

```

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
import geopandas as gpd
import folium

# Example: Analyzing Health Equity and Identifying Disparities in Healthcare Access

# Step 1: Load the Data
# Dataset contains the following columns:
# 'Region': Geographic region name
# 'Population': Population size of the region
# 'Healthcare_Facilities': Number of healthcare facilities in the region
# 'Median_Income': Median income of the population
# 'Disease_Prevalence': Percentage of population with a specific disease
# 'Uninsured_Rate': Percentage of population without health insurance

data = {
    'Region': ['Region A', 'Region B', 'Region C', 'Region D', 'Region E'],
    'Population': [50000, 30000, 40000, 60000, 25000],
    'Healthcare_Facilities': [5, 3, 4, 6, 2],
    'Median_Income': [55000, 35000, 45000, 65000, 30000],
    'Disease_Prevalence': [15, 25, 20, 10, 30],

```

```
'Uninsured_Rate': [10, 20, 15, 5, 25]  
}
```

```
# Convert data to a DataFrame
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```
df = pd.DataFrame(data)
```

```
# Step 2: Calculate Access-to-Care Metrics
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```
df['Facilities_Per_Capita'] = df['Healthcare_Facilities'] / df['Population']
```

```
df['Disease_Burden'] = df['Disease_Prevalence'] * df['Population'] / 100
```

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# Step 3: Exploratory Data Analysis
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print("Summary Statistics:")
```

```
print(df.describe())
```

```
# Correlation Heatmap
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```
plt.figure(figsize=(8, 6))
```

```
sns.heatmap(df.corr(), annot=True, cmap="coolwarm")
```

```
plt.title("Correlation Matrix")
```

```
plt.show()
```

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# Step 4: Clustering to Identify Underserved Regions
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# Prepare data for clustering
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```
features = ['Facilities_Per_Capita', 'Median_Income', 'Uninsured_Rate',  
'Disease_Prevalence']
```

```
scaler = StandardScaler()
```

```
X = scaler.fit_transform(df[features])
```

```

# Apply K-Means Clustering

kmeans = KMeans(n_clusters=2, random_state=42)

df['Cluster'] = kmeans.fit_predict(X)


# Visualize Clusters

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

sns.scatterplot(x=df['Median_Income'], y=df['Facilities_Per_Capita'], hue=df['Cluster'],
palette='viridis')

plt.title("Clusters of Regions Based on Healthcare Equity")

plt.xlabel("Median Income")

plt.ylabel("Facilities Per Capita")

plt.show()


# Step 5: Geospatial Analysis (Mapping Underserved Regions)

# Assume we have a shapefile of the regions (mock data)

region_geo = gpd.GeoDataFrame({
    'Region': ['Region A', 'Region B', 'Region C', 'Region D', 'Region E'],
    'geometry': gpd.points_from_xy([0, 1, 2, 3, 4], [0, 1, 2, 3, 4]) # Mock coordinates
})

region_geo = region_geo.merge(df, on='Region')


# Create an interactive map

m = folium.Map(location=[2, 2], zoom_start=6)

for _, row in region_geo.iterrows():
    folium.CircleMarker(

```

```

location=(row.geometry.y, row.geometry.x),
radius=10,
color='red' if row['Cluster'] == 1 else 'blue',
fill=True,
fill_opacity=0.6,

popup=f"Region: {row['Region']}<br>Facilities Per Capita:
{row['Facilities_Per_Capita']:.2f}<br>Cluster: {row['Cluster']}"

).add_to(m)

m.save("health_equity_map.html")

```

# Step 6: Insights and Recommendations

# Highlight regions in Cluster 1 (potentially underserved)

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underserved = df[df['Cluster'] == 1]
```

```
print("Underserved Regions:")
```

```
print(underserved[['Region', 'Population', 'Facilities_Per_Capita', 'Median_Income',
'Uninsured_Rate']])
```

# Example Insight:

# Regions in Cluster 1 tend to have lower median incomes, higher uninsured rates, and fewer facilities per capita.

# Recommendation: Prioritize these regions for healthcare resource allocation and policy interventions.

# Save results for reporting

```
df.to_csv("health_equity_analysis_results.csv", index=False)
```