

# Configuration Manual

MSc Research Project  
Masters in Artificial Intelligence

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# Configuration Manual

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## 1 System Requirements

### Hardware

**Device:** Apple MacBook Pro equipped with an M4 chip

**Processor:** M2 Chip, 8-Core CPU

**Memory:** 16 GB Unified RAM

**Storage:** 512 GB SSD

**GPU:** Integrated Apple M2 GPU

### Software

**Operating System:** macOS Sonoma 14.x

**Python Version:** 3.11.9 (managed with pyenv)

**Integrated Development Environment (IDE):** Visual Studio Code (latest version)

### Libraries & Frameworks

TensorFlow / Keras, NumPy, Pandas, Matplotlib, Seaborn, OpenCV, Scikit-learn, pyenv + virtualenv, Jupyter / Colab (for experiments)

### Dataset Source

**Name:** Brain Tumor Segmentation Dataset

**Source:** Kaggle: Brain MRI with Masks (Figshare) 3,000 image-mask pairs (binary segmentation)

## 2 Installation & Environment Setup

All steps were executed on macOS Sonoma via Terminal and VS Code.

1. I installed Homebrew, a macOS package manager, to facilitate the installation of Python and the other dependencies. `./bin/bash -c "$(curl -fsSL https://raw.githubusercontent.com/Homebrew/install/HEAD/install.sh)"`

**2. I installed pyenv to manage multiple Python versions and subsequently installed Python 3.11.9.**

```
brew install pyenv  
pyenv install 3.11.9  
pyenv global 3.11.9
```

**3. A virtual environment was created and activated to ensure the isolation of project dependencies.**

```
python3 -m venv venv  
source venv/bin/activate
```

**4. All necessary Python libraries pertinent to deep learning, fairness evaluation, and explainability were installed.**

```
pip install tensorflow keras numpy pandas matplotlib seaborn opencv-python shap lime  
scikit-learn
```

**5. Dataset:** Extracted into archive\_1/images/ and archive\_1/masks/ folders.

### 3 Execution Instructions

The following Python scripts were used to build and evaluate the project:

**1. preprocessing.py**

Prepares the input MRI images and segmentation masks.  
Converts to grayscale, resizes to 128x128, normalizes pixel values.

**Output:**

X.npy and y.npy → preprocessed data arrays.

**2. unet\_train.py**

Implements and trains a U-Net architecture for segmentation.  
Model is trained on ~2,500 images with 500 used for validation.  
Metrics: Accuracy, Dice Score, Intersection-over-Union (IoU).

**Output:**

unet\_model.keras → Trained U-Net model  
Training summary, model architecture  
Final validation results

**3. explain\_gradcam.py**

Applies Grad-CAM on the trained U-Net model to visualize focused regions.  
Used final convolutional layer (e.g., conv2d\_13) for heatmap generation.

**Output:**

gradcam\_overlay.png  
Heatmaps superimposed on original MRI inputs  
Highlights tumor focus zone

#### 4. `transformer_train.py` (*Optional – for model comparison*)

Under development: Implements Swin-UNet or Vision Transformer (ViT).

Expected to follow the same evaluation pipeline as U-Net.

## References

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