

Lab5: Image segmentation (MONAI)

3099704 AI for Digital Health (2025/2)

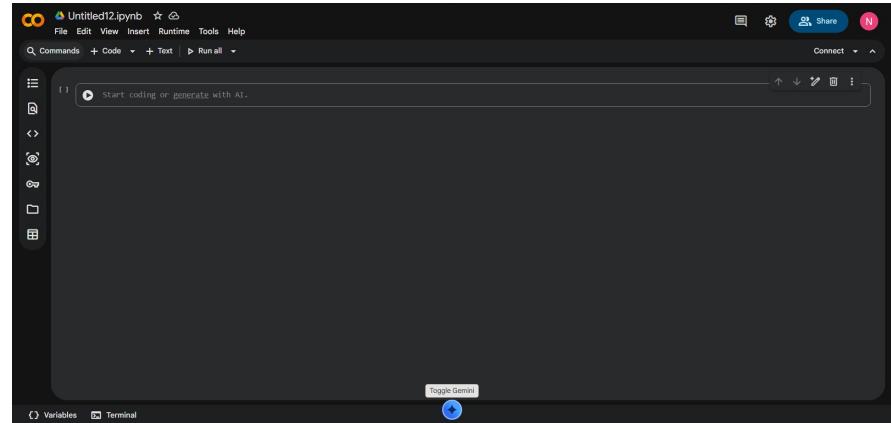
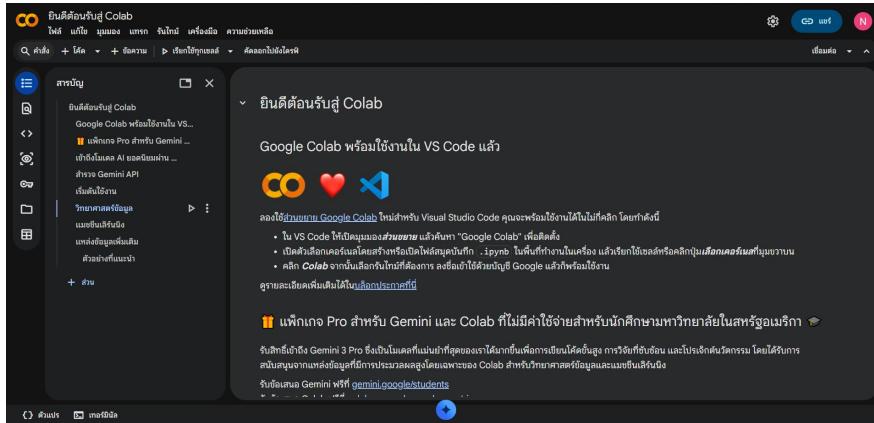
Objective

- Create segmentation model
- Use the **MONAI library** to build deep learning model (UNet)



Material

- With **Google Colab**, you don't need to install any software. All you need is a Google account, and you can start using it right away. Simply visit: <https://colab.research.google.com/> or select NEW NOTEBOOK to start a new file.



Pytorch VS MONAI

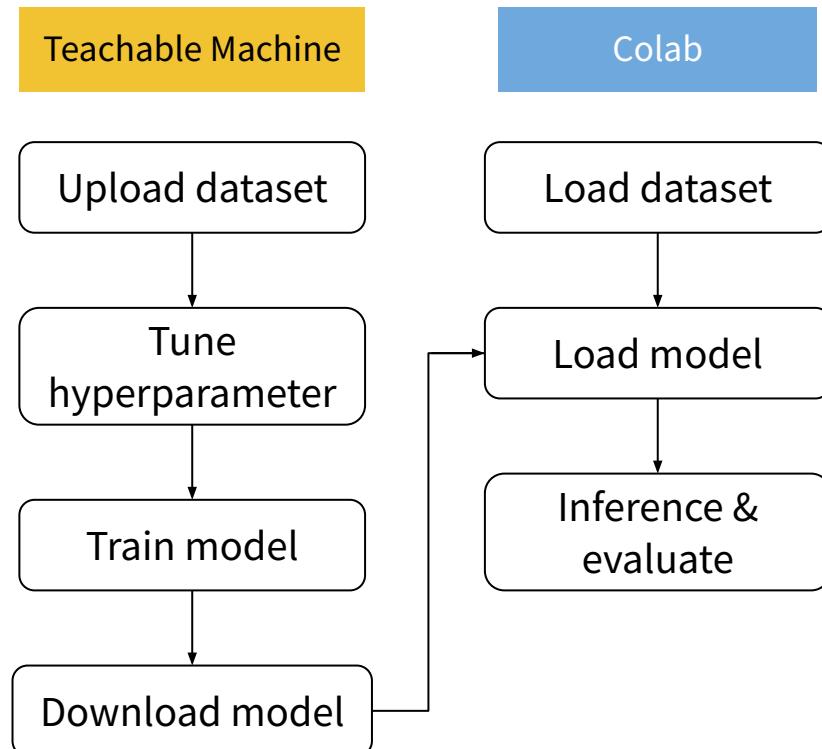
PyTorch is a general-purpose framework for deep learning, while MONAI is built on PyTorch but adds specialized functions for medical imaging tasks, such as

- **Data Loading:** CacheDataset supports big data and caching
- **Transforms:** monai.transforms supports 3D medical volume
- **Networks:** build-in models, such as UNet and SegResNet
- **Inferers:** sliding_window_inference enable inference 3D volumes by 2D models



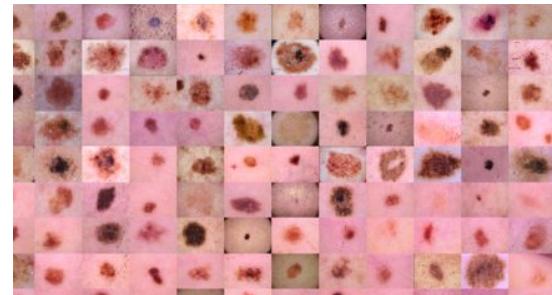
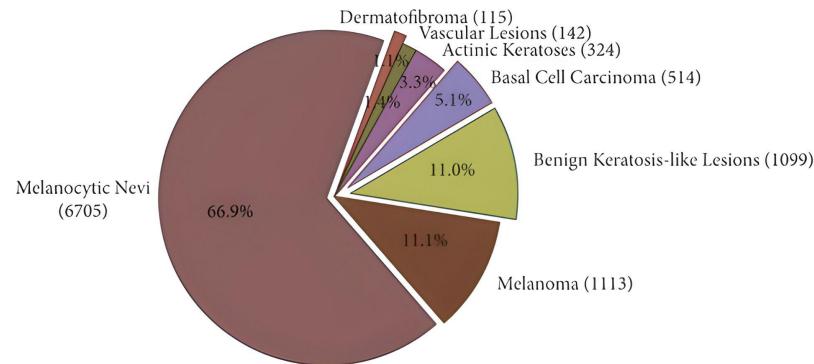
Lab5.1: MobileNet (2D segmentation)

In this lab, you will create an image classification model using **Teachable Machine** and then evaluate its performance using a **Google Colab notebook**.



Dataset: Skin Cancer MNIST (HAM10000)

- The dataset consists of 10015 images with 10013 labeled objects belonging to 7 skin cancer classes.
- The data contains image in JPG format and documents in JSON format
- In the experiment, we reduced the amount of data and formatted it to simplify the experiment.



Lab5.1: UNet (2D segmentation)

In this lab, you will create and evaluate an skin cancer segmentation model (UNet) using the **MONAI library**. Code can be executed in [Lab_5_1_MONAI\(2Dsegmentaton\)](#) on Google Colab.

This notebook will consist of:

- 1) Setup
- 2) Load Data & Set Transforms
- 3) Define Model & Set Parameter
- 4) Train Model
- 5) Inference & Evaluate & save

The screenshot shows a Google Colab interface with a notebook titled "Lab_5_1_MONAI(2Dsegmentaton).ipynb". The first cell contains the following text and code:

Lab 5.1 – Skin cancer segmenataion: UNet (MONAI)

This notebook implements training of a 2D UNet from MONAI library (https://monai-dev.readthedocs.io/en/stable/_modules/monai/networks/nets/unet.html#UNet) to segment skin cancer. The training code is also customizable to enable training with a different target. In this notebook, we are using the HAM10000 Dataset (<https://www.kaggle.com/datasets/surajhuwalela/ham10000-segmentation-and-classification?select=masks>)

This example code will consist of:

1. Setup
2. Load Data & Set Transforms
3. Define Model & Set Hyper Parameter
4. Train Model
5. Inference & Evaluate & Save

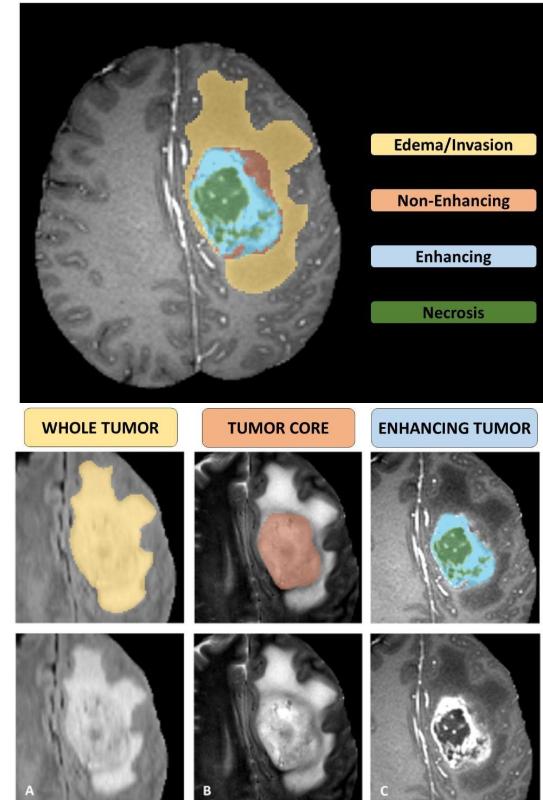
1) Setup

The code below download dataset, imports all required libraries and defines utility functions that will be used in the rest of this notebook.

```
# Download Dataset
!wget https://github.com/pvateekul/digitalhealth-ai2025/raw/main/dataset/Ham10000_segment.zip
!unzip -q -o 'Ham10000_segment.zip'
```

Dataset: Brain Tumor Segmentation 2020 Dataset (BraTS2020)

- BraTS 2020 is dataset for brain tumor segmentation, consisting of T1, T1ce, T2, FLAIR with tumors, which contains image in [nii.gz](#) format
- BraTS 2020 includes **369 training cases**, with further validation/test subjects provided for evaluation.
- In the experiment, we reduced the amount of data and formatted it to simplify the experiment.

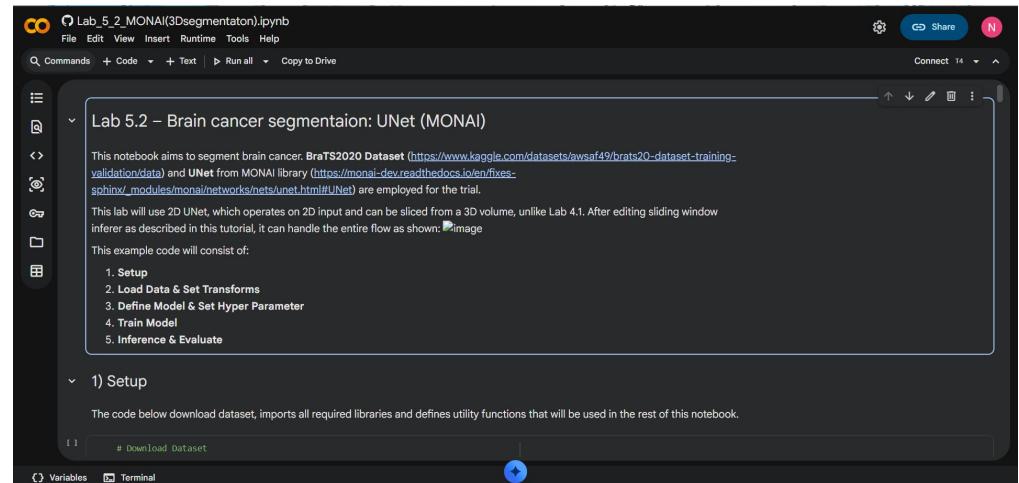


Lab5.2: UNet (3D segmentation)

In this lab, you will create and evaluate an brain tumor segmentation model (UNet) using the **MONAI library**. Code can be executed in [Lab_5_2_MONAI\(3Dsegmentaton\)](#) on Google Colab.

This notebook will consist of:

- 1) Setup
- 2) Load Data & Set Transforms
- 3) Define Model & Set Parameter
- 4) Train Model
- 5) Inference & Evaluate & save



The screenshot shows a Google Colab interface with a dark theme. The title bar reads "Lab_5_2_MONAI(3Dsegmentaton).ipynb". The main content area displays a section titled "Lab 5.2 – Brain cancer segmentaion: UNet (MONAI)". It includes a brief description of the task, mentioning the Brats2020 Dataset and MONAI library. Below this, a list of steps is provided: 1) Setup, 2) Load Data & Set Transforms, 3) Define Model & Set Hyper Parameter, 4) Train Model, 5) Inference & Evaluate. The "1) Setup" section is expanded, showing a code cell starting with "# Download Dataset". At the bottom, there are tabs for "Variables" and "Terminal".