

# Guide: Preparing Data and Running the Thin-Section Mineral Segmentation Workflow

## Critical Requirement

Optical input images and corresponding labels must be **aligned pixel-by-pixel**, and labels must be ready for training, either as class IDs or as colors consistently mapped to class IDs.

## 1 What This Workflow Does

This workflow trains a deep-learning model to generate mineral maps from optical thin-section images. The model learns a mapping between:

- **Input:** Optical microscopy images (single-angle or multi-angle polarized light)
- **Output:** Mineral label masks (ground truth)

Once trained, the model can predict mineral distributions for new thin-section images or for extracted image patches.

## 2 Hardware Recommendation for Training

Model training is computationally intensive, particularly when using high-resolution patches, multiple optical angles, and deep encoder backbones. For efficient training, the use of a high-end GPU is strongly recommended.

- GPUs with at least **16–24 GB of VRAM** are recommended for multi-angle training.
- Systems with limited GPU memory may require reducing batch size, patch size, or the number of optical angles.
- CPU-only training is supported just for debugging or small-scale tests.

### 3 What You Need Before Running Anything

Before running any code, ensure that the following requirements are met:

- Python version 3.9 or newer
- All required Python packages installed (see `requirements.txt`)
- A dataset organized by sample
- At least **one aligned optical angle per sample** (minimum requirement)
- A defined list of mineral classes to be predicted
- Labels that are either already aligned or can be generated using the provided scripts

**Important:** If alignment or label generation is incorrect, model results will be meaningless, even if the code executes without errors.

### 4 Dataset Layout on Disk

The dataset must follow the structure shown below:

```
data/  
  sample_name_1/  
    image/    # optical images (*.tif)  
    label/    # mineral label masks (*.tif)  
  sample_name_2/  
    image/  
    label/
```

#### Notes:

- The folder name `data` must match the value specified for `data_root` in `config.yaml`.
- Sample folder names must exactly match those listed under `samples` in the `data` section of `config.yaml`. Names are case-sensitive.

Each sample folder represents one thin section (or one physical specimen). The `image/` folder contains optical images (PPL, XPL, or multiple angles), while the `label/` folder contains a single-channel label mask used for training.

### 5 Data Preparation: Conceptual Explanation

The model learns pixel-level correspondences between optical images and mineral labels. Therefore:

- Each pixel in the label must correspond to the same physical location in the optical image.
- Labels must be aligned in scale, orientation, and spatial resolution.

## 5.1 Alignment Requirement (Very Important)

If the data are not already aligned, the following scripts must be used during data preparation:

- `input_alignments.py`: aligns optical inputs and reference maps
- `labelgeneration.py`: generates training-ready label masks

**Minimum requirement:** At least one optical angle with ground truth labels must be aligned. This aligned angle can then be used to align additional angles or to generate the final label mask.

## 6 Label Generation and Color Mapping

In many cases, labels originate from sources such as  $\mu$ XRF or mineral maps that use colors or categorical values. These labels must be converted into a **single-channel class-ID mask** prior to training.

This conversion is handled by `labelgeneration.py`.

### Required Edits in `labelgeneration.py`

Users must:

- Define input paths, output visualization paths, and output label paths
- Edit `class_names` and corresponding `hex_colors` (or values) to match the label data
- Ensure that each mineral class maps to a unique integer class ID

Incorrect color or class mapping will result in mislabeled training data and invalid model predictions.

## 7 Editing `config.yaml`

All workflow settings are controlled through a single file, `config.yaml`. Most users do not need to modify any Python source code.

Section	Purpose	Options
project	Experiment metadata	–
data	Dataset and loading	stack, separate
split	Data splitting strategy	fraction, sample_holdout, fixed_manifest
model	Architecture and classes	UNet, UNet++, DeepLabV3+
train	Training control	–
eval	Metrics and visuals	–
inference	Prediction settings	–

More details are provided in the repository README.

## 8 Running the Code

### Step A: Install Dependencies

```
pip install -r requirements.txt
```

### Step B: Run a Quick Smoke Test

```
python traintest.py --config config.yaml
```

### Step C: Train the Model

```
python -m src.train --config config.yaml
```

### Step D: Evaluate the Model

```
python -m src.evaluate --config config.yaml --checkpoint models/best.ckpt
```

### Step E: Run Inference

```
python inference.py --config config.yaml
```

## 9 Patch Naming for Inference

```
patch_0001_PPL.tif  
patch_0001_XPL40.tif  
patch_0001_XPL50.tif  
patch_0001_XPL60.tif  
...
```

## 10 Troubleshooting Checklist

- **Files not found:** Verify dataset paths and sample names
- **Incorrect results:** Verify alignment and label color mapping
- **Out-of-memory:** Reduce batch size or workers

**Best practice:** Maintain a small debug dataset and validate the full pipeline before scaling up.