

README

1. A single python file is used for all three parts.
2. Place the python file and the 'breast-cancer.data' dataset in the same directory.
3. For running the file, type `python3 <filename>.py` on the terminal.
4. The output for these codes will be printed directly on the terminal. The plots will be displayed automatically.
5. However, close the graph after it's printed to get further results such as printing the final tree.