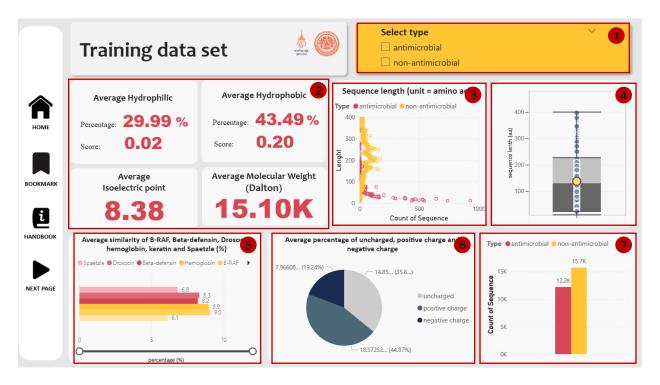
Handbook for dashboard



This main dashboard provides a comprehensive overview of the dataset, offering visual summaries and key statistics for both **antimicrobial** and **non-antimicrobial** types. Users can filter between the two types using the selection panel at the top right (1). The dashboard includes the following components:

1. Filter Panel (1):

Enables users to select and compare data between antimicrobial and non-antimicrobial types.

2. Key Statistics (2):

Displays overall averages, including:

- o Hydrophilic percentage and score
- Hydrophobic percentage and score
- Average isoelectric point
- Average molecular weight (in Daltons)

3. Sequence Length Distribution (3):

A scatter plot illustrating the distribution of sequence lengths for both types across the dataset.

4. Box Plot of Sequence Length (4):

Highlights the statistical range and distribution of amino acid sequence lengths, including median, quartiles, and outliers.

5. Similarity Comparison (5):

A bar chart comparing average similarity (%) to known proteins such as B-RAF, Beta-defensin, Drosocin, Hemoglobin, Keratin, and Spaetzle.

6. Charge Composition (6):

A pie chart showing the average proportion of uncharged, positively charged, and negatively charged amino acids in the sequences.

7. Sequence Count by Type (7):

A bar chart summarizing the total number of sequences for each type.



This dashboard presents insights focused on target bacteria types (Gram-, Gram+, and both). Key components include:

8. Bacteria Type Filter:

Users can select between Gram-, Gram+, or both groups to filter the data.

9. Hydrophilic vs. Hydrophobic Percentage:

A bar chart is used to compare the average percentage of hydrophilic and hydrophobic sequences across each bacteria group.

10. Key Averages:

Displays the overall average isoelectric point and molecular weight (in Daltons) for the selected bacteria group.

11. Sequence Distribution Tree:

This tree diagram shows the number of sequences categorized by antimicrobial type and bacteria group.

12. Tabbed Score Comparisons:

a. Average scores of hydrophilic and hydrophobic properties.

- b. Graph showing the relationship between the percentage of positive charge and the number of peptides.
- c. Graph showing the relationship between the percentage of negative charge and the number of peptides.
- d. Graph showing the relationship between the percentage of uncharged residues and the number of peptides.

13. Sequence Length Scatter Plot:

The scatter plot displays sequence lengths by bacteria group, showing spread and density.

14. Sequence Length Box Plot:

Summarizes the distribution (median, quartiles, outliers) of amino acid sequence lengths.