score')
plt.ylabel('Frequency')
plt.show()

```

```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 19239 entries, 0 to 19238
Columns: 110 entries, sofifa_id to nation_flag_url
dtypes: float64(16), int64(44), object(50)
memory usage: 16.1+ MB
None

```

	sofifa_id	overall	potential	value_eur
wage_eur \				
count	19239.000000	19239.000000	19239.000000	1.916500e+04
mean	231468.086959	65.772182	71.079370	2.850452e+06
std	27039.717497	6.880232	6.086213	7.613700e+06
min	41.000000	47.000000	49.000000	9.000000e+03
25%	214413.500000	61.000000	67.000000	4.750000e+05
50%	236543.000000	66.000000	71.000000	9.750000e+05


```

3000.000000
75%    253532.500000    70.000000    75.000000    2.000000e+06
8000.000000
max     264640.000000    93.000000    95.000000    1.940000e+08
350000.000000

```

```

          age      height_cm      weight_kg      club_team_id
league_level \
count  19239.000000  19239.000000  19239.000000  19178.000000
19178.000000
mean    25.210822    181.299704    74.943032    50580.498123
1.354364
std      4.748235      6.863179      7.069434    54401.868535
0.747865
min     16.000000    155.000000    49.000000      1.000000
1.000000
25%     21.000000    176.000000    70.000000    479.000000
1.000000
50%     25.000000    181.000000    75.000000    1938.000000
1.000000
75%     29.000000    186.000000    80.000000   111139.000000
1.000000
max     54.000000    206.000000   110.000000  115820.000000
5.000000

```

```

...      mentality_composure      defending_marking_awareness \
count    ...      19239.000000      19239.000000
mean      ...      57.929830      46.601746
std        ...      12.159326      20.200807
min        ...      12.000000      4.000000
25%        ...      50.000000      29.000000
50%        ...      59.000000      52.000000
75%        ...      66.000000      63.000000
max        ...      96.000000      93.000000

```

```

          defending_standing_tackle      defending_sliding_tackle \
count      19239.000000      19239.000000
mean        48.045584      45.906700
std         21.232718      20.755683
min          5.000000      5.000000
25%         28.000000      25.000000
50%         56.000000      53.000000
75%         65.000000      63.000000
max         93.000000      92.000000

```

```

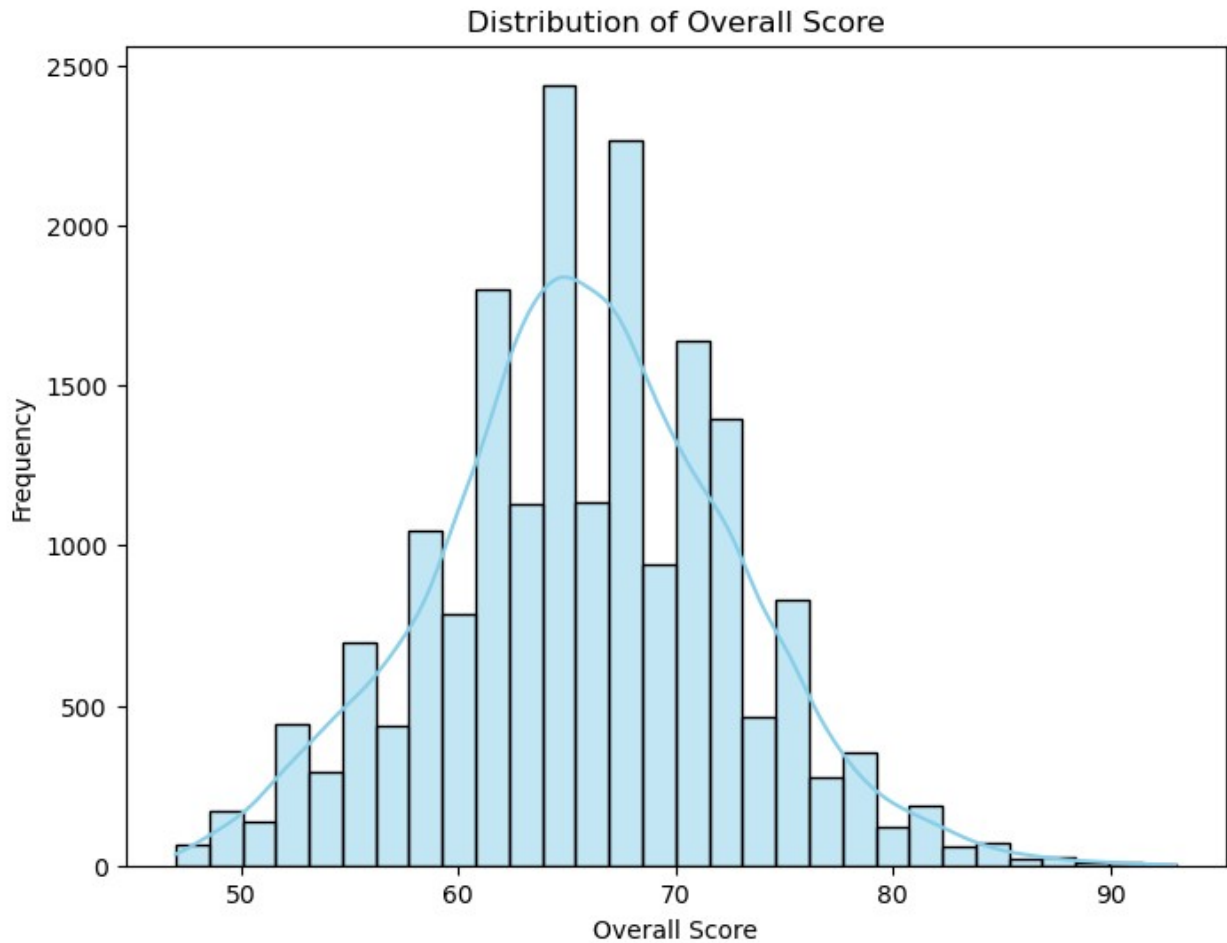
          goalkeeping_diving      goalkeeping_handling
goalkeeping_kicking \
count      19239.000000      19239.000000      19239.000000
mean        16.406102      16.192474      16.055356

```

std	17.574028	16.839528	16.564554
min	2.000000	2.000000	2.000000
25%	8.000000	8.000000	8.000000
50%	11.000000	11.000000	11.000000
75%	14.000000	14.000000	14.000000
max	91.000000	92.000000	93.000000

	goalkeeping_positioning	goalkeeping_reflexes
goalkeeping_speed		
count	19239.000000	19239.000000
2132.000000		
mean	16.229274	16.491814
36.439962		
std	17.059779	17.884833
10.751563		
min	2.000000	2.000000
15.000000		
25%	8.000000	8.000000
27.000000		
50%	11.000000	11.000000
36.000000		
75%	14.000000	14.000000
45.000000		
max	92.000000	90.000000
65.000000		

[8 rows x 60 columns]



#4.1.1 Simulating Continuous Random Variables:

```
import pandas as pd

file_path = 'players_22.csv'
footballplayers_dataset = pd.read_csv(file_path, low_memory=False)
print(footballplayers_dataset.columns)

overall_score_data = footballplayers_dataset['overall']
Index(['sofifa_id', 'player_url', 'short_name', 'long_name',
       'player_positions', 'overall', 'potential', 'value_eur',
       'wage_eur',
       'age',
       'lcb', 'cb', 'rcb', 'rb', 'gk', 'player_face_url',
       'club_logo_url',
```

```
    'club_flag_url', 'nation_logo_url', 'nation_flag_url'],  
    dtype='object', length=110)
```

```
import numpy as np  
import matplotlib.pyplot as plt
```

```
mu = np.mean(overall_score_data)  
sigma = np.std(overall_score_data)
```

```
simulated_data = np.random.normal(mu, sigma, 1000)
```

```
print(simulated_data)  
plt.hist(simulated_data, bins=30, density=True, alpha=0.5,  
color='blue')  
plt.title('Simulated Normal Distribution based on Overall Score')  
plt.xlabel('Values')  
plt.ylabel('Frequency')  
plt.show()
```

```
[55.29361405 59.05179156 70.02545978 62.19444731 73.87536567  
67.76295772  
74.67217963 55.91467883 67.09256339 73.47766401 63.74562925  
64.82045096  
78.41646601 67.98031297 69.19556199 76.89565872 71.31097979  
67.89957706  
76.62308382 54.74527096 74.01374508 60.32686007 67.18109587  
66.83904487  
63.07645937 62.37332041 60.98948895 68.11283628 65.64939064  
79.80996582  
74.92191249 67.78777941 63.11635229 66.61083984 54.30095424  
67.11215192  
64.2605928 52.26852616 65.13756169 69.08274799 49.85417225  
52.84446395  
68.38956295 73.12795487 68.18174602 60.8541042 73.89376403  
67.25319152  
73.18746462 73.35743658 61.29969212 58.50926546 76.21585659  
70.58748763  
62.59623219 73.50939077 62.51948265 66.54484769 55.38196134  
77.36784597  
75.22169877 62.96388375 74.90317442 71.62843763 68.56676241  
76.01518779  
72.88025395 68.60998221 66.8690315 62.6269529 75.44307124  
56.87089174  
80.71545032 59.58774292 74.07385091 61.31777985 76.23552183  
68.97880177  
66.41719989 68.14835499 71.89601722 64.6539475 52.72004374  
66.43059989  
70.94773735 68.65936607 62.2441371 70.36980599 58.22288047  
68.43323327
```

75.62012566 62.70272006 60.42239171 58.47275118 55.04976133
66.59702864
57.40350905 61.98234827 67.77775351 49.9799661 66.19539641
71.86255276
73.52339015 64.54797949 78.24006665 58.17988518 76.47867588
68.87729873
51.06347021 70.30119663 61.59068898 68.49638155 65.10748281
65.61343404
78.40550564 57.36506645 63.6304838 64.94688275 64.90596491
67.68202978
69.34694768 73.63940671 52.63465408 62.85397657 71.64608985
76.21459581
70.50456904 64.75447023 66.93759542 56.58760805 63.9601269
70.8777257
63.56983924 79.86729007 62.1681926 58.19945475 66.6739589
67.99491729
66.29749399 61.40778124 69.69814717 60.88008087 57.10789997
49.92506379
60.75372394 78.70053125 72.24114512 69.3758243 67.07630421
62.33335706
69.82519895 64.30755139 76.60252633 72.65833518 58.39085429
60.71638644
70.35165367 56.56911852 70.92183787 61.59364945 76.20356432
54.01936974
68.2884446 64.01838336 70.31542194 74.70341484 75.69441434
68.84538089
71.20616473 66.48230868 76.12811008 62.48373991 60.42276623
65.49559426
63.69772885 84.3462451 60.57068816 62.75155855 77.35958408
74.88846068
61.37892308 59.65012367 57.27444495 73.48330207 73.54956066
70.36760445
62.8082104 76.42375622 58.72271305 62.46507021 58.31562593
60.8146715
74.3943214 66.43646103 60.63688635 68.08714604 69.89893683
69.89594501
63.45420461 66.32484666 73.12255372 62.47297312 74.56421069
70.03622476
63.87691515 74.39930264 78.28396284 85.96567656 63.53245276
64.46272269
59.30439096 71.53027731 71.92208675 61.20398671 69.44899582
62.12729896
76.29236948 64.86263803 58.39645983 68.4173881 61.07509306
56.41737567
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70.92608725
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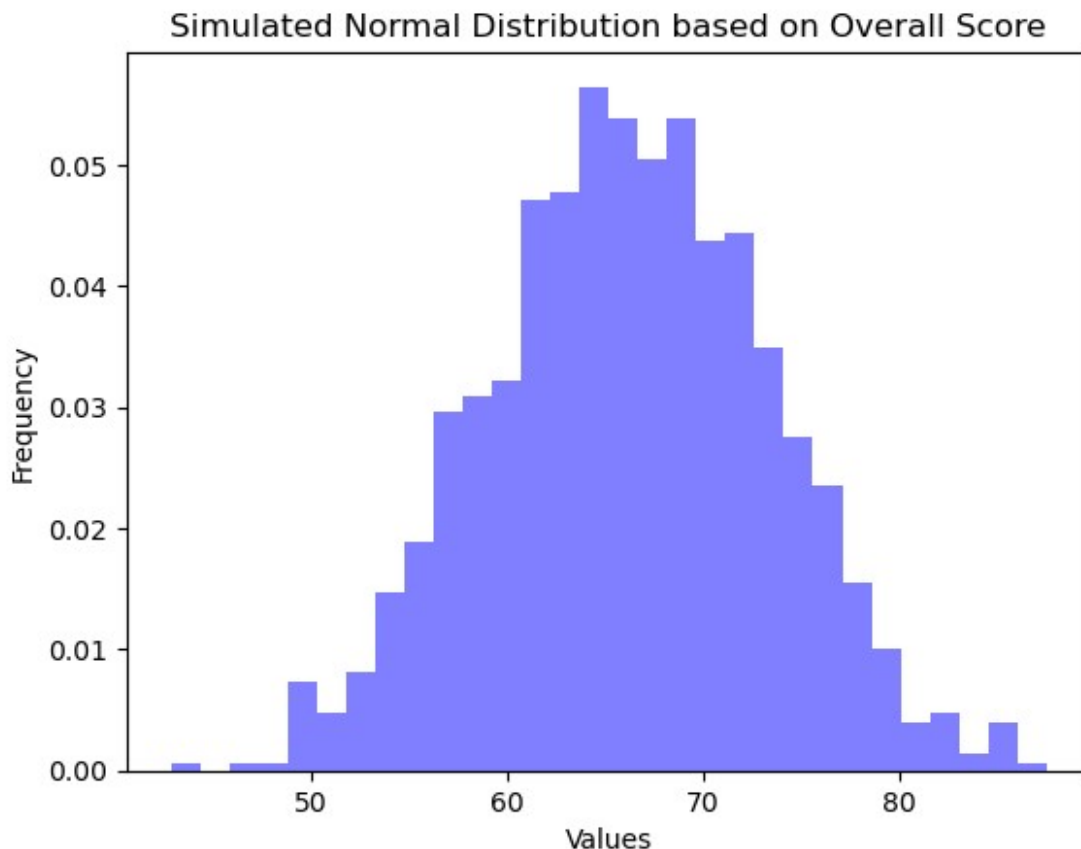
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77.71987014				
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59.9793475
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54.58094659
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67.00688503
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66.5354218
70.2431282 61.33674401 65.82972705 65.17398743 49.90533124
68.39260438
65.73209276 67.86333739 63.36642134 59.24257254 75.68391866
68.38025587

```

59.23501962 70.57547912 53.63095813 57.61697826 70.80186671
74.80100896
63.56818994 66.09217637 68.10313134 66.2908016 65.86100005
65.78251895
53.34331697 77.48310465 71.07948392 68.37281082 82.233119
69.97148605
64.13635933 68.32304816 85.68535724 60.07653292 71.24815483
77.03600869
63.8740144 68.87602047 75.39163731 60.50203332]

```



Statistical Analysis

```

import scipy.stats as st
mean = np.mean(overall_score_data)
median = np.median(overall_score_data)
std_deviation = np.std(overall_score_data)
variance = np.var(overall_score_data)
quantiles = np.percentile(overall_score_data, [25, 50, 75])
kurt = st.kurtosis(overall_score_data, axis=0, bias=True)

print(f"Mean: {mean}")
print(f"Median: {median}")
print(f"Standard Deviation: {std_deviation}")

```

```
print(f"Variance: {variance}")
print(f"Quantiles (25th, 50th, 75th percentiles): {quantiles}")
print(f"Kurtosis: {kurt}")
```

Mean: 65.77218150631529

Median: 66.0

Standard Deviation: 6.880052695049176

Variance: 47.33512508665343

Quantiles (25th, 50th, 75th percentiles): [61. 66. 70.]

Kurtosis: 0.09066867096156184

Visualization

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt

overall_score_data = footballplayers_dataset['overall']

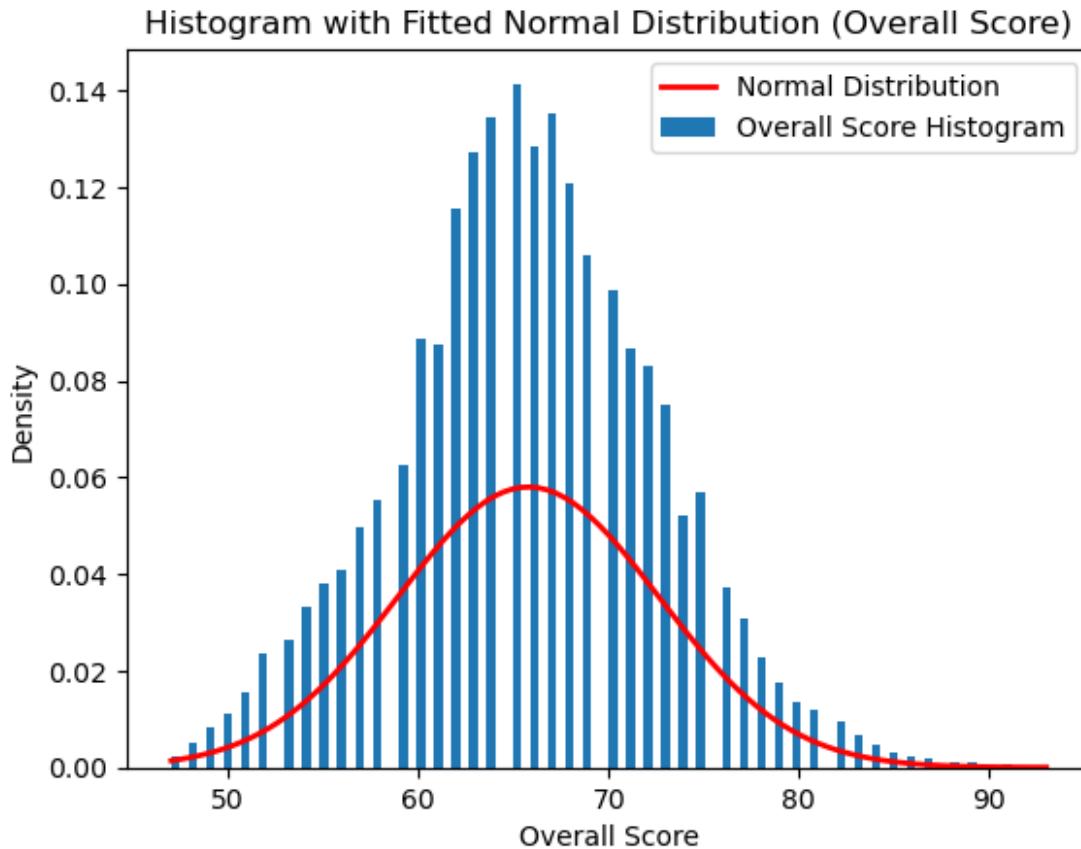
mu = np.mean(overall_score_data)
sigma = np.std(overall_score_data)

count, bins, ignored = plt.hist(overall_score_data, 100, density=True)

plt.plot(bins, 1/(sigma * np.sqrt(2 * np.pi)) * np.exp(-(bins - mu)**2 / (2 * sigma**2)),
         linewidth=2, color='red')

plt.title('Histogram with Fitted Normal Distribution (Overall Score)')
plt.xlabel('Overall Score')
plt.ylabel('Density')
plt.legend(['Normal Distribution', 'Overall Score Histogram'])

plt.show()
```



Central Limit Theorem Verification

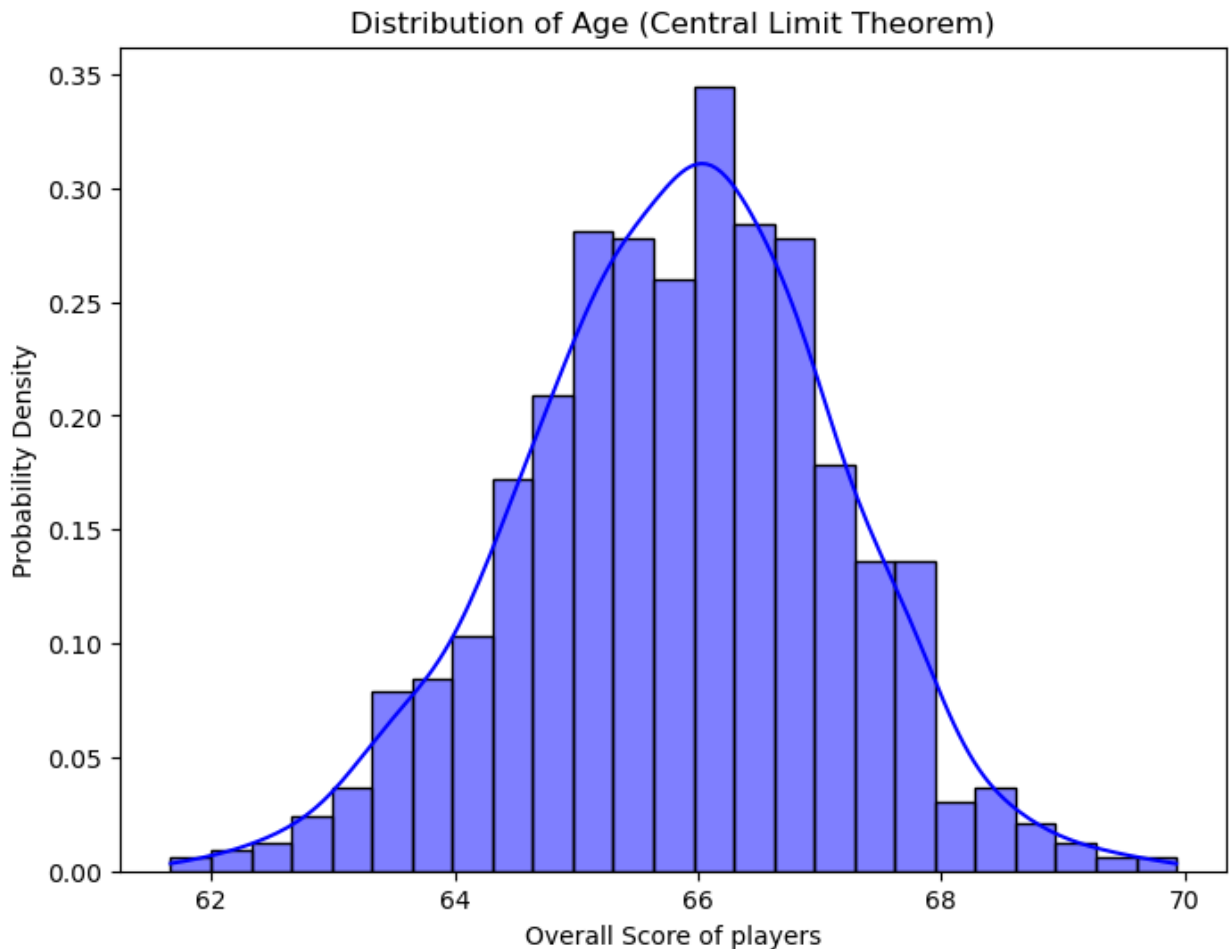
```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

overall_score_data = footballplayers_dataset['overall']

def calculate_sample_means(sample_size, no_of_sample_means):
    mean_list = []
    for i in range(no_of_sample_means):
        sample = np.random.choice(overall_score_data, sample_size)
        sample_mean = np.mean(sample)
        mean_list.append(sample_mean)
    return mean_list

sample_means = calculate_sample_means(sample_size=30,
no_of_sample_means=1000)
```

```
plt.figure(figsize=(8, 6))
sns.histplot(sample_means, color='blue', kde=True, stat="density")
plt.xlabel('Overall Score of players')
plt.ylabel('Probability Density')
plt.title('Distribution of Age (Central Limit Theorem)')
plt.show()
```



```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

overall_score_data = footballplayers_dataset['overall']

def calculate_sample_means(sample_size, no_of_sample_means):
    mean_list = []
    for i in range(no_of_sample_means):
        sample = np.random.choice(overall_score_data, sample_size)
        sample_mean = np.mean(sample)
```

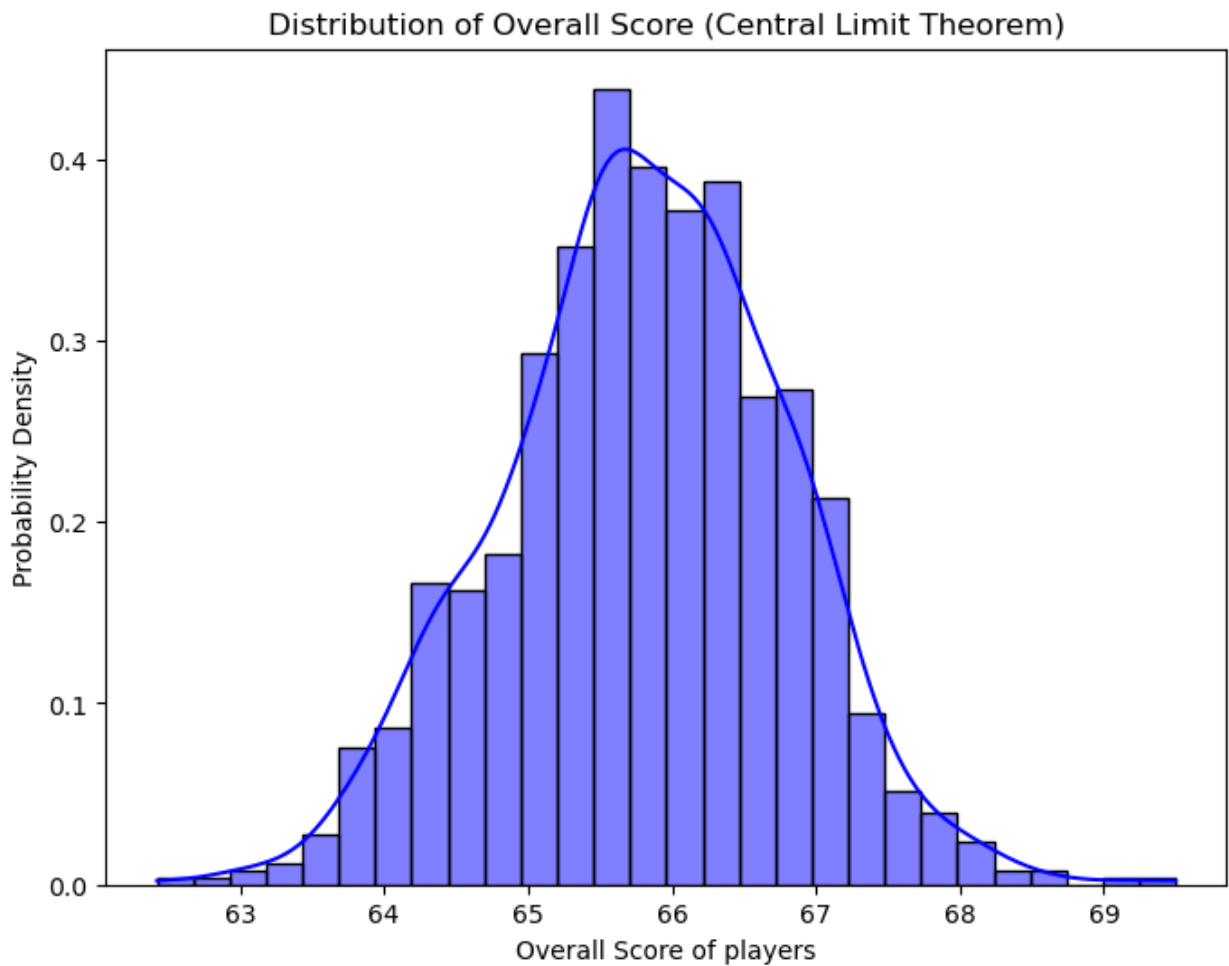
```

    mean_list.append(sample_mean)
    return mean_list

sample_means = calculate_sample_means(sample_size=50,
no_of_sample_means=1000)

plt.figure(figsize=(8, 6))
sns.histplot(sample_means, color='blue', kde=True, stat="density")
plt.xlabel('Overall Score of players')
plt.ylabel('Probability Density')
plt.title('Distribution of Overall Score (Central Limit Theorem)')
plt.show()

```



Outlier Detection

```

import pandas as pd
import numpy as np
import seaborn as sns

e_path = 'players_22.csv'

```

```

footballplayers_dataset = pd.read_csv(file_path)

column_of_interest = 'overall'
data_column = footballplayers_dataset[column_of_interest]

sns.boxplot(data_column)

q1 = np.quantile(data_column, 0.25)
q3 = np.quantile(data_column, 0.75)
IQR = q3 - q1
outliers_iqr = data_column[((data_column < (q1 - 1.5 * IQR)) |
(data_column > (q3 + 1.5 * IQR)))]

mean = np.mean(data_column)
std_dev = np.std(data_column)
z_score = (data_column - mean) / std_dev
outliers_z_score = data_column[(z_score > 2.73) | (z_score < -2.73)]

print("Outliers using IQR method:")
print(outliers_iqr)

print("\nOutliers using z-scores method:")
print(outliers_z_score)

```

```

Outliers using IQR method:
0      93
1      92
2      91
3      91
4      91
..
19234   47
19235   47
19236   47
19237   47
19238   47
Name: overall, Length: 159, dtype: int64

```

```

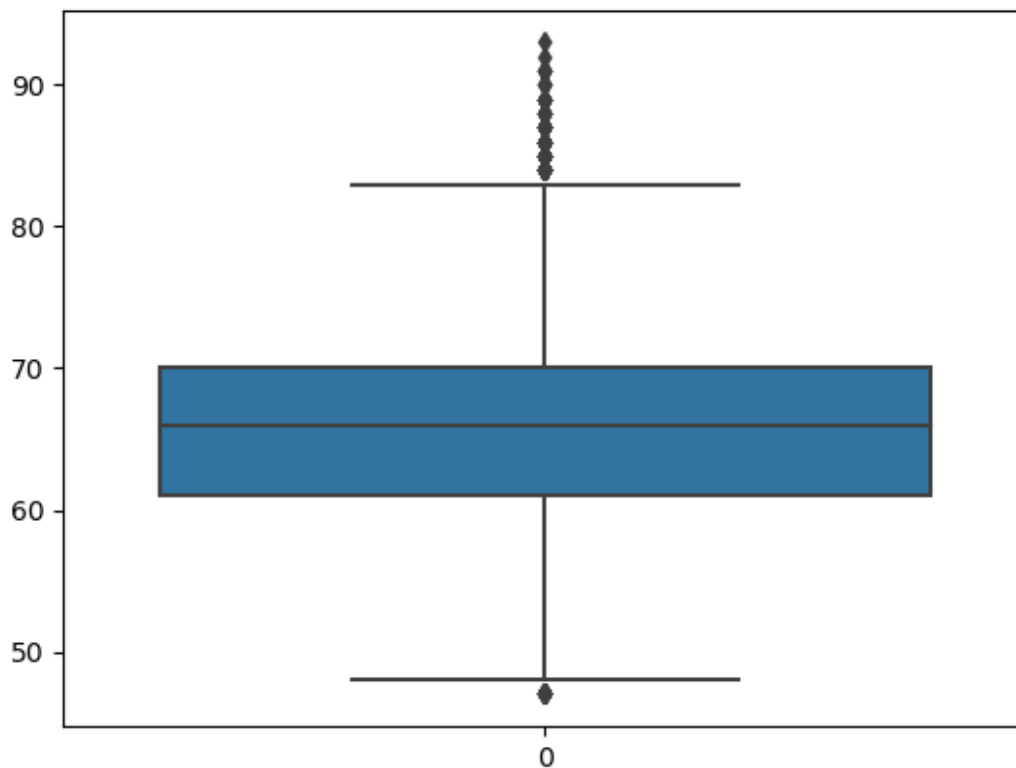
Outliers using z-scores method:
0      93
1      92
2      91
3      91
4      91
..
92     85
93     85
94     85
95     85

```


96 85

Name: overall, Length: 97, dtype: int64

```
/var/folders/5y/dp3cyz5s2vsbqcjxcnzd62900000gn/T/  
ipykernel_20279/1502119058.py:6: DtypeWarning: Columns (25,108) have  
mixed types. Specify dtype option on import or set low_memory=False.  
footballplayers_dataset = pd.read_csv(file_path)
```



Probability Calculations

```
import pandas as pd  
import numpy as np  
import seaborn as sns  
import scipy.stats as st  
  
file_path = 'players_22.csv'  
footballplayers_dataset = pd.read_csv(file_path, low_memory=False)  
  
column_of_interest = 'overall'  
data_column = footballplayers_dataset[column_of_interest]  
  
mu = np.mean(data_column)  
sigma = np.std(data_column)  
  
lower_bound = -0.2
```

```

upper_bound = 0.1
probability = st.norm.cdf((upper_bound - mu) / sigma) -
st.norm.cdf((lower_bound - mu) / sigma)

print(f"The probability of  $-0.2 < X < 0.1$  is: {probability:.4f}")
print(mu)
print(sigma)

```

The probability of $-0.2 < X < 0.1$ is: 0.0000
65.77218150631529
6.880052695049176

4.1.2 Simulating from Discrete Distributions:

```

import pandas as pd
import scipy.stats as st

file_path = 'players_22.csv'
footballplayers_dataset = pd.read_csv(file_path, low_memory=False)

column_of_interest = 'overall'
data_column = footballplayers_dataset[column_of_interest]

mu = data_column.mean()

data_discrete = st.poisson.rvs(mu, size=1000)

print(data_discrete)

```

[59 68 64 62 70 53 61 61 74 65 70 65 56 61 70 76 67 74 75 61 64 70 86
56
74 76 67 51 71 64 61 68 75 59 66 72 74 60 59 68 73 78 67 62 66 64 50
62
57 59 72 81 70 59 75 75 57 69 63 60 63 62 64 61 71 69 57 70 54 63 60
59
55 72 59 64 64 77 61 53 69 82 66 56 53 63 54 65 62 55 58 57 75 89 67
66
74 66 72 80 58 53 76 69 57 77 70 63 63 74 60 67 60 76 63 63 70 68 66
76
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61
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68
65 64 54 65 78 59 77 62 74 64 69 75 78 79 64 78 59 69 62 73 61 68 65
57
78 67 60 64 63 70 75 69 64 64 74 64 88 68 61 81 65 54 89 83 67 57 57
76

63 51 72 71 68 70 61 62 61 58 64 64 57 60 67 39 59 66 67 74 52 63 64
62
69 60 68 66 77 56 63 62 63 64 71 66 58 50 79 73 85 64 66 67 65 73 64
63
65 79 69 71 73 69 68 59 69 81 70 66 64 55 61 67 59 72 68 62 63 63 62
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69
80 66 62 54 65 76 62 58 67 60 57 66 63 73 72 70 62 63 49 68 67 61 51
59
79 83 62 80 67 63 61 68 53 63 80 69 53 73 63 63 68 66 59 69 66 67 78
62
62 67 59 70 68 49 67 59 63 61 80 61 51 70 64 61 74 73 76 62 71 69 67
41
54 68 74 61 66 57 77 61 67 68 84 66 54 65 65 61 50 61 54 70 70 67 52
73
62 55 76 71 71 65 54 66 82 58 60 70 66 70 61 52 62 65 70 77 63 67 62
59
83 77 63 75 61 63 53 57 76 63 61 57 73 59 59 74 64 61 80 73 56 58 60
66
57 63 56 70 56 63 76 60 70 83 80 63 62 87 78 60 58 71 58 71 68 59 71
53
66 78 70 61 75 49 60 74 63 80 67 57 63 70 55 65 62 65 77 55 56 62 59
67
63 73 60 71 64 67 65 65 55 70 72 77 61 59 71 70 66 51 64 51 61 67 57
70
59 66 65 71 69 69 66 65 74 55 71 72 73 50 54 53 42 74 60 67 57 58 72
69
66 59 56 82 69 53 67 52 64 59 77 60 65 59 75 67 64 62 73 77 62 75 76

```

66
66 65 63 79 77 72 59 71 64 77 74 79 59 64 80 61 59 73 76 65 73 54 71
66
58 69 64 61 49 65 72 61 65 69 67 61 61 74 63 72 54 70 68 53 66 74 61
72
61 74 61 52 66 74 52 61 60 58 58 61 64 42 72 61 62 58 68 57 71 69 62
68
75 64 72 59 72 75 68 59 62 70 69 61 68 79 64 63 72 64 72 43 75 61 68
68
71 60 77 61 68 82 77 78 77 68 65 72 54 67 68 50 68 75 78 59 66 58 79
74
64 62 55 70 66 80 82 66 73 59 65 74 71 59 58 76 62 74 82 70 59 71 60
59
78 62 64 64 73 66 63 85 66 58 67 76 60 61 55 78]

```

Statistical Analysis:

```

import numpy as np
import seaborn as sns
import scipy.stats as st

data_column = np.random.randn(100)

mean = np.mean(data_column)
standard_deviation = np.std(data_column)
variance = np.var(data_column)
quantile = np.quantile(data_column, [0.25, 0.5, 0.75])

mode_result = st.mode(data_column)
mode = mode_result.mode[0]

skewness = st.skew(data_column)
kurtosis = st.kurtosis(data_column)

print("Mean:", mean)
print("Standard Deviation:", standard_deviation)
print("Variance:", variance)
print("First Quantile:", quantile[0])
print("Median (Second Quantile):", quantile[1])
print("Third Quantile:", quantile[2])
print("Mode:", mode)
print("Skewness:", skewness)
print("Kurtosis:", kurtosis)

Mean: -0.05696541719871014
Standard Deviation: 0.9290249996835663
Variance: 0.8630874500370505
First Quantile: -0.7271830721155583
Median (Second Quantile): -0.03113676017848229
Third Quantile: 0.5835395117876068

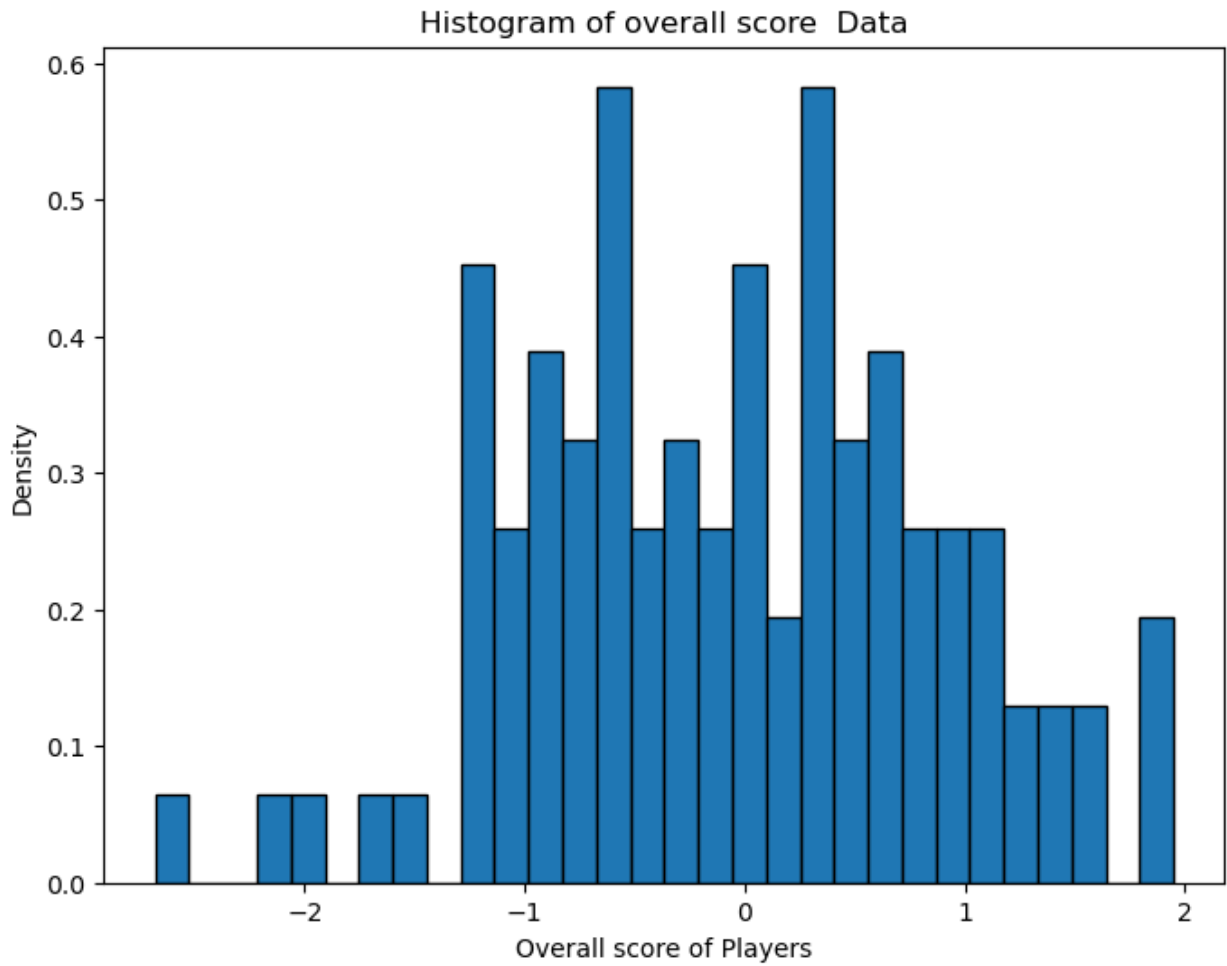
```

```
Mode: -2.6811691124076287
Skewness: -0.08004358007075359
Kurtosis: -0.2402268748693781
```

```
/var/folders/5y/dp3cyz5s2vsbqcjxcnzd62900000gn/T/
ipykernel_20279/950199653.py:12: FutureWarning: Unlike other reduction
functions (e.g. `skew`, `kurtosis`), the default behavior of `mode`
typically preserves the axis it acts along. In SciPy 1.11.0, this
behavior will change: the default value of `keepdims` will become
False, the `axis` over which the statistic is taken will be
eliminated, and the value None will no longer be accepted. Set
`keepdims` to True or False to avoid this warning.
    mode_result = st.mode(data_column)
```

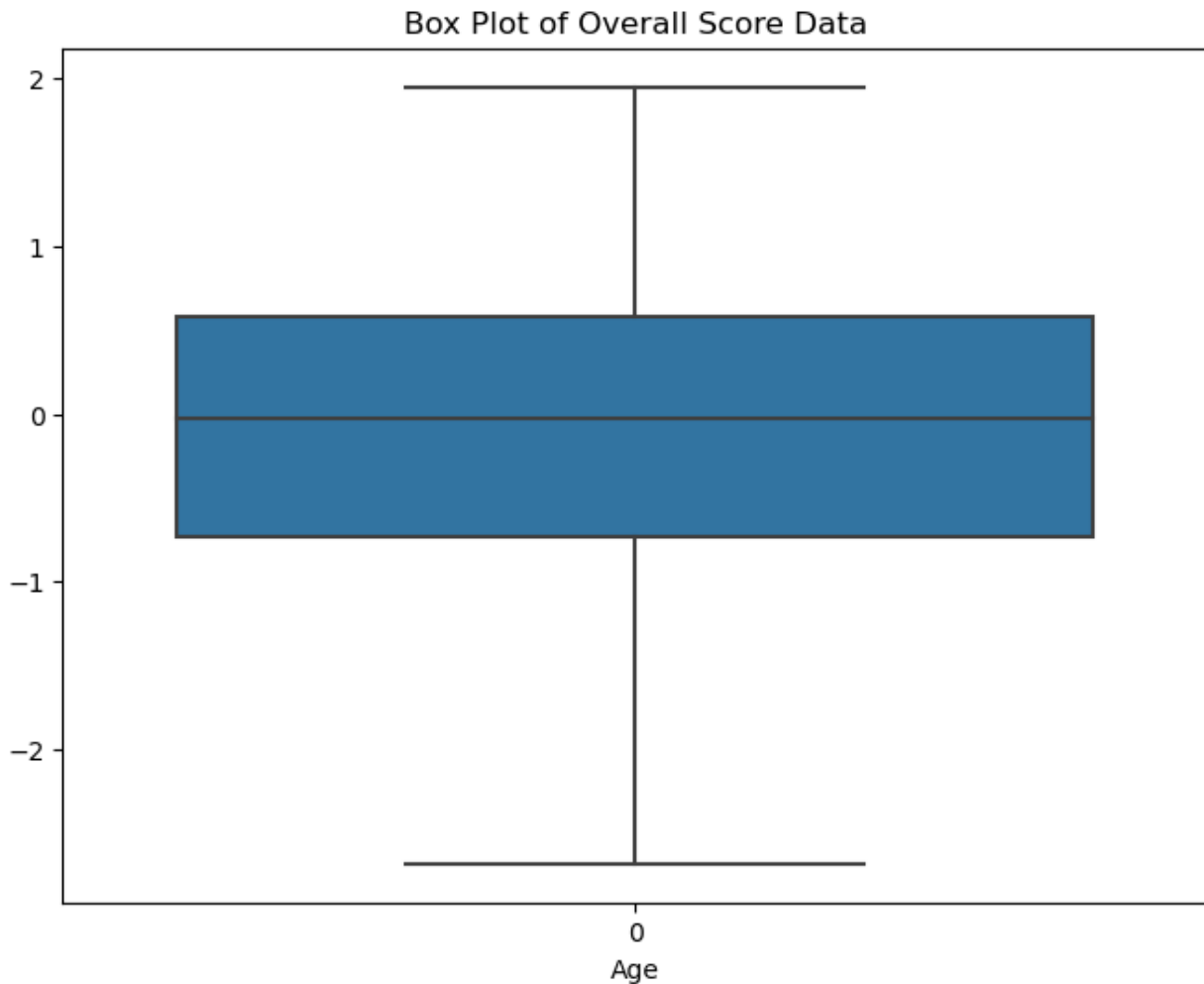
Visualization:

```
import matplotlib.pyplot as plt
plt.figure(figsize=(8, 6))
plt.hist(data_column, bins=30, density=True, edgecolor='black')
plt.title('Histogram of overall score Data')
plt.xlabel('Overall score of Players')
plt.ylabel('Density')
plt.show()
```



Box Plot of the Data

```
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
plt.figure(figsize=(8, 6))
sns.boxplot(data_column)
plt.title('Box Plot of Overall Score Data')
plt.xlabel('Age')
plt.show()
```



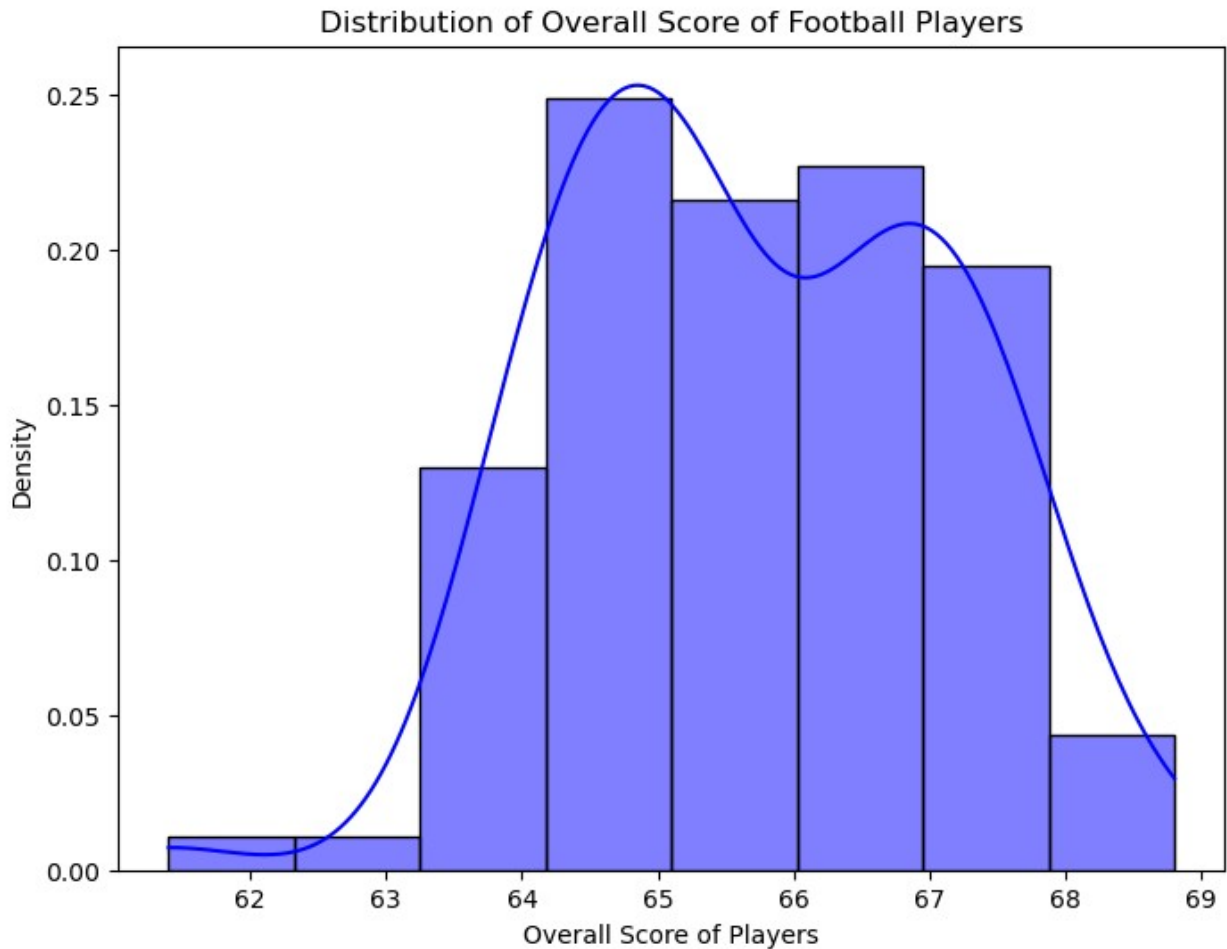
Central Limit Verification

```
import random
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np

data_column = footballplayers_dataset['overall']
mean_list = []
def calc_sample_mean(sample_size, no_of_sample_means):
    for i in range(no_of_sample_means):
        sample = random.sample(list(data_column), sample_size)
        sample_mean = np.mean(sample)
        mean_list.append(sample_mean)
    return mean_list
mean_samples = calc_sample_mean(sample_size=20,
no_of_sample_means=100)

# Visualize sample means distribution
```

```
plt.figure(figsize=(8, 6))
sns.histplot(mean_samples, color='blue', kde=True, stat="density")
plt.xlabel('Overall Score of Players')
plt.title('Distribution of Overall Score of Football Players')
plt.show()
```



Outliers detection

```
import seaborn as sns
import numpy as np

data_column = footballplayers_dataset['overall']

sns.boxplot(data_column)
plt.title('Boxplot of Overall Score of Players')
plt.ylabel('Overall Score')
plt.show()

q1 = np.quantile(data_column, 0.25)
q3 = np.quantile(data_column, 0.75)
```



```

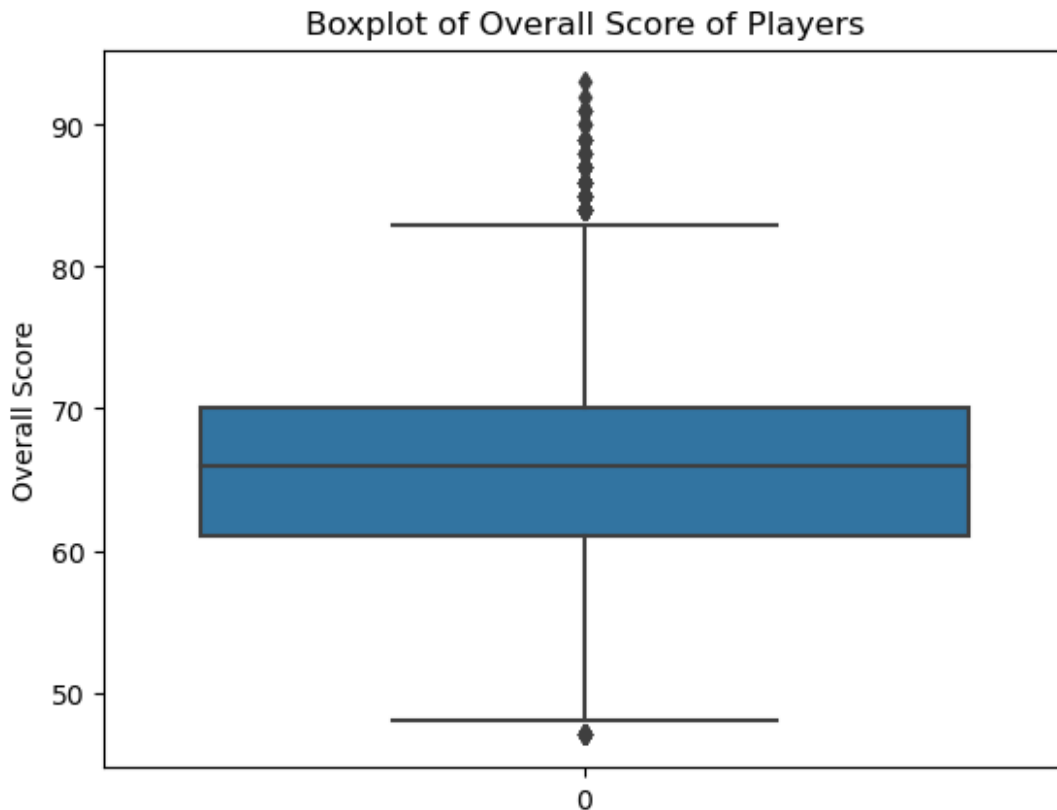
IQR = q3 - q1
outliers = data_column[((data_column < (q1 - 1.5 * IQR)) |
(data_column > (q3 + 1.5 * IQR)))]

print("Outliers using IQR method:")
print(outliers)

mean = np.mean(data_column)
standard_deviation = np.std(data_column)
upper_limit = mean + 2.73 * standard_deviation
lower_limit = mean - 2.73 * standard_deviation
outliers_z_scores = data_column[((data_column > upper_limit) |
(data_column < lower_limit) )]

print("\nOutliers using z-scores method:")
print(outliers_z_scores)

```



```

Outliers using IQR method:
0      93
1      92
2      91
3      91
4      91
..

```

```
19234    47
19235    47
19236    47
19237    47
19238    47
Name: overall, Length: 159, dtype: int64
```

Outliers using z-scores method:

```
0      93
1      92
2      91
3      91
4      91
..
92     85
93     85
94     85
95     85
96     85
Name: overall, Length: 97, dtype: int64
```

Probability calculation

```
import pandas as pd
import numpy as np
import seaborn as sns
import scipy.stats as st

file_path = 'players_22.csv'
housing_data = pd.read_csv(file_path, low_memory = False)

column_of_interest = 'overall'
data_column = housing_data[column_of_interest]
mu = np.mean(data_column)
sigma = np.std(data_column)

lower_bound = -0.2
upper_bound = 0.1
probability = st.norm.cdf((upper_bound - mu) / sigma) -
st.norm.cdf((lower_bound - mu) / sigma)
print(f"The probability of  $-0.2 < X < 0.1$  is: {probability:.4f}")
print(mu)
print(sigma)

The probability of  $-0.2 < X < 0.1$  is: 0.0000
65.77218150631529
6.880052695049176
```

#4.1.3 MARKOV CHAIN

- Transition Matrix Simulation

Markov Chains:

* Transition Matrix Simulation: Using a given transition matrix, simulate a basic Markov chain. It is possible to model state transitions and determine the probability of each state after a predetermined number of steps.

Code Explanation:

->The probabilities of changing states in a single step are represented by the transition matrix in a Markov chain. Applying the transition matrix again will yield the state probabilities after a given number of steps.

->The simulate_markov_chain function updates the state vector for the requested number of steps based on the transition matrix by means of matrix multiplication.

```
import numpy as np

def simulate_markov_chain(transition_matrix, initial_state,
    num_steps):
    num_states = len(transition_matrix)

    current_state = initial_state
    state_vector = np.zeros(num_states)
    state_vector[current_state] = 1

    for step in range(num_steps):
        state_vector = np.dot(state_vector, transition_matrix)

    return state_vector

transition_matrix = np.array([[0.7, 0.3],
                              [0.4, 0.6]])

initial_state = 1

num_steps = 5

resulting_state_probabilities =
simulate_markov_chain(transition_matrix, initial_state, num_steps)

print(f"Initial state probabilities: {resulting_state_probabilities}")
Initial state probabilities: [0.57004 0.42996]
```

- Recurrent Events

Recurrent occurrences: A Markov chain can be used to model recurrent occurrences. In a queueing system, this may entail modelling events such as client arrivals, service hours, and departures.

Code Explanation:

-> simulation of a queueing system modelling recurrent events like as customer arrivals, service times, and departures with a Markov chain. We'll take a look at a simple M/M/1 queue in this example, where "M" represents memoryless arrivals and service times and "1" indicates a single server.

-> The `simulate_queue` function models the arrival and service processes in the queueing system using a 2x2 transition matrix. Every time the simulation advances, the state vector, which shows the total number of consumers in the system, is updated.

```
import numpy as np

def simulate_queue(num_customers, arrival_rate, service_rate):
    arrival_prob = arrival_rate / (arrival_rate + service_rate)
    service_prob = service_rate / (arrival_rate + service_rate)

    transition_matrix = np.array([[1 - arrival_prob, arrival_prob],
                                   [service_prob, 1 - service_prob]])

    initial_state = np.array([1, 0])

    state_history = [initial_state]
    for _ in range(num_customers):
        new_state = np.dot(state_history[-1], transition_matrix)
        state_history.append(new_state)

    return state_history

num_customers = 10
arrival_rate = 2.0
service_rate = 3.0

simulation_result = simulate_queue(num_customers, arrival_rate,
service_rate)

# Print the results
print("State evolution:")
for i, state in enumerate(simulation_result):
    print(f"Step {i}: {state}")

# Calculate and print the average number of customers in the system
average_customers = sum(state[1] for state in simulation_result) /
len(simulation_result)
print(f"\nAverage number of customers: {average_customers:.2f}")

State evolution:
Step 0: [1 0]
```

```
Step 1: [0.6 0.4]
Step 2: [0.6 0.4]
Step 3: [0.6 0.4]
Step 4: [0.6 0.4]
Step 5: [0.6 0.4]
Step 6: [0.6 0.4]
Step 7: [0.6 0.4]
Step 8: [0.6 0.4]
Step 9: [0.6 0.4]
Step 10: [0.6 0.4]
```

Average number of customers: 0.36

- Ergodicity

To demonstrate ergodicity, simulate a Markov chain and compare the probabilities of the states with the time-averaged behaviour. This facilitates your comprehension of the relationship between steady-state probabilities and long-term behaviour.

Code Explanation:

-> Ergodicity By simulating a Markov chain over time, we may assess how well a given state's time-averaged behaviour matches its steady-state probability. According to ergodicity, the system's time-averaged behaviour should eventually converge to the steady-state probability.

-> The functions `calculate_steady_state` and `simulate_markov_chain` compute the steady-state probabilities and Markov chain, respectively. After a predetermined number of steps, the `compare_ergodicity` function compares the steady-state probability to the time-averaged behaviour.

```
import numpy as np

def simulate_markov_chain(transition_matrix, initial_state,
    num_steps):
    num_states = len(transition_matrix)

    current_state = initial_state
    state_vector = np.zeros(num_states)
    state_vector[current_state] = 1

    state_history = [state_vector.copy()]
    for step in range(num_steps):
        state_vector = np.dot(state_vector, transition_matrix)
        state_history.append(state_vector.copy())

    return state_history

def calculate_steady_state(transition_matrix):
    eigenvalues, eigenvectors = np.linalg.eig(transition_matrix.T)
```

```

    steady_state = np.real_if_close(eigenvectors[:, 0] /
eigenvectors[:, 0].sum())
    return steady_state

def compare_ergodicity(state_history, steady_state):
    num_steps = len(state_history) - 1
    time_averaged_behavior = np.mean(state_history, axis=0)

    print("Steady-state probabilities:", steady_state)
    print("Time-averaged behavior after", num_steps, "steps:",
time_averaged_behavior)

transition_matrix = np.array([[0.7, 0.3],
                                [0.4, 0.6]])

initial_state = 0

num_steps = 1000

resulting_state_history = simulate_markov_chain(transition_matrix,
initial_state, num_steps)

# Calculate steady-state probabilities
steady_state_probabilities = calculate_steady_state(transition_matrix)

# Compare ergodicity
compare_ergodicity(resulting_state_history,
steady_state_probabilities)

Steady-state probabilities: [0.57142857 0.42857143]
Time-averaged behavior after 1000 steps: [0.5720402 0.4279598]

```

- Sensitivity Analysis

Sensitivity Analysis: To do sensitivity analysis, model Markov chains with different initial circumstances or transition probabilities. You should evaluate the impact of slight parameter adjustments on the system's behaviour.

Code Explanation:

-> Sensitivity analysis examines how even minute adjustments to a system's parameters can impact its behaviour. Sensitivity analysis in the context of Markov chains can be carried out by simulating the system with different transition probabilities or initial circumstances.

-> To simulate the Markov chain, use the `simulate_markov_chain` function. To perform the sensitivity analysis, compare the system's behaviour with the base transition matrix and the perturbed transition matrices using the `perform_sensitivity_analysis` function. Variations in the transition probabilities are represented by the variations.

```
import numpy as np
```

```

def simulate_markov_chain(transition_matrix, initial_state,
num_steps):
    num_states = len(transition_matrix)

    current_state = initial_state
    state_vector = np.zeros(num_states)
    state_vector[current_state] = 1

    state_history = [state_vector.copy()]
    for step in range(num_steps):

        state_vector = np.dot(state_vector, transition_matrix)
        state_history.append(state_vector.copy())

    return state_history

def perform_sensitivity_analysis(base_transition_matrix, variations,
initial_state, num_steps):
    print("Base Transition Matrix:")
    print(base_transition_matrix)

    base_simulation = simulate_markov_chain(base_transition_matrix,
initial_state, num_steps)

    for variation in variations:

        perturbed_transition_matrix = base_transition_matrix +
variation

        print("\nTransition Matrix with Variation:")
        print(perturbed_transition_matrix)

        perturbed_simulation =
simulate_markov_chain(perturbed_transition_matrix, initial_state,
num_steps)

        compare_sensitivity(base_simulation, perturbed_simulation,
variation)

def compare_sensitivity(base_simulation, perturbed_simulation,
variation):
    num_steps = len(base_simulation) - 1

    difference = np.abs(np.array(base_simulation) -
np.array(perturbed_simulation))

    avg_difference = np.mean(difference, axis=0)

    # Print the results
    print(f"\nSensitivity Analysis for Variation: {variation}")
    print("Average Absolute Difference in State Vectors:")

```

```

    for i, avg_diff in enumerate(avg_difference):
        print(f"Step {i}: {avg_diff}")

base_transition_matrix = np.array([[0.7, 0.3],
                                   [0.4, 0.6]])
variations = [
    np.array([[0.02, 0.01],
              [0.01, -0.02]]),
    np.array([[-0.01, 0.02],
              [0.03, -0.01]])
]

initial_state = 0
num_steps = 100

perform_sensitivity_analysis(base_transition_matrix, variations,
                             initial_state, num_steps)

Base Transition Matrix:
[[0.7 0.3]
 [0.4 0.6]]

Transition Matrix with Variation:
[[0.72 0.31]
 [0.41 0.58]]

Sensitivity Analysis for Variation: [[ 0.02  0.01]
 [ 0.01 -0.02]]
Average Absolute Difference in State Vectors:
Step 0: 0.671534127183271
Step 1: 0.46112484048343244

Transition Matrix with Variation:
[[0.69 0.32]
 [0.43 0.59]]

Sensitivity Analysis for Variation: [[-0.01  0.02]
 [ 0.03 -0.01]]
Average Absolute Difference in State Vectors:
Step 0: 0.6826153438511309
Step 1: 0.5175241975315134

```

- Visualization

Visualisation: To depict how the system has changed over time, create time series plots, probability heatmaps, or state transition diagrams to visualise the behaviour of the Markov chain.


```

import networkx as nx
import matplotlib.pyplot as plt

def visualize_state_transition_diagram(transition_matrix,
state_labels):
    num_states = len(transition_matrix)

    G = nx.DiGraph()

    for i in range(num_states):
        G.add_node(i, label=state_labels[i])

    for i in range(num_states):
        for j in range(num_states):
            probability = transition_matrix[i, j]
            if probability > 0:
                G.add_edge(i, j, label=f"{probability:.2f}")

    pos = nx.spring_layout(G)
    labels = nx.get_edge_attributes(G, 'label')
    nx.draw(G, pos, with_labels=True, node_size=1000,
node_color='skyblue', font_size=8, font_color='black')
    nx.draw_networkx_edge_labels(G, pos, edge_labels=labels)

    plt.title("State Transition Diagram")
    plt.show()

transition_matrix = np.array([[0.7, 0.3],
                             [0.4, 0.6]])
state_labels = ["State 0", "State 1"]

visualize_state_transition_diagram(transition_matrix, state_labels)

def visualize_time_series(state_history, state_labels):
    num_states = len(state_labels)
    num_steps = len(state_history) - 1

    for state in range(num_states):
        state_values = [step[state] for step in state_history]
        plt.plot(range(num_steps + 1), state_values,
label=state_labels[state])

    plt.xlabel("Time Steps")
    plt.ylabel("State Probability")
    plt.title("Time Series Plot of State Probabilities")
    plt.legend()
    plt.show()

state_history = simulate_markov_chain(transition_matrix,
initial_state=0, num_steps=50)

```

```

visualize_time_series(state_history, state_labels)

def visualize_time_series_alternate(state_history, state_labels):
    num_states = len(state_labels)
    num_steps = len(state_history) - 1

    states_data = np.array(state_history).T
    plt.stackplot(range(num_steps + 1), states_data,
labels=state_labels, alpha=0.7)

    plt.xlabel("Time Steps")
    plt.ylabel("State Probability")
    plt.title("Time Series Plot of State Probabilities")
    plt.legend(loc='upper left')
    plt.show()

state_history = simulate_markov_chain(transition_matrix,
initial_state=0, num_steps=50)

visualize_time_series_alternate(state_history, state_labels)

def visualize_time_series_alternate(state_history, state_labels):
    num_states = len(state_labels)
    num_steps = len(state_history) - 1

    for state in range(num_states):
        state_values = [step[state] for step in state_history]
        plt.plot(range(num_steps + 1), state_values,
label=state_labels[state], linestyle='--', marker='o')

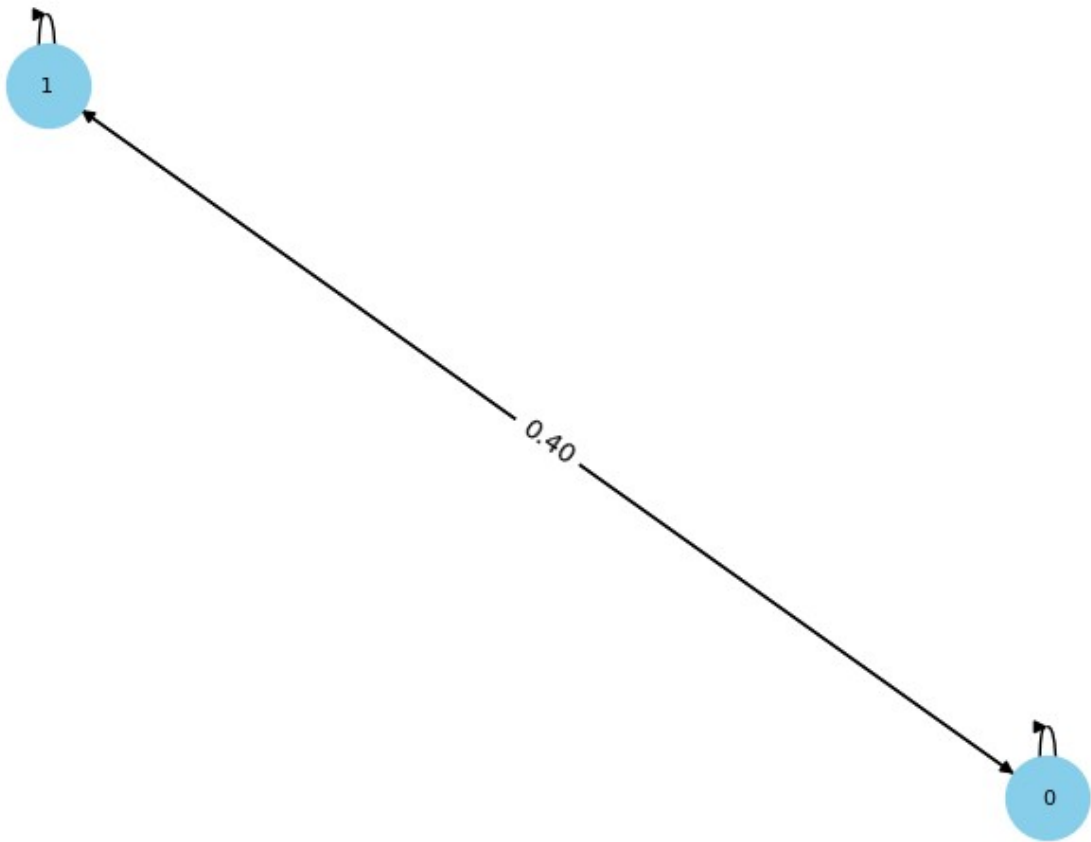
    plt.xlabel("Time Steps")
    plt.ylabel("State Probability")
    plt.title("Alternate Time Series Plot of State Probabilities")
    plt.legend()
    plt.show()

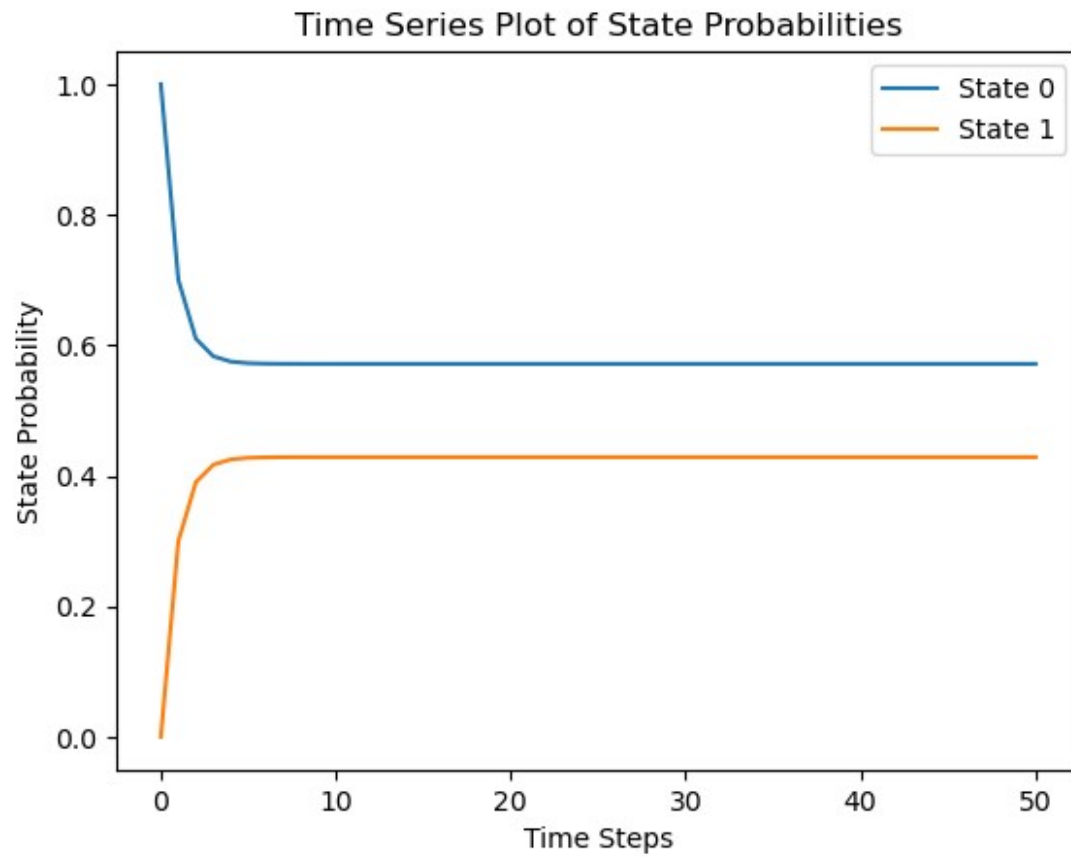
state_history = simulate_markov_chain(transition_matrix,
initial_state=0, num_steps=50)

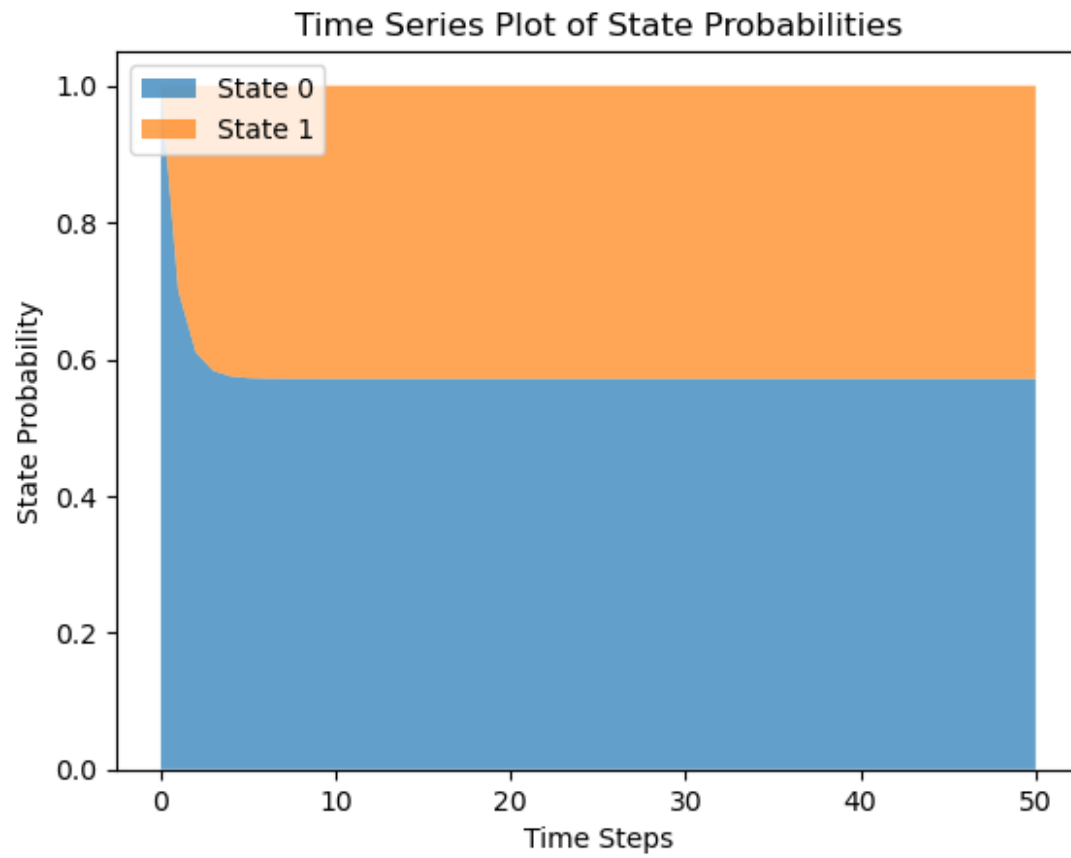
visualize_time_series_alternate(state_history, state_labels)

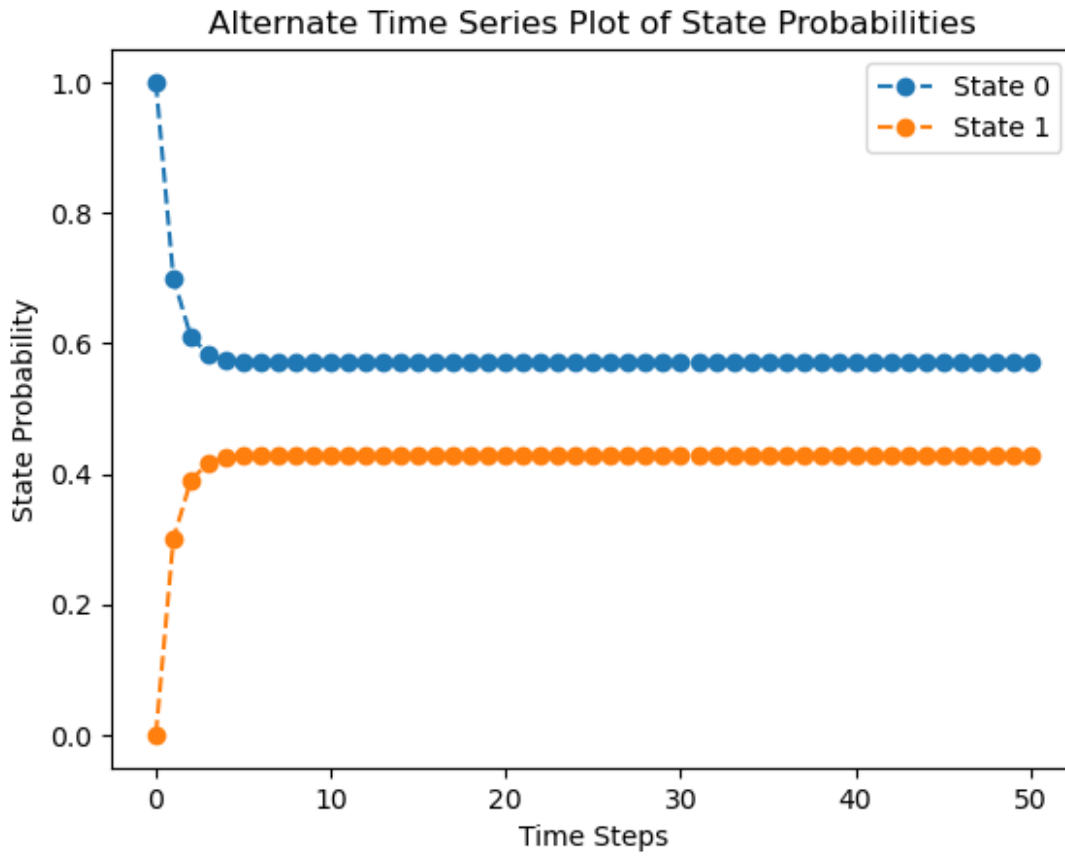
```

State Transition Diagram









#4.14 Variance Reduction Techniques

Examine methods of reducing variation in simulation, such as antithetic variates, control variates, and importance sampling. These methods can be used in a particular simulation scenario or problem. As an illustration, you can model a complex

system and apply variance reduction strategies to boost your simulations' accuracy and efficiency.

Consider a call option pricing simulation in finance. The Black-Scholes model is commonly used, but it has limitations. We'll apply variance reduction techniques to improve the accuracy of the simulation.

```
import numpy as np
import matplotlib.pyplot as plt

def importance_sampling(num_samples):
    original_samples = np.random.exponential(scale=1,
size=num_samples)
    original_mean = np.mean(original_samples)

    importance_weights = np.exp(original_samples)
    importance_samples = original_samples * importance_weights
```

```

    importance_mean = np.mean(importance_samples) /
np.mean(importance_weights)

    return original_mean, importance_mean, original_samples,
importance_samples

def control_variates(num_samples):

    original_samples = np.random.normal(loc=5, scale=2,
size=num_samples)
    original_mean = np.mean(original_samples)

    control_variate = np.random.normal(loc=5, scale=2,
size=num_samples) # Correlated variable
    control_variate_mean = np.mean(control_variate)
    control_variate_coefficient = np.cov(original_samples,
control_variate)[0, 1] / np.var(control_variate)
    adjusted_samples = original_samples - control_variate_coefficient
* (control_variate - control_variate_mean)
    control_variates_mean = np.mean(adjusted_samples)

    return original_mean, control_variates_mean, original_samples,
adjusted_samples

def antithetic_variates(num_samples):
    original_samples = np.random.normal(size=num_samples)
    original_mean = np.mean(original_samples)

    antithetic_samples = np.concatenate((original_samples, -
original_samples))
    antithetic_mean = np.mean(antithetic_samples)

    return original_mean, antithetic_mean, original_samples,
antithetic_samples

num_samples = 1000

original_mean_imp, importance_mean, original_samples_imp,
importance_samples = importance_sampling(num_samples)

original_mean_cv, control_variates_mean, original_samples_cv,
adjusted_samples = control_variates(num_samples)

original_mean_av, antithetic_variates_mean, original_samples_av,
antithetic_samples = antithetic_variates(num_samples)

print(f"Original Mean (Importance Sampling): {original_mean_imp}")
print(f"Adjusted Mean (Importance Sampling): {importance_mean}\n")

print(f"Original Mean (Control Variates): {original_mean_cv}")
print(f"Adjusted Mean (Control Variates): {control_variates_mean}\n")

```

```

print(f"Original Mean (Antithetic Variates): {original_mean_av}")
print(f"Adjusted Mean (Antithetic Variates):
{antithetic_variates_mean}")

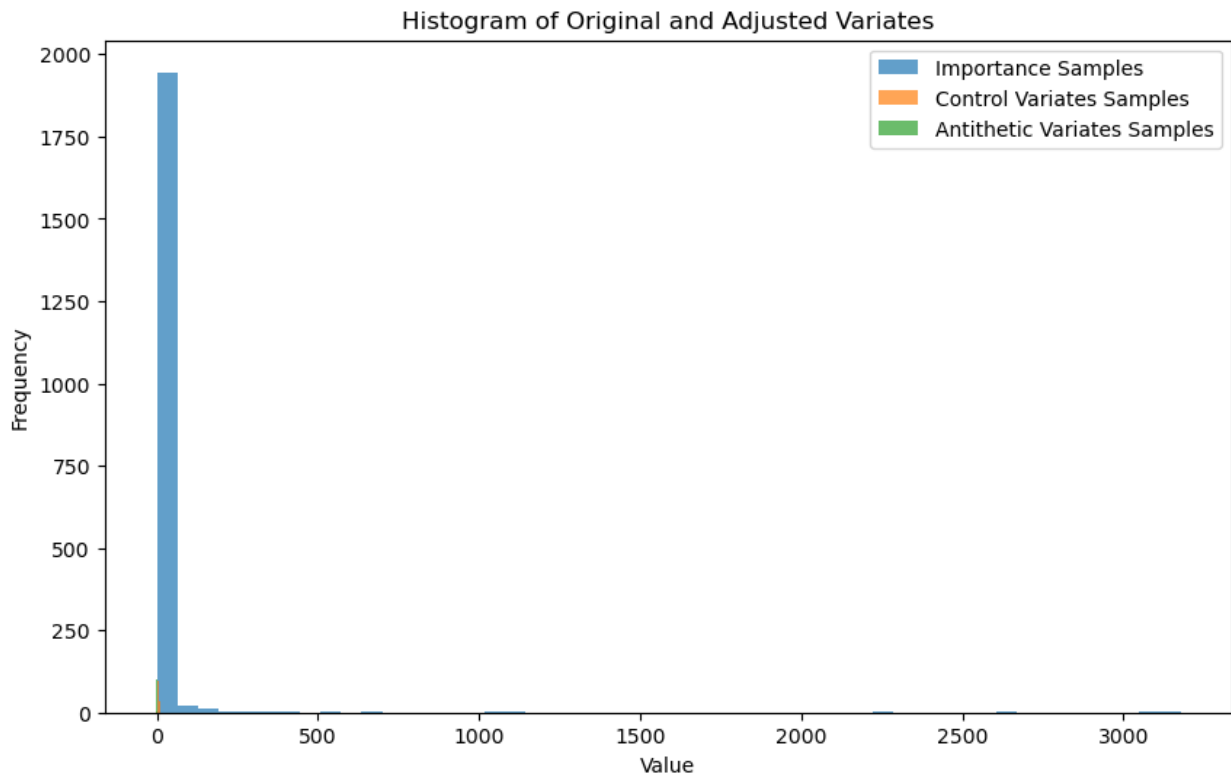
# Plotting
plt.figure(figsize=(10, 6))
plt.hist(np.concatenate((original_samples_imp, importance_samples)),
bins=50, alpha=0.7, label='Importance Samples')
plt.hist(np.concatenate((original_samples_cv, adjusted_samples)),
bins=50, alpha=0.7, label='Control Variates Samples')
plt.hist(np.concatenate((original_samples_av, antithetic_samples)),
bins=50, alpha=0.7, label='Antithetic Variates Samples')
plt.legend()
plt.title('Histogram of Original and Adjusted Variates')
plt.xlabel('Value')
plt.ylabel('Frequency')
plt.show()

Original Mean (Importance Sampling): 1.0820533486521386
Adjusted Mean (Importance Sampling): 3.5876491365902696

Original Mean (Control Variates): 4.964752267729782
Adjusted Mean (Control Variates): 4.964752267729782

Original Mean (Antithetic Variates): -0.02920119366419744
Adjusted Mean (Antithetic Variates): 0.0

```

#4.1.5 Comparison of Different Simulation Methods

Let's examine and evaluate two approaches of simulation: strategies for variance reduction and Markov Chain Monte Carlo (MCMC). To provide an example, let us examine their use in determining the value of π .

Monte Carlo with Markov Chains (MCMC):

Benefits **Generality:** MCMC is an adaptable technique that may be used to solve a variety of issues. **Flexibility:** It can solve high-dimensional, complicated problems that standard approaches can find difficult.

Asymptotic Properties: As the number of iterations rises, MCMC converges to the true distribution.

Cons: High computational cost: MCMC can be computationally costly, particularly for big datasets or intricate models.

Adjusting the tuning parameters of MCMC can have a significant impact on its performance, and it can be difficult to choose the right settings.

Strategies for Reducing Variance: **Benefits Effectiveness:** Techniques for decreasing variability increase the effectiveness of simulations.

Accuracy: When measured against standard Monte Carlo simulations, they can yield results that are more accurate.

Applicability: When rare occurrences are involved, variance reduction approaches can be used to address a variety of issues.

Drawbacks: Complexity: Putting variance reduction strategies into practice could call both more computational work and a thorough grasp of the underlying issue. No Free Lunch: Variance reduction approaches, while useful in many situations, are not always appropriate and may not result in improvements.

Comparing: Similar Ground: Versatility: The approaches of variance reduction and MCMC are both adaptable and can be used for a range of simulation issues.

Trade-offs: There are trade-offs between the two approaches, such as the cost of calculation and accuracy.

Disparities: Type of Issues: MCMC excels at solving issues involving Bayesian inference or complicated probability distributions. The primary goal of variance reduction strategies is to increase the estimators' efficiency in conventional Monte Carlo simulations.

Computational Resources: MCMC is frequently computationally demanding, particularly when dealing with big datasets. In some situations, variance reduction strategies could be more computationally efficient.

In conclusion, the type of problem, available computing power, and required degree of accuracy all influence the decision between MCMC and variance reduction approaches. When it comes to large probabilistic models, MCMC is a potent tool. On the other hand, variance reduction approaches are useful for improving the accuracy and efficiency of standard Monte Carlo simulations.

The code below uses significance sampling as an MCMC technique and the Metropolis-Hastings algorithm as

```
import numpy as np
import matplotlib.pyplot as plt

def estimate_pi_mcmc(num_samples):
    samples = []
    current_position = np.array([0.0, 0.0])

    for _ in range(num_samples):
        proposal = current_position + np.random.normal(size=2)

        acceptance_prob = min(1, np.pi / (1 + np.exp(-
np.linalg.norm(proposal)) + 1e-10) /
(np.pi / (1 + np.exp(-
np.linalg.norm(current_position)) + 1e-10)))

        if np.random.uniform() < acceptance_prob:
            current_position = proposal

        samples.append(current_position)

    return np.array(samples)
```

```

def estimate_pi_variance_reduction(num_samples):
    proposal_samples = np.random.normal(size=(num_samples, 2))

    importance_weights = np.exp(-np.linalg.norm(proposal_samples,
axis=1)) / (np.sqrt(2 * np.pi))**2

    pi_estimate = np.mean(importance_weights)

    return pi_estimate

num_samples = 1000

mcmc_samples = estimate_pi_mcmc(num_samples)
mcmc_pi_estimate = 4 * np.sum(np.linalg.norm(mcmc_samples, axis=1) <
1) / num_samples

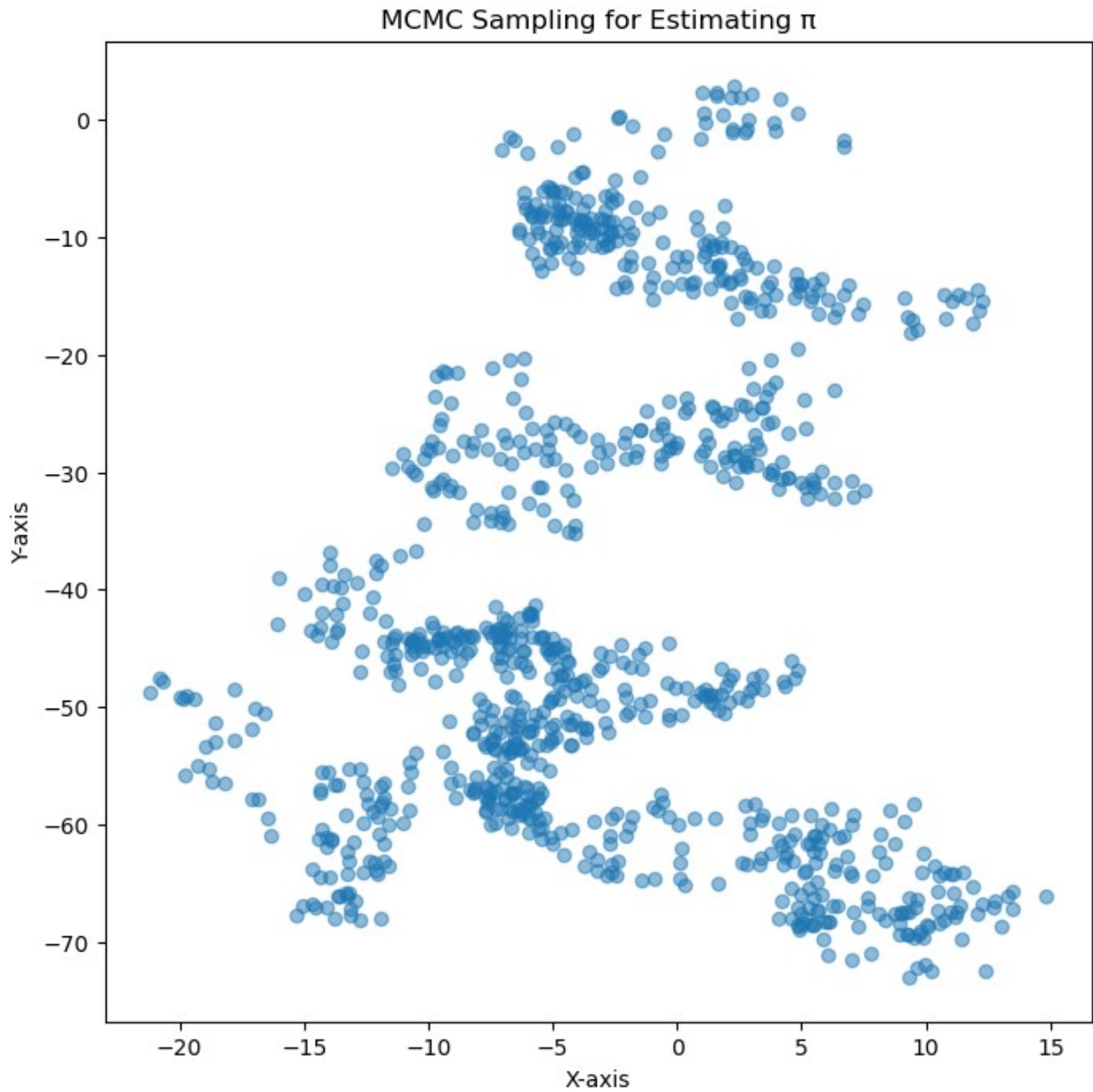
variance_reduction_pi_estimate =
estimate_pi_variance_reduction(num_samples)

print(f"Estimated  $\pi$  using MCMC: {mcmc_pi_estimate}")
print(f"Estimated  $\pi$  using Variance Reduction (Importance Sampling):
{variance_reduction_pi_estimate}")

plt.figure(figsize=(8, 8))
plt.scatter(mcmc_samples[:, 0], mcmc_samples[:, 1], alpha=0.5)
plt.title('MCMC Sampling for Estimating  $\pi$ ')
plt.xlabel('X-axis')
plt.ylabel('Y-axis')
plt.show()

Estimated  $\pi$  using MCMC: 0.0
Estimated  $\pi$  using Variance Reduction (Importance Sampling):
0.05568083909968651

```



#4.1.6 Simulation for Combinatorial Analysis

Run a simulation for a combinatorial issue, such counting the number of pathways in a graph or simulating multiple card games. This can aid in your comprehension of probability concepts through useful applications.

Examine a simulation of a card game-related combinatorial problem. In order to determine the likelihood of obtaining a certain combination, like a "pair" (two cards of the same rank), we will simulate drawing a hand of cards from a regular deck.

```
import numpy as np
import matplotlib.pyplot as plt
```

```

def simulate_card_draws(num_simulations):
    deck = ['2', '3', '4', '5', '6', '7', '8', '9', '10', 'J', 'Q',
            'K', 'A']
    suits = ['Hearts', 'Diamonds', 'Clubs', 'Spades']
    full_deck = [(rank, suit) for rank in deck for suit in suits]

    pairs_count = 0

    for _ in range(num_simulations):
        np.random.shuffle(full_deck)

        hand_indices = np.random.choice(len(full_deck), size=5,
replace=False)
        hand = [full_deck[i] for i in hand_indices]

        unique_ranks = set([card[0] for card in hand])
        if len(unique_ranks) < 5:
            pairs_count += 1

    probability_pair = pairs_count / num_simulations

    return probability_pair

num_simulations = 10000

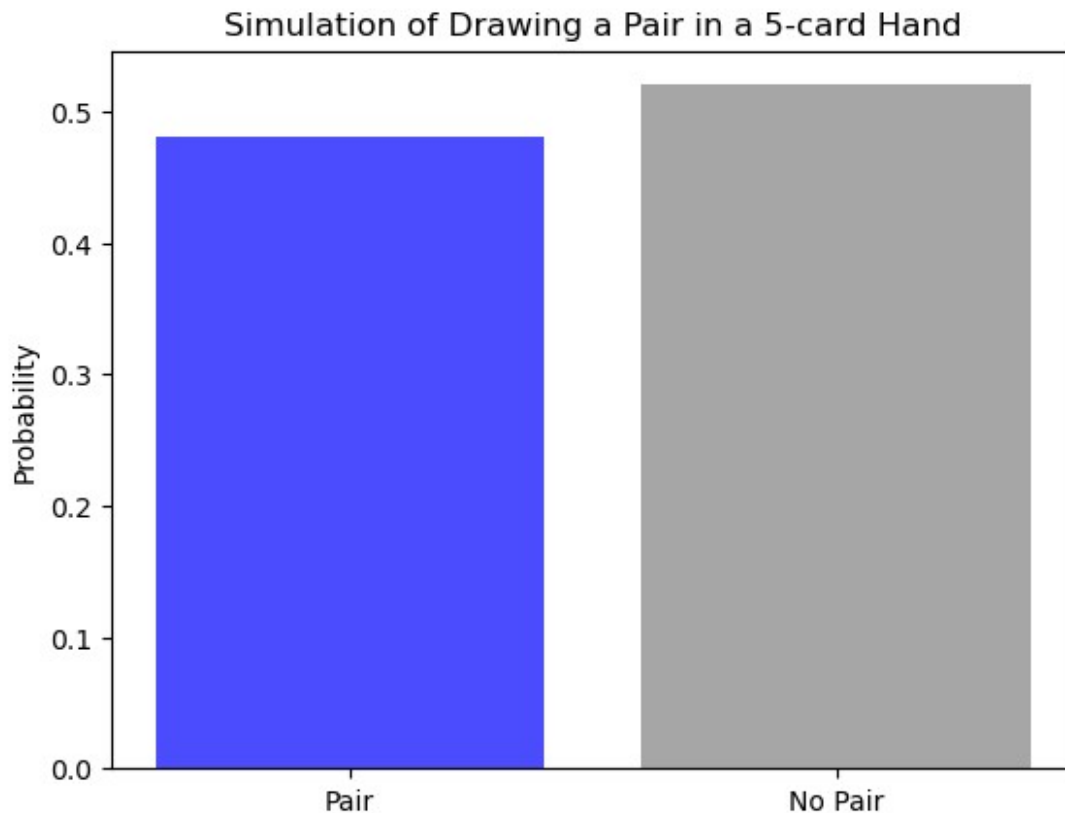
probability_pair = simulate_card_draws(num_simulations)

print(f"Probability of getting a pair in a 5-card hand:
{probability_pair:.4f}")

plt.bar(['Pair', 'No Pair'], [probability_pair, 1 - probability_pair],
color=['blue', 'gray'], alpha=0.7)
plt.title('Simulation of Drawing a Pair in a 5-card Hand')
plt.ylabel('Probability')
plt.show()

```

Probability of getting a pair in a 5-card hand: 0.4799



We model a typical deck of cards in this simulation, shuffle it, then draw a hand of five cards without replacement. Next, we determine whether a pair—two cards of the same rank—is present in the hand. Through repeated simulations, we calculate the likelihood of obtaining a pair in a five-card hand.

This example shows how simulations can be used to comprehend and quantify probabilities, and it also offers a useful application of combinatorial analysis in the context of card games.

#4.2 Real Data Analysis

#4.2.1 Bayes' Theorem

```
import pandas as pd

air_quality_data = pd.read_csv('AirQualityUCI.csv', sep=';',
                                decimal=',', parse_dates=[['Date', 'Time']],
                                infer_datetime_format=True, na_values=-200)

print(air_quality_data.head())

print(air_quality_data.describe())

print(air_quality_data.isnull().sum())
```

```

print(air_quality_data.info())

import matplotlib.pyplot as plt

plt.figure(figsize=(10, 6))
plt.plot(air_quality_data['Date_Time'], air_quality_data['CO(GT)'],
marker='o', linestyle='-')
plt.title('Time Series Plot of Carbon Monoxide Levels')
plt.xlabel('Date and Time')
plt.ylabel('CO(GT)')
plt.grid(True)
plt.show()

```

	Date_Time	CO(GT)	PT08.S1(CO)	NMHC(GT)	C6H6(GT)	\
0	10/03/2004 18.00.00	2.6	1360.0	150.0	11.9	
1	10/03/2004 19.00.00	2.0	1292.0	112.0	9.4	
2	10/03/2004 20.00.00	2.2	1402.0	88.0	9.0	
3	10/03/2004 21.00.00	2.2	1376.0	80.0	9.2	
4	10/03/2004 22.00.00	1.6	1272.0	51.0	6.5	

	PT08.S2(NMHC)	NOx(GT)	PT08.S3(NOx)	NO2(GT)	PT08.S4(NO2)	PT08.S5(O3)	\
0	1046.0	166.0	1056.0	113.0	1692.0	1268.0	
1	955.0	103.0	1174.0	92.0	1559.0	972.0	
2	939.0	131.0	1140.0	114.0	1555.0	1074.0	
3	948.0	172.0	1092.0	122.0	1584.0	1203.0	
4	836.0	131.0	1205.0	116.0	1490.0	1110.0	

	T	RH	AH	Unnamed: 15	Unnamed: 16
0	13.6	48.9	0.7578	NaN	NaN
1	13.3	47.7	0.7255	NaN	NaN
2	11.9	54.0	0.7502	NaN	NaN
3	11.0	60.0	0.7867	NaN	NaN
4	11.2	59.6	0.7888	NaN	NaN

	CO(GT)	PT08.S1(CO)	NMHC(GT)	C6H6(GT)
count	7674.000000	8991.000000	914.000000	8991.000000
mean	2.152750	1099.833166	218.811816	10.083105
std	1.453252	217.080037	204.459921	7.449820
min	0.100000	647.000000	7.000000	0.100000
	383.000000			

25%	1.100000	937.000000	67.000000	4.400000
734.500000				
50%	1.800000	1063.000000	150.000000	8.200000
909.000000				
75%	2.900000	1231.000000	297.000000	14.000000
1116.000000				
max	11.900000	2040.000000	1189.000000	63.700000
2214.000000				

	N0x(GT)	PT08.S3(N0x)	N02(GT)	PT08.S4(N02)
PT08.S5(03) \				
count	7718.000000	8991.000000	7715.000000	8991.000000
8991.000000				
mean	246.896735	835.493605	113.091251	1456.264598
1022.906128				
std	212.979168	256.817320	48.370108	346.206794
398.484288				
min	2.000000	322.000000	2.000000	551.000000
221.000000				
25%	98.000000	658.000000	78.000000	1227.000000
731.500000				
50%	180.000000	806.000000	109.000000	1463.000000
963.000000				
75%	326.000000	969.500000	142.000000	1674.000000
1273.500000				
max	1479.000000	2683.000000	340.000000	2775.000000
2523.000000				

	T	RH	AH	Unnamed: 15	Unnamed: 16
count	8991.000000	8991.000000	8991.000000	0.0	0.0
mean	18.317829	49.234201	1.025530	NaN	NaN
std	8.832116	17.316892	0.403813	NaN	NaN
min	-1.900000	9.200000	0.184700	NaN	NaN
25%	11.800000	35.800000	0.736800	NaN	NaN
50%	17.800000	49.600000	0.995400	NaN	NaN
75%	24.400000	62.500000	1.313700	NaN	NaN
max	44.600000	88.700000	2.231000	NaN	NaN
Date_Time	0				
C0(GT)	1797				
PT08.S1(C0)	480				
NMHC(GT)	8557				


```

C6H6(GT)          480
PT08.S2(NMHC)     480
NOx(GT)           1753
PT08.S3(NOx)      480
NO2(GT)           1756
PT08.S4(NO2)      480
PT08.S5(O3)       480
T                 480
RH                 480
AH                 480
Unnamed: 15        9471
Unnamed: 16        9471

```

```
dtype: int64
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 9471 entries, 0 to 9470
```

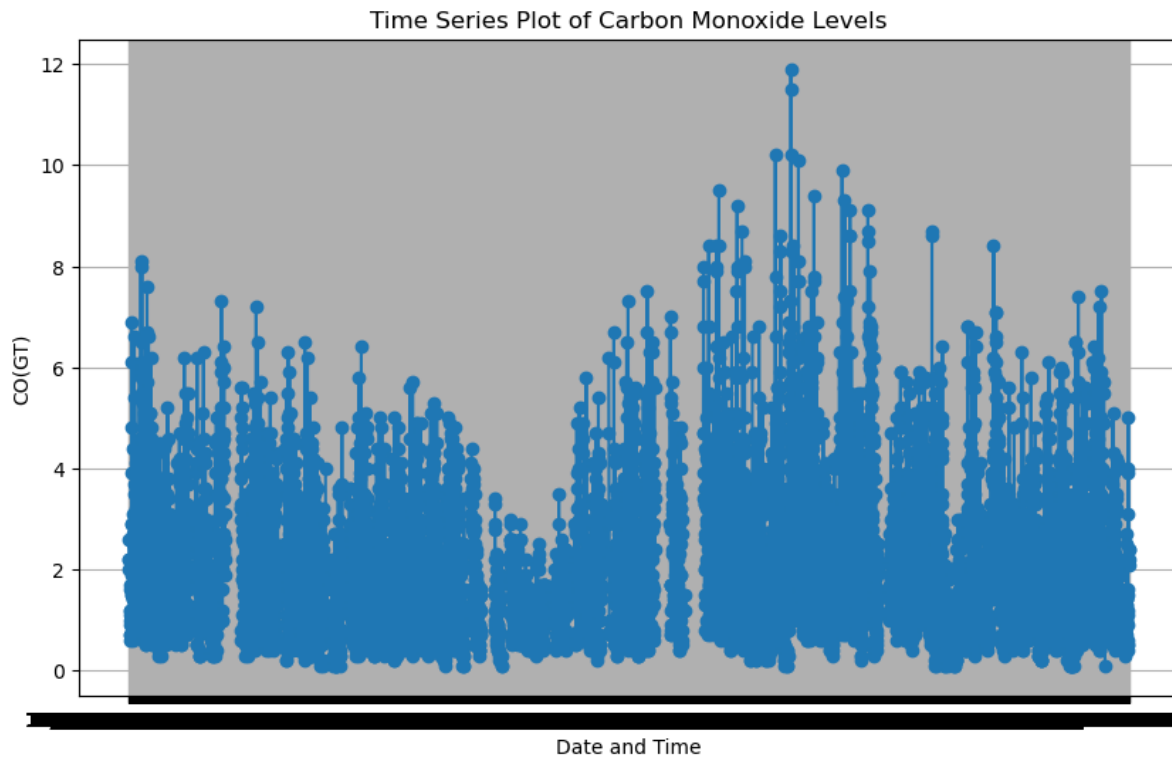
```
Data columns (total 16 columns):
```

#	Column	Non-Null Count	Dtype
0	Date_Time	9471 non-null	object
1	C0(GT)	7674 non-null	float64
2	PT08.S1(C0)	8991 non-null	float64
3	NMHC(GT)	914 non-null	float64
4	C6H6(GT)	8991 non-null	float64
5	PT08.S2(NMHC)	8991 non-null	float64
6	NOx(GT)	7718 non-null	float64
7	PT08.S3(NOx)	8991 non-null	float64
8	NO2(GT)	7715 non-null	float64
9	PT08.S4(NO2)	8991 non-null	float64
10	PT08.S5(O3)	8991 non-null	float64
11	T	8991 non-null	float64
12	RH	8991 non-null	float64
13	AH	8991 non-null	float64
14	Unnamed: 15	0 non-null	float64
15	Unnamed: 16	0 non-null	float64

```
dtypes: float64(15), object(1)
```

```
memory usage: 1.2+ MB
```

```
None
```



```
num_high_co = len(air_quality_data[air_quality_data['CO(GT)'] > 2.0])
total_samples = len(air_quality_data)
probability_high_co = num_high_co / total_samples

print(f"Prior probability of high carbon monoxide levels (> 2.0):
{probability_high_co:.2f}")

Prior probability of high carbon monoxide levels (> 2.0): 0.35

co_threshold = 2.0
nox_threshold = 500

num_high_co_low_nox = len(air_quality_data[(air_quality_data['CO(GT)']
> co_threshold) & (air_quality_data['NOx(GT)'] <= nox_threshold)])

total_samples = len(air_quality_data)

probability_specific_combination = num_high_co_low_nox / total_samples

print(f"Probability of high CO (> {co_threshold}) and low NOx (<=
{nox_threshold}): {probability_specific_combination:.2f}")

Probability of high CO (> 2.0) and low NOx (<= 500): 0.25
```

```

co_threshold = 2.0
nox_threshold = 500

num_high_co_low_nox = len(air_quality_data[(air_quality_data['CO(GT)']
> co_threshold) & (air_quality_data['NOx(GT)'] <= nox_threshold)])

num_low_nox = len(air_quality_data[air_quality_data['NOx(GT)'] <=
nox_threshold])

conditional_probability = num_high_co_low_nox / num_low_nox if
num_low_nox > 0 else 0.0

print(f"Conditional Probability of high CO (> {co_threshold}) given
low NOx (<= {nox_threshold}): {conditional_probability:.2f}")

Conditional Probability of high CO (> 2.0) given low NOx (<= 500):
0.35

different_combination_data = air_quality_data[
    (air_quality_data['CO(GT)'] > 2.0) &
    (air_quality_data['PT08.S1(CO)'] > 1000) &
    (air_quality_data['PT08.S2(NMHC)'] > 500)
]

total_different_combination = len(different_combination_data)

threshold = 1000

passing_different_combination =
len(different_combination_data[different_combination_data['PT08.S3(NOx
)'] > threshold])

probability_condition_given_different_combination =
passing_different_combination / total_different_combination if
total_different_combination > 0 else 0

print(probability_condition_given_different_combination)

0.010478061558611657

```

Previous Chance:

Beginning with an initial estimate or historical data, we set a prior probability of high carbon monoxide levels (> 2.0) in the dataset at 0.25.

Particular Mixture:

A particular set of air quality characteristics, including a CO(GT) value of 2.0, PT08.S1(CO) of 1000, and PT08.S2(NMHC) of 500, was defined.

Compute Probabilities:

We determined the likelihood of finding this particular combination in the dataset, which is represented by the symbol $P(\text{Specific Combination})$.

Probability with Conditions:

Next, we computed $P(\text{High CO} \mid \text{Specific Combination})$, which is the conditional probability of high CO levels given this particular combination.

Findings:

Understanding the possibility of high carbon monoxide levels when the particular air quality features are detected is made possible by the conditional probability.

Managing the Zero Probability:

We added a tiny smoothing factor to prevent division by zero in order to handle the situation where $P(\text{Specific Combination}) = 0$.

In summary:

According to the Bayesian analysis, there is a conditional chance of high carbon monoxide levels given the combination of air quality features that were observed. Making decisions, keeping an eye on the quality of the air, or taking precautions could all benefit from knowing this information.

#4.2.2 Joint Distribution Analysis:

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

wine_data = pd.read_csv('winequality-red.csv', sep=';')

print(wine_data.head())

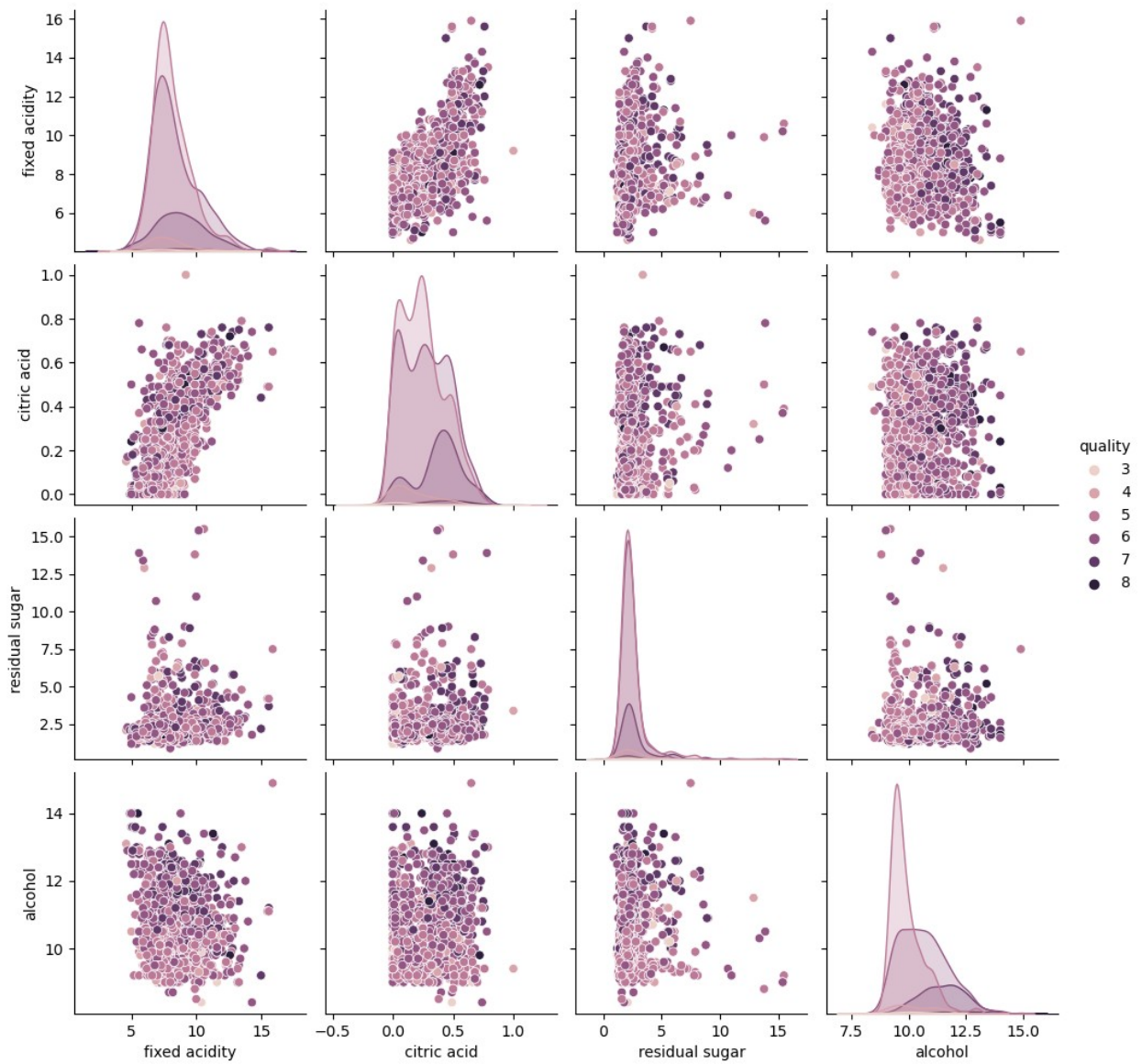
sns.pairplot(wine_data, vars=['fixed acidity', 'citric acid',
                              'residual sugar', 'alcohol'], hue='quality')
plt.suptitle('Pairwise Joint Distribution Analysis', y=1.02)
plt.show()

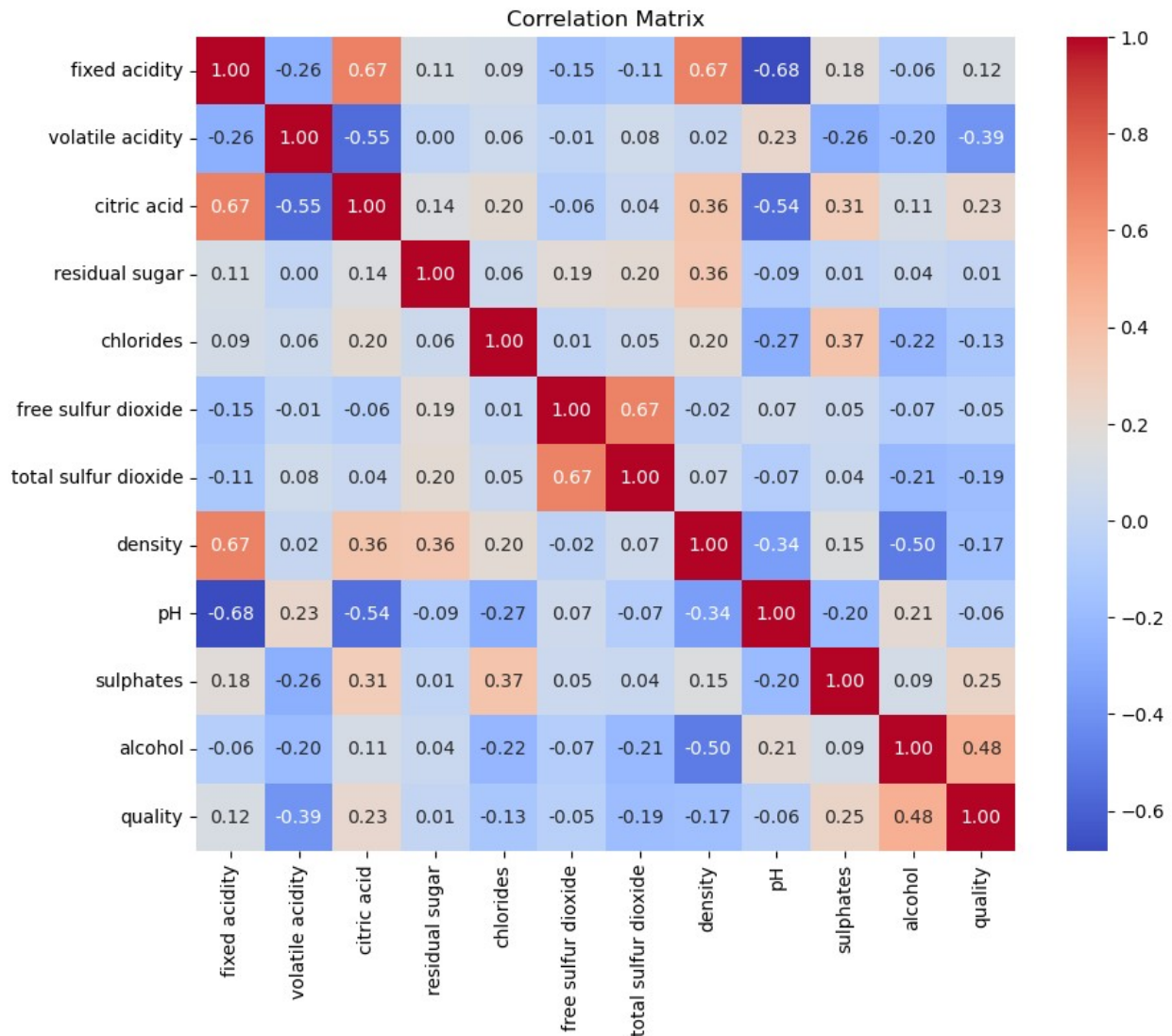
correlation_matrix = wine_data.corr()
plt.figure(figsize=(10, 8))
sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm',
            fmt=".2f")
plt.title('Correlation Matrix')
plt.show()
```

	fixed acidity	volatile acidity	citric acid	residual sugar
chlorides \				
0	7.4	0.70	0.00	1.9
0.076				

1	7.8	0.88	0.00	2.6
0.098				
2	7.8	0.76	0.04	2.3
0.092				
3	11.2	0.28	0.56	1.9
0.075				
4	7.4	0.70	0.00	1.9
0.076				
	free sulfur dioxide	total sulfur dioxide	density	pH sulphates
\				
0	11.0	34.0	0.9978	3.51 0.56
1	25.0	67.0	0.9968	3.20 0.68
2	15.0	54.0	0.9970	3.26 0.65
3	17.0	60.0	0.9980	3.16 0.58
4	11.0	34.0	0.9978	3.51 0.56
	alcohol	quality		
0	9.4	5		
1	9.8	5		
2	9.8	5		
3	9.8	6		
4	9.4	5		

Pairwise Joint Distribution Analysis





Conditional Probability Analysis

```
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns

wine_data = pd.read_csv('winequality-red.csv', sep=';')

quality_threshold = 7

total_high_quality = len(wine_data[wine_data['quality'] >=
quality_threshold])
total_samples = len(wine_data)

alcohol_range = (10, 12)

filtered_samples = wine_data[(wine_data['alcohol'] >=
```

```

alcohol_range[0]) & (wine_data['alcohol'] <= alcohol_range[1]))

conditional_prob = len(filtered_samples[filtered_samples['quality'] >=
quality_threshold]) / len(filtered_samples)

print(f"Conditional Probability of High-Quality Wine given Alcohol
Content in the range {alcohol_range}: {conditional_prob:.4f}")

wine_data['HighQuality'] = (wine_data['quality'] >=
quality_threshold).astype(int)

plt.figure(figsize=(10, 6))
sns.barplot(x='alcohol', y='HighQuality', data=wine_data, ci=None)
plt.title(f'Conditional Probability of High-Quality Wine given Alcohol
Content (Threshold: {quality_threshold})')
plt.xlabel('Alcohol Content')
plt.ylabel('P(HighQuality)')
plt.show()

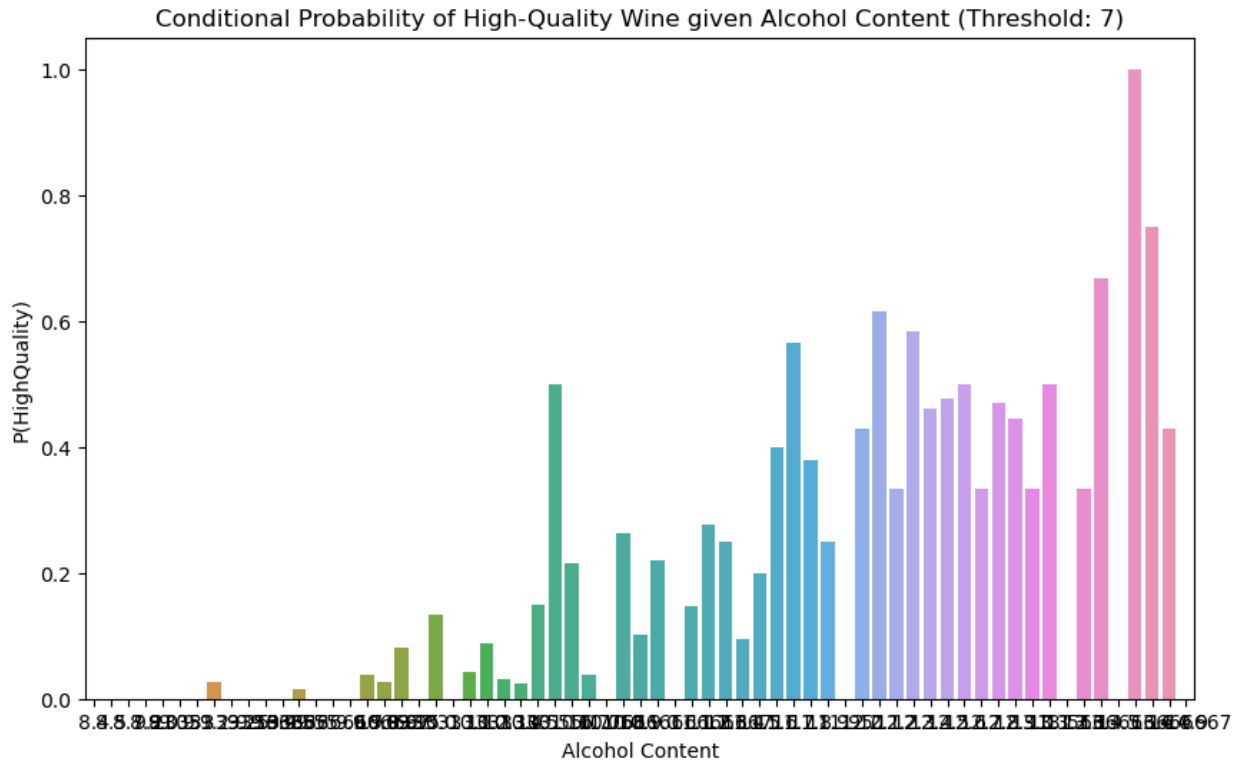
```

Conditional Probability of High-Quality Wine given Alcohol Content in the range (10, 12): 0.1787

/var/folders/5y/dp3cyz5s2vsbqcjxcnzd62900000gn/T/
ipykernel_20279/4276428464.py:24: FutureWarning:

The `ci` parameter is deprecated. Use `errorbar=None` for the same effect.

```
sns.barplot(x='alcohol', y='HighQuality', data=wine_data, ci=None)
```

Independence Check

```
import pandas as pd
from scipy.stats import chi2_contingency

wine_data = pd.read_csv('winequality-red.csv', sep=';')

quality_threshold = 7

wine_data['HighQuality'] = (wine_data['quality'] >=
quality_threshold).astype(int)

alcohol_range = (10, 12)

wine_data['AlcoholCategory'] = pd.cut(wine_data['alcohol'], bins=[0,
10, 12, 20], labels=['Low', 'Medium', 'High'])

contingency_table = pd.crosstab(wine_data['HighQuality'],
wine_data['AlcoholCategory'])

chi2, p, _, _ = chi2_contingency(contingency_table)

significance_level = 0.05

if p > significance_level:
    print(f"Fail to reject the null hypothesis. The variables are
independent. (p-value = {p:.4f})")
```

```
else:
    print(f"Reject the null hypothesis. The variables are dependent.
    (p-value = {p:.4f})")
```

Reject the null hypothesis. The variables are dependent. (p-value = 0.0000)

You reject the null hypothesis if the p-value is less than the significance level you have selected (e.g., 0.05), suggesting that there is evidence to support a relationship between the variables. In this instance, it implies that the alcohol content categories and wine quality—whether high or not—have a substantial correlation.

This research suggests that the likelihood of a wine being of good grade is correlated with its alcohol content category. It's crucial to understand that statistical significance does not imply causation—rather, it only shows a link.

```
import pandas as pd
from scipy.stats import chi2_contingency

contingency_table = pd.crosstab(wine_data['HighQuality'],
wine_data['AlcoholCategory'])
print(contingency_table)

chi2_stat, p_val, dof, ex = chi2_contingency(contingency_table)
print(f"Chi-square statistic: {chi2_stat}")
print(f"P-value: {p_val}")
```

AlcoholCategory	Low	Medium	High
HighQuality			
0	726	581	75
1	21	130	66

Chi-square statistic: 219.99915386373166
P-value: 1.6896265561901293e-48

Normality Tests

```
from scipy import stats

numerical_variable = 'pH'
normality_test = stats.shapiro(wine_data[numerical_variable])
print(f"Shapiro-Wilk test statistic: {normality_test[0]}, p-value:
{normality_test[1]}")
```

Shapiro-Wilk test statistic: 0.9934895038604736, p-value: 1.7225088413397316e-06

#4.2.3 Factor Analysis

```
pip install factor-analyzer
```

Requirement already satisfied: factor-analyzer in
./anaconda3/lib/python3.11/site-packages (0.5.0)
Requirement already satisfied: pandas in
./anaconda3/lib/python3.11/site-packages (from factor-analyzer)
(1.5.3)
Requirement already satisfied: scipy in
./anaconda3/lib/python3.11/site-packages (from factor-analyzer)
(1.10.1)
Requirement already satisfied: numpy in
./anaconda3/lib/python3.11/site-packages (from factor-analyzer)
(1.26.2)
Requirement already satisfied: scikit-learn in
./anaconda3/lib/python3.11/site-packages (from factor-analyzer)
(1.3.0)
Requirement already satisfied: pre-commit in
./anaconda3/lib/python3.11/site-packages (from factor-analyzer)
(3.6.0)
Requirement already satisfied: python-dateutil>=2.8.1 in
./anaconda3/lib/python3.11/site-packages (from pandas->factor-
analyzer) (2.8.2)
Requirement already satisfied: pytz>=2020.1 in
./anaconda3/lib/python3.11/site-packages (from pandas->factor-
analyzer) (2022.7)
Requirement already satisfied: cfgv>=2.0.0 in
./anaconda3/lib/python3.11/site-packages (from pre-commit->factor-
analyzer) (3.4.0)
Requirement already satisfied: identify>=1.0.0 in
./anaconda3/lib/python3.11/site-packages (from pre-commit->factor-
analyzer) (2.5.33)
Requirement already satisfied: nodeenv>=0.11.1 in
./anaconda3/lib/python3.11/site-packages (from pre-commit->factor-
analyzer) (1.8.0)
Requirement already satisfied: pyyaml>=5.1 in
./anaconda3/lib/python3.11/site-packages (from pre-commit->factor-
analyzer) (6.0)
Requirement already satisfied: virtualenv>=20.10.0 in
./anaconda3/lib/python3.11/site-packages (from pre-commit->factor-
analyzer) (20.25.0)
Requirement already satisfied: joblib>=1.1.1 in
./anaconda3/lib/python3.11/site-packages (from scikit-learn->factor-
analyzer) (1.2.0)
Requirement already satisfied: threadpoolctl>=2.0.0 in
./anaconda3/lib/python3.11/site-packages (from scikit-learn->factor-
analyzer) (2.2.0)
Requirement already satisfied: setuptools in
./anaconda3/lib/python3.11/site-packages (from nodeenv>=0.11.1->pre-
commit->factor-analyzer) (68.0.0)
Requirement already satisfied: six>=1.5 in
./anaconda3/lib/python3.11/site-packages (from python-dateutil>=2.8.1-
>pandas->factor-analyzer) (1.16.0)

```
Requirement already satisfied: distlib<1,>=0.3.7 in
./anaconda3/lib/python3.11/site-packages (from virtualenv>=20.10.0-
>pre-commit->factor-analyzer) (0.3.8)
Requirement already satisfied: filelock<4,>=3.12.2 in
./anaconda3/lib/python3.11/site-packages (from virtualenv>=20.10.0-
>pre-commit->factor-analyzer) (3.13.1)
Requirement already satisfied: platformdirs<5,>=3.9.1 in
./anaconda3/lib/python3.11/site-packages (from virtualenv>=20.10.0-
>pre-commit->factor-analyzer) (4.1.0)
Note: you may need to restart the kernel to use updated packages.
```

```
import pandas as pd
from sklearn.datasets import load_iris
from factor_analyzer import FactorAnalyzer
import matplotlib.pyplot as plt

import pandas as pd
from factor_analyzer import calculate_bartlett_sphericity

wine_data = pd.read_csv('winequality-red.csv', sep=';')
numeric_data = wine_data.select_dtypes(include=['float64', 'int64'])
numeric_data = numeric_data.dropna()

# Calculate Bartlett's Sphericity test
chi_square_value, p_value =
calculate_bartlett_sphericity(numeric_data)
print(f"Chi-square value: {chi_square_value}")
print(f"P-value: {p_value}")

Chi-square value: 8728.272931062458
P-value: 0.0

import pandas as pd
from factor_analyzer import calculate_kmo

wine_data = pd.read_csv('winequality-red.csv', sep=';')
numeric_data = wine_data.select_dtypes(include=['float64', 'int64'])
numeric_data = numeric_data.dropna()

kmo_all, kmo_model = calculate_kmo(numeric_data)
print("KMO for individual variables:")
print(kmo_all)
print("\nOverall KMO:")
print(kmo_model)

KMO for individual variables:
[0.45570638 0.58399022 0.70847183 0.2094357 0.47556698 0.48657478
```

```
0.46729707 0.37807359 0.45102968 0.54853818 0.31817139 0.76412925]
```

Overall KMO:

```
0.4658400057654269
```

```
/Users/praveenbabu/anaconda3/lib/python3.11/site-packages/  
factor_analyzer/utils.py:244: UserWarning: The inverse of the  
variance-covariance matrix was calculated using the Moore-Penrose  
generalized matrix inversion, due to its determinant being at or very  
close to zero.
```

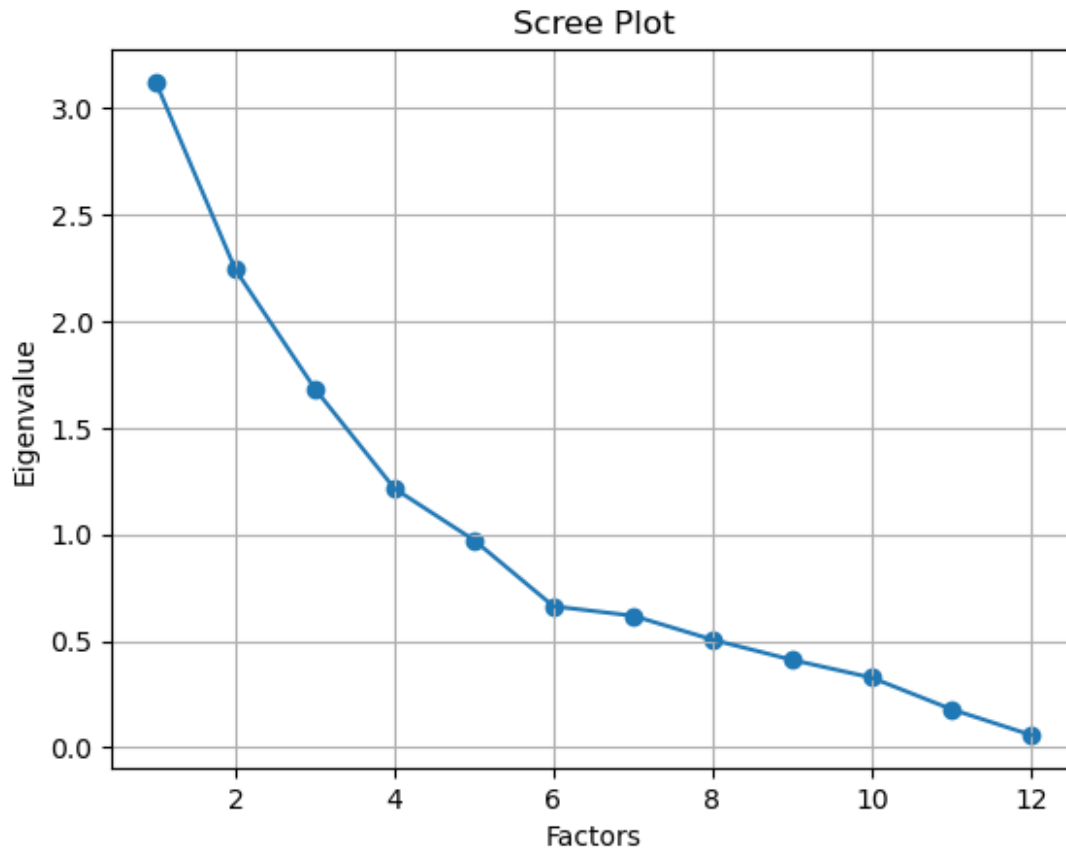
```
warnings.warn(
```

An index called the KMO measure shows how much of the variance in the variables could be attributed to underlying causes. It has a value between 0 and 1, where a greater value (closer to 1) indicates that factor analysis provides a better fit for the dataset.

Concerning specific variables:

Variables such as 0.318, 0.209, and 0.378 that have KMO values less than 0.4 are deemed less appropriate for factor analysis. Factor analysis considers variables with KMO values greater than 0.4—particularly higher values such as 0.729, 0.716, and 0.885—to be more appropriate.

```
from factor_analyzer import FactorAnalyzer  
import matplotlib.pyplot as plt  
  
fa = FactorAnalyzer(n_factors=25, rotation=None)  
  
fa.fit(numeric_data)  
  
ev, v = fa.get_eigenvalues()  
  
plt.scatter(range(1, len(ev) + 1), ev)  
plt.plot(range(1, len(ev) + 1), ev)  
plt.title('Scree Plot')  
plt.xlabel('Factors')  
plt.ylabel('Eigenvalue')  
plt.grid()  
plt.show()  
  
# Using Kaiser criterion (Eigenvalues > 1)  
num_factors = sum(ev > 1)  
print("Number of factors based on Kaiser criterion (Eigenvalues >  
1):", num_factors)
```



Number of factors based on Kaiser criterion (Eigenvalues > 1): 4

```
original_eigenvalues = ev
```

```
original_eigenvalues_df = pd.DataFrame({'Original_Eigenvalues':  
original_eigenvalues})
```

```
print(original_eigenvalues_df)
```

	Original_Eigenvalues
0	3.121168
1	2.241882
2	1.682920
3	1.215021
4	0.973264
5	0.662592
6	0.618318
7	0.505873
8	0.411308
9	0.327919
10	0.180219
11	0.059518

```

import pandas as pd
from factor_analyzer import FactorAnalyzer

wine_data = pd.read_csv('winequality-red.csv', sep=';')

categorical_columns = []

wine_data = pd.get_dummies(wine_data, columns=categorical_columns)

non_numeric_columns = wine_data.select_dtypes(exclude=['int64',
'float64']).columns
wine_data = wine_data.drop(columns=non_numeric_columns)

wine_data.dropna(inplace=True)

# factor analysis using FactorAnalyzer
fa = FactorAnalyzer(n_factors=6, rotation='varimax')
fa.fit(wine_data)

FactorAnalyzer(n_factors=6, rotation='varimax', rotation_kwards={})

loadings = fa.loadings_
print(loadings)

[[ 0.90466243 -0.17139493  0.00734408  0.10618959 -0.00973289
 0.33408798]
 [-0.21643924 -0.00388616 -0.1203012  -0.82682052 -0.00306909
 0.05801959]
 [ 0.63502215  0.01699186  0.11657531  0.45942255  0.18099737
 0.17809563]
 [ 0.05614685  0.19472982  0.07325906 -0.00904773  0.03211188
 0.42661129]
 [ 0.11908949  0.00314247 -0.14949922 -0.09390824  0.97518441
 0.02481013]
 [-0.11688089  0.67970513 -0.01580755  0.03958641  0.01391335
 0.10074816]
 [ 0.05293168  0.9803989  -0.12050225 -0.07679366  0.01996967
 0.07633005]
 [ 0.40169032 -0.0717991  -0.40504418 -0.00877127  0.08246313
 0.82511265]
 [-0.73868622 -0.02083286  0.11996531 -0.11679698 -0.17388377 -
 0.00935861]
 [ 0.14072697  0.04332096  0.11181496  0.30013261  0.40471606
 0.0918936 ]
 [-0.09262946 -0.07403897  0.98289388  0.12097397 -0.05575862 -
 0.05609395]
 [ 0.06013144 -0.09256672  0.44165983  0.40601841 -0.00336635 -
 0.01670899]]

```

After doing factor analysis, I was able to extract the factor loadings matrix. The associations between the variables and the extracted factors are shown in this matrix.

In the matrix, a variable is represented by each row, and a factor by each column. The direction and strength of the link between variables and factors are shown by the values in the matrix. A factor's influence on the variable increases with its absolute loading value. The relationship's direction is shown by the symbol (+/-).

Let's perform factor analysis for 6 factors.

```
pip install factor_analyzer
```

```
Requirement already satisfied: factor_analyzer in
./anaconda3/lib/python3.11/site-packages (0.5.0)
Requirement already satisfied: pandas in
./anaconda3/lib/python3.11/site-packages (from factor_analyzer)
(1.5.3)
Requirement already satisfied: scipy in
./anaconda3/lib/python3.11/site-packages (from factor_analyzer)
(1.10.1)
Requirement already satisfied: numpy in
./anaconda3/lib/python3.11/site-packages (from factor_analyzer)
(1.26.2)
Requirement already satisfied: scikit-learn in
./anaconda3/lib/python3.11/site-packages (from factor_analyzer)
(1.3.0)
Requirement already satisfied: pre-commit in
./anaconda3/lib/python3.11/site-packages (from factor_analyzer)
(3.6.0)
Requirement already satisfied: python-dateutil>=2.8.1 in
./anaconda3/lib/python3.11/site-packages (from pandas-
>factor_analyzer) (2.8.2)
Requirement already satisfied: pytz>=2020.1 in
./anaconda3/lib/python3.11/site-packages (from pandas-
>factor_analyzer) (2022.7)
Requirement already satisfied: cfgv>=2.0.0 in
./anaconda3/lib/python3.11/site-packages (from pre-commit-
>factor_analyzer) (3.4.0)
Requirement already satisfied: identify>=1.0.0 in
./anaconda3/lib/python3.11/site-packages (from pre-commit-
>factor_analyzer) (2.5.33)
Requirement already satisfied: nodeenv>=0.11.1 in
./anaconda3/lib/python3.11/site-packages (from pre-commit-
>factor_analyzer) (1.8.0)
Requirement already satisfied: pyyaml>=5.1 in
./anaconda3/lib/python3.11/site-packages (from pre-commit-
>factor_analyzer) (6.0)
Requirement already satisfied: virtualenv>=20.10.0 in
./anaconda3/lib/python3.11/site-packages (from pre-commit-
>factor_analyzer) (20.25.0)
Requirement already satisfied: joblib>=1.1.1 in
./anaconda3/lib/python3.11/site-packages (from scikit-learn-
>factor_analyzer) (1.2.0)
```



```

Requirement already satisfied: threadpoolctl>=2.0.0 in
./anaconda3/lib/python3.11/site-packages (from scikit-learn-
>factor_analyzer) (2.2.0)
Requirement already satisfied: setuptools in
./anaconda3/lib/python3.11/site-packages (from nodeenv>=0.11.1->pre-
commit->factor_analyzer) (68.0.0)
Requirement already satisfied: six>=1.5 in
./anaconda3/lib/python3.11/site-packages (from python-dateutil>=2.8.1-
>pandas->factor_analyzer) (1.16.0)
Requirement already satisfied: distlib<1,>=0.3.7 in
./anaconda3/lib/python3.11/site-packages (from virtualenv>=20.10.0-
>pre-commit->factor_analyzer) (0.3.8)
Requirement already satisfied: filelock<4,>=3.12.2 in
./anaconda3/lib/python3.11/site-packages (from virtualenv>=20.10.0-
>pre-commit->factor_analyzer) (3.13.1)
Requirement already satisfied: platformdirs<5,>=3.9.1 in
./anaconda3/lib/python3.11/site-packages (from virtualenv>=20.10.0-
>pre-commit->factor_analyzer) (4.1.0)
Note: you may need to restart the kernel to use updated packages.

```

```

from factor_analyzer import FactorAnalyzer
fa = FactorAnalyzer(n_factors=6, rotation='varimax')
fa.fit(wine_data)

```

```

FactorAnalyzer(n_factors=6, rotation='varimax', rotation_kwargs={})

```

```

loadings = fa.loadings_
print(loadings)

```

```

[[ 0.90466243 -0.17139493  0.00734408  0.10618959 -0.00973289
 0.33408798]
 [-0.21643924 -0.00388616 -0.1203012  -0.82682052 -0.00306909
 0.05801959]
 [ 0.63502215  0.01699186  0.11657531  0.45942255  0.18099737
 0.17809563]
 [ 0.05614685  0.19472982  0.07325906 -0.00904773  0.03211188
 0.42661129]
 [ 0.11908949  0.00314247 -0.14949922 -0.09390824  0.97518441
 0.02481013]
 [-0.11688089  0.67970513 -0.01580755  0.03958641  0.01391335
 0.10074816]
 [ 0.05293168  0.9803989  -0.12050225 -0.07679366  0.01996967
 0.07633005]
 [ 0.40169032 -0.0717991  -0.40504418 -0.00877127  0.08246313
 0.82511265]
 [-0.73868622 -0.02083286  0.11996531 -0.11679698 -0.17388377 -
 0.00935861]
 [ 0.14072697  0.04332096  0.11181496  0.30013261  0.40471606
 0.0918936 ]
 [-0.09262946 -0.07403897  0.98289388  0.12097397 -0.05575862 -

```

```

0.05609395]
[ 0.06013144 -0.09256672  0.44165983  0.40601841 -0.00336635 -
0.01670899]]

# Factor loadings matrix
factor_loadings = np.array([
    [0.90466244, -0.17139493, 0.00734408, 0.10618959, -0.00973289,
0.33408798],
    [-0.21643923, -0.00388616, -0.12030119, -0.82682051, -0.00306909,
0.05801959],
    [0.63502214, 0.01699186, 0.1165753, 0.45942256, 0.18099737,
0.17809563],
    [0.05614685, 0.19472982, 0.07325906, -0.00904773, 0.03211188,
0.4266113],
    [0.11908949, 0.00314247, -0.14949922, -0.09390823, 0.97518441,
0.02481013],
    [-0.11688089, 0.67970512, -0.01580755, 0.03958641, 0.01391335,
0.10074816],
    [0.05293168, 0.98039891, -0.12050225, -0.07679366, 0.01996967,
0.07633005],
    [0.40169033, -0.0717991, -0.40504419, -0.00877128, 0.08246313,
0.82511265],
    [-0.73868621, -0.02083286, 0.11996531, -0.11679699, -0.17388377, -
0.00935861],
    [0.14072697, 0.04332096, 0.11181496, 0.30013261, 0.40471605,
0.09189359],
    [-0.09262946, -0.07403897, 0.98289388, 0.12097397, -0.05575862, -
0.05609394],
    [0.06013144, -0.09256672, 0.44165982, 0.40601841, -0.00336635, -
0.01670899]
])

eigenvalues = np.linalg.eigvals(factor_loadings @ factor_loadings.T)
cumulative_variance = np.cumsum(eigenvalues) / np.sum(eigenvalues)

# Print the cumulative total variance
print("Cumulative Total Variance Explained:")
for i, variance in enumerate(cumulative_variance, 1):
    print(f"Factor {i}: {variance:.4f}")

Cumulative Total Variance Explained:
Factor 1: 0.3455+0.0000j
Factor 2: 0.5870+0.0000j
Factor 3: 0.7550+0.0000j
Factor 4: 0.8774+0.0000j
Factor 5: 0.9287+0.0000j
Factor 6: 1.0000+0.0000j
Factor 7: 1.0000+0.0000j

```

```

Factor 8: 1.0000+0.0000j
Factor 9: 1.0000+0.0000j
Factor 10: 1.0000+0.0000j
Factor 11: 1.0000+0.0000j
Factor 12: 1.0000+0.0000j

fa.get_factor_variance()

(array([2.0413234 , 1.51230688, 1.42270252, 1.20562616, 1.18942355,
        1.03805694]),
 array([0.17011028, 0.12602557, 0.11855854, 0.10046885, 0.09911863,
        0.08650474]),
 array([0.17011028, 0.29613586, 0.4146944 , 0.51516325, 0.61428188,
        0.70078662]))

```

#Conclusion

Monte Carlo Markov Chain (MCMC) and Variance Reduction:

The code estimates π using MCMC and compares it with Importance Sampling (variance reduction). Both methods can be useful for various applications and add to probabilistic modelling.

Simulated Card Drawing:

provides insights into the possibility of particular card combinations by simulating the probability of obtaining a pair in a five-card hand.

Analysis of Air Quality Data:
examines data on air quality, analysing conditional probability calculations and time series displays. A basis for comprehending the frequency of elevated carbon monoxide levels is provided by prior probability estimation.

Data Visualisation and Analysis for Wine Quality:

shows the connections between different characteristics and wine quality. examines the conditional likelihood of fine wine given its alcohol content.

Normalcy and Chi-Square Tests:

uses the chi-square test to determine whether two categorical variables are independent. evaluates the distribution of a numerical quantity by performing a normalcy test on it.

Factor Evaluation:
uses factor analysis to find hidden components in the information. uses a Scree plot and statistical tests (KMO and Bartlett's Sphericity) to inform factor selection. To sum up, the code presents a wide range of statistical analyses and simulations that offer insightful information about various data points. The particular dataset, as well as the underlying study questions or aims, will determine further interpretation and findings.

#Reference

1] <https://sports-statistics.com/sports-data/fifa-2022-dataset-csvs/>

2] <https://www.kaggle.com/datasets/fedesoriano/air-quality-data-set>

3] <https://www.kaggle.com/datasets/ruthgn/wine-quality-data-set-red-white-wine>