

Data Visualization of MIMIC Demo Dataset

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UT AUSTIN: AI IN HEALTHCARE



Overview



GRAPH 1: MOST
COMMON
MICROORGANISM IN
BLOOD CULTURES



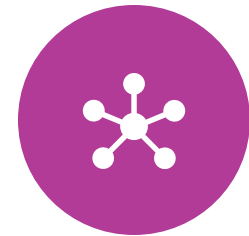
GRAPH 2: TREND OF
PRESCRIPTIONS OVER
TIME



GRAPH 3: TOP 10
PRESCRIBED DRUGS

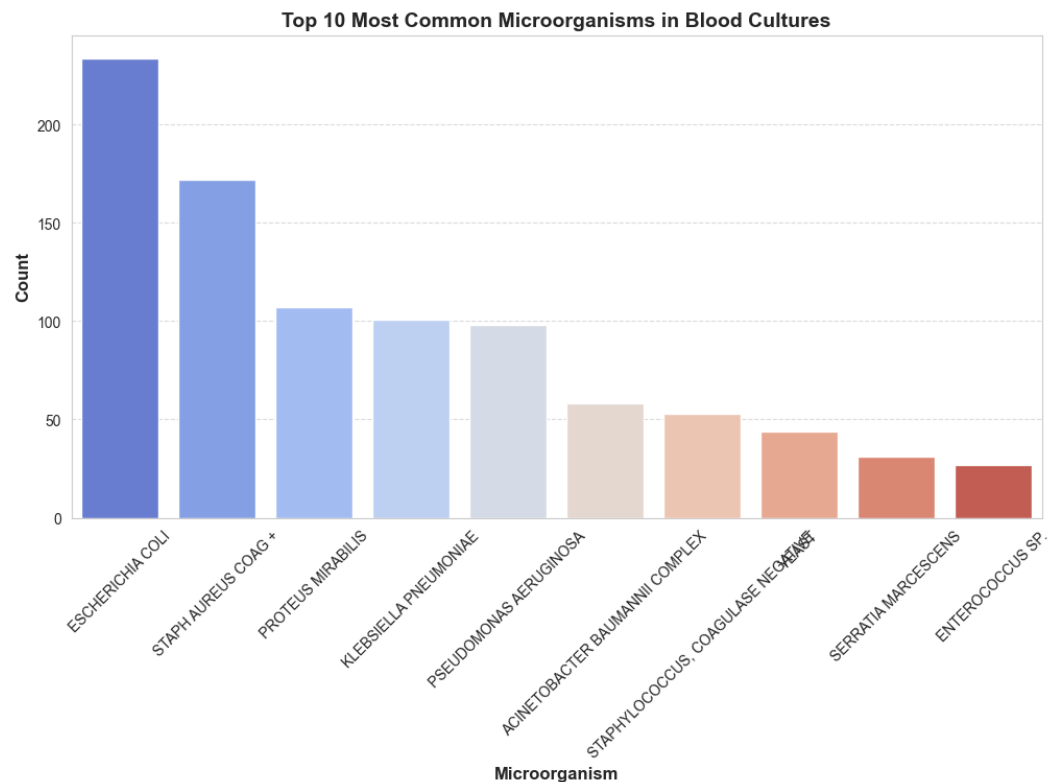


GRAPH 4: ANTIBIOTIC
RESISTANCE
PATTERNS



GRAPH 5: DIAGNOSIS-
BASED PRESCRIPTION
NETWORK GRAPH

Graph 1: Most Common Microorganisms in Blood Cultures



- This visualization highlights the most frequently detected microorganisms in hospital blood cultures. Identifying these microorganisms is critical for guiding treatment strategies and antibiotic stewardship. By analyzing this data, hospitals can proactively prepare for common infections and mitigate the spread of resistant bacteria.

1

Load the dataset: Read the `MICROBIOLOGYEVENTS.csv` file using *pandas*.

2

Process the data: Extract microorganism names and count their occurrences.

3

Select the top 10: Identify the 10 most frequently occurring microorganisms.

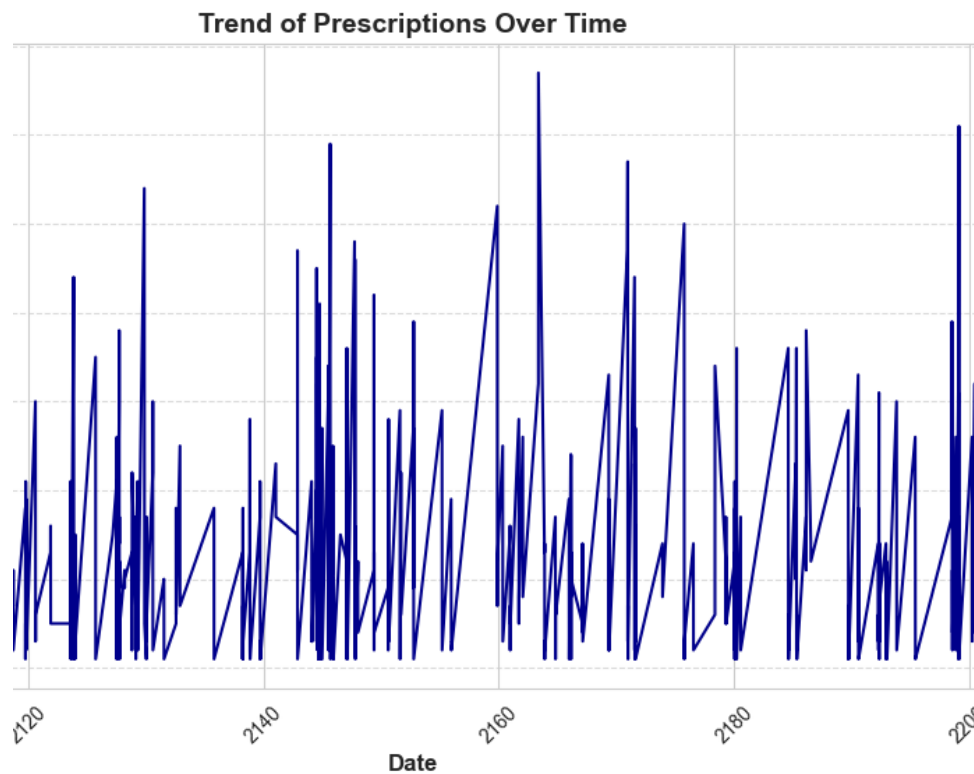
4

Generate the bar chart:

- Use `sns.barplot()` to plot the data.
- Set appropriate labels and a title.
- Rotate x-axis labels for readability.

Graph 1: Step by Step

GRAPH 2: TREND OF PRESCRIPTIONS OVER TIME



- This line graph presents the variation in prescription rates over time. Seasonal trends, public health interventions, and hospital protocols can influence these fluctuations. Understanding these trends assists in inventory management and policy adjustments to optimize patient care.

Graph 2: Step by Step

1

Load the dataset: Read the `PRESCRIPTIONS.csv` file.

2

Convert time format: Ensure the `startdate` column is in datetime format.

3

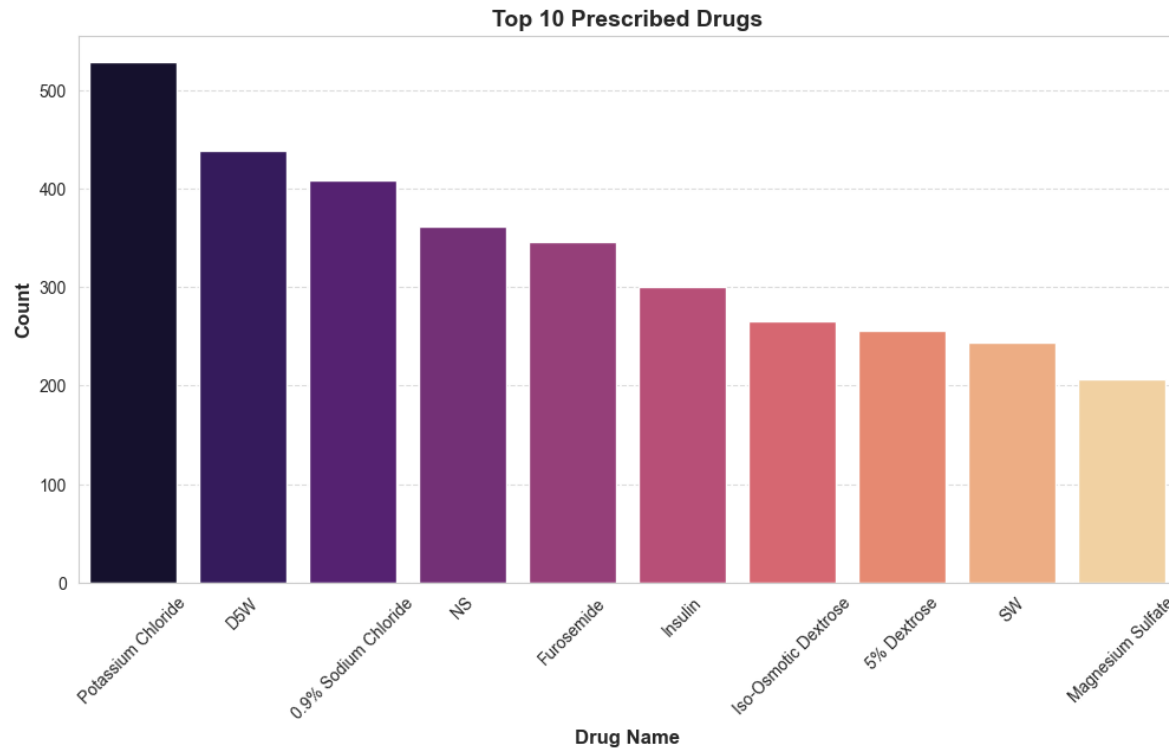
Aggregate prescription counts: Group data by date to compute daily prescription totals.

4

Generate the line plot:

- Use `sns.lineplot()` to plot the trend over time.
- Format the x-axis and y-axis labels appropriately.
- Rotate x-axis labels for clarity.

GRAPH 3: TOP 10 PRESCRIBED DRUGS



- This bar chart illustrates the most frequently prescribed medications. Identifying these drugs helps hospitals manage stock levels and analyze treatment patterns for prevalent conditions.

1

Load the dataset: Read the PRESCRIPTIONS.csv file.

2

Extract prescription data: Identify unique drug names and their frequency.

3

Select the top 10 drugs: Find the most frequently prescribed medications.

4

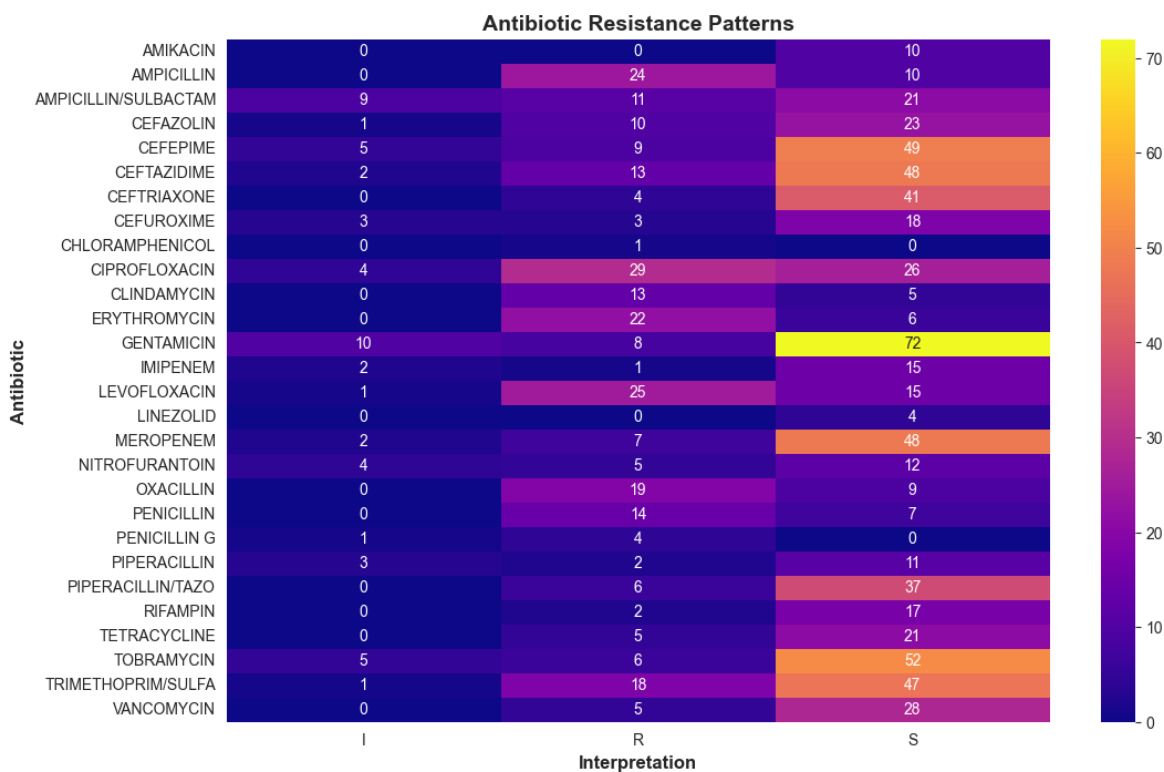
Generate the bar chart:

- Use `sns.barplot()` to plot the drug frequencies.
- Format the visualization with proper labels and title.
- Rotate x-axis labels for clarity.

Graph 3: Step by Step

Graph 4: Antibiotic Resistance Patterns

- This heatmap visualizes antibiotic resistance patterns by displaying interactions between different antibiotics and bacterial resistance interpretations. Hospitals use this data to optimize treatment protocols and mitigate antibiotic resistance



Graph 4: Step by Step

1

Load the dataset: Read the `MICROBIOLOGYEVENTS.csv` file.

2

Filter relevant columns: Keep only antibiotic names and resistance classifications.

3

Create a pivot table:

- Summarize resistance occurrences.
- Index by antibiotic name and classify resistance on columns.

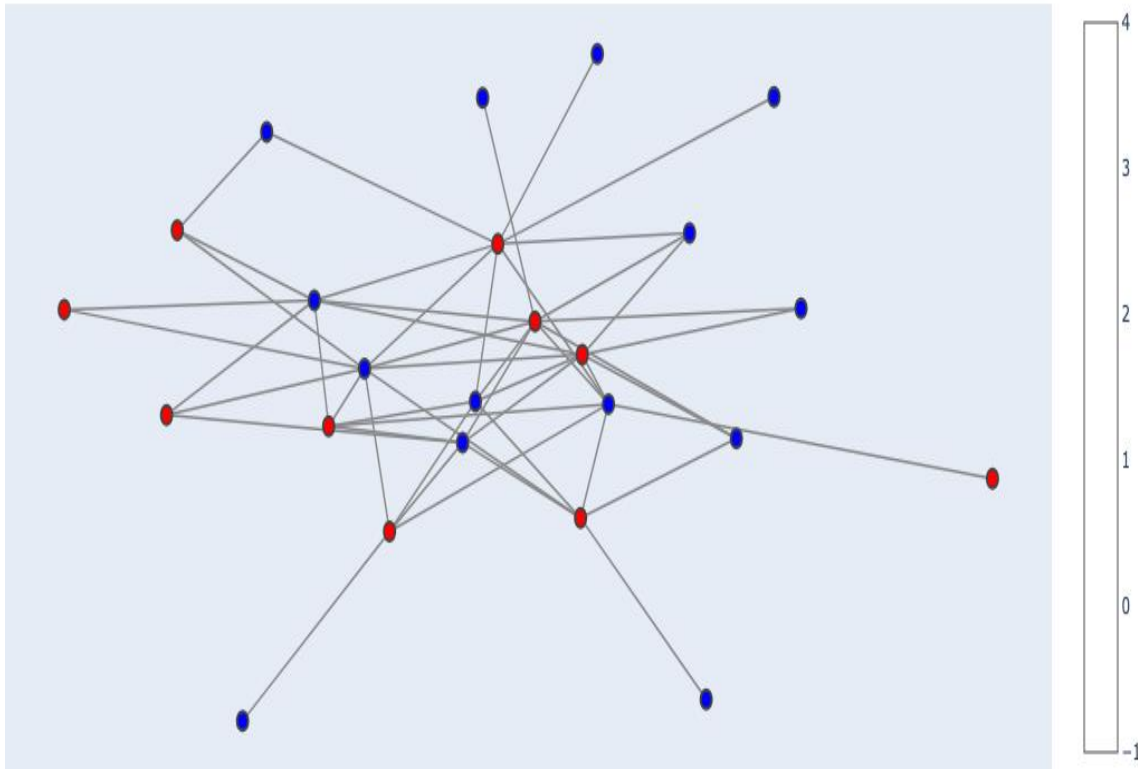
4

Generate the heatmap:

- Use `sns.heatmap()` to visualize the data.
- Annotate the heatmap for better clarity.
- Apply an appropriate color scheme.

Graph 5: Diagnosis-Based Prescription Network Graph

Top 50 Prescription-Diagnosis Network Graph



- A network graph shows drug relationships, co-prescriptions, and interactions, highlighting common treatment patterns. Red nodes represent diagnoses, blue nodes are prescriptions, and thicker lines indicate stronger links. Some drugs treat many conditions, while others are for specific cases.

Graph 5: Step by Step

1

Load the dataset: Read the `PRESCRIPTIONS.csv` file using pandas.

2

Extract relevant data: Identify co-prescribed drugs within patient stays.

3

Create an adjacency list: Construct a list of drug pairs that are frequently prescribed together.

4

Generate the network graph:

- Use `networkx` to construct the graph structure.
- Apply `nx.spring_layout()` for node positioning.
- Use `nx.draw()` to visualize the connections.