



**UTM**  
UNIVERSITI TEKNOLOGI MALAYSIA

**FACULTY OF COMPUTING**  
UTM Johor Bahru

## **SECB3203-01(PROGRAMMING FOR BIOINFORMATIC)**

Semester 01 2025/2026  
Section 01

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### **Progress 1**

**Faculty of Computing**

**Dataset:**

<https://www.kaggle.com/datasets/miadul/tuberculosis-x-ray-dataset-synthetic>

**Project Title:**

**Tuberculosis Disease Classification Using Synthetic Chest X-Ray  
Images and Machine Learning Techniques**

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# **1. Software and Hardware Requirements**

## **1.1 Software Requirements**

The following software will be used to develop and implement the tuberculosis classification system:

- Operating System:  
Windows 10 / Windows 11
- Programming Language:  
Python 3.x
- Development Environment:
  - Jupyter Notebook
  - Anaconda Navigator
- Python Libraries:
  - NumPy – numerical computations
  - Pandas – data manipulation and preprocessing
  - Matplotlib – data visualization
  - Scikit-learn – machine learning models and evaluation
  - OpenCV / PIL – image processing
  - TensorFlow / Keras (if deep learning is applied)
- Version Control:
  - GitHub for code management and documentation
- Dataset Source:
  - Synthetic Tuberculosis Chest X-Ray Dataset from Kaggle

## **1.2 Hardware Requirements**

The project will be implemented using student-level computing resources:

- Processor:  
Intel Core i5 or equivalent
- RAM:  
Minimum 8 GB
- Storage:  
At least 20 GB free disk space
- Display:  
Standard monitor with minimum  $1366 \times 768$  resolution
- Internet Connection:  
Required for dataset download, library installation, and GitHub access

## **2. Flowchart of the Proposed Approach**

The flowchart below shows the overall workflow of the project from data collection to model evaluation.

