SRSML24: STM Machine Learning Module

Steven R. Schofield

April 21, 2025

Overview

This module provides tools for machine learning analysis of scanning tunnelling microscopy (STM) data, including autoencoder models, clustering tools, and STM-specific preprocessing.

Getting the Code

Clone the repository from GitHub:

```
git clone https://github.com/srschofield/SRSML24.git
```

Installation

It is recommended to create a clean Python environment using conda. The following steps assume you are working on a macOS system with Apple Silicon:

```
# create and activate environment
conda create --name srsml24 python=3.8 -y
conda activate srsml24

# install packages
pip install -r requirements-macos.txt
```

Known Working Configuration

This module has been tested and is known to work with the following configuration on macOS 15.0.1 (Apple Silicon, M3 Pro chip):

| Package | Version |
|------------------|-----------|
| python | 3.8 |
| tensorflow-macos | 2.13.0 |
| tensorflow-metal | 1.0.1 |
| numpy | 1.24.3 |
| pandas | 2.0.3 |
| matplotlib | 3.7.5 |
| scikit-learn | 1.3.2 |
| scipy | 1.10.1 |
| opencv-python | 4.11.0.86 |
| Pillow | 10.4.0 |
| joblib | 1.4.2 |
| jupyter | 1.1.1 |
| ipykernel | 6.29.5 |
| keras-core | 0.1.5 |
| spiepy | 0.2.1 |
| access2thematrix | 0.4.4 |

Table 1: Verified package versions for macOS (Apple Silicon) environment

These packages can be installed using the requirements-macos.txt file. The Python version is critical: other versions may cause compatibility issues with TensorFlow or other packages on Apple Silicon.

Python Files

- data_prep.py Functions for data preparation, including slicing STM images into windows and saving them in efficient formats.
- model.py Defines convolutional autoencoder and UNET-style models.
- utils.py Utility functions for loading/saving models, feature arrays, and results.

License

This work is licensed under the Creative Commons Attribution-NonCommercial-ShareAlike 4.0 International License (CC BY-NC-SA 4.0). You may share and adapt the material for non-commercial purposes, provided that appropriate credit is given and any derivatives are licensed under identical terms.

Parameter Summary

| Parameter | Description |
|---|---|
| General | |
| job_name | Label for the run, it will be the folder name for output. |
| verbose | If True, enables more detailed print output. |
| Matrix data file processing | 5 |
| flatten_method | Method used to flatten STM images before analysis. Options |
| | are 'none', 'iterate_mask', 'poly_xy'. |
| pixel_density | All images will be converted to this pixel density (px/nm). |
| pixel_ratio | Images that have ratio of fast/slow scan direction less than this |
| | will be discared. Setting to 1 means only complete (square) |
| | images are kept. |
| data_scaling | Multiplicative factor for z-height data. Setting to 1.e9 means |
| | that the range 0–1 (used for training) corresponds to 1 nm. |
| Window generation | |
| window_size | Side length of square image windows (in pixels). |
| window_pitch | Spacing between adjacent windows during tiling. |
| Data saving | |
| (Should remain defaults but options can be useful for examining data manually.) | |
| save_windows | If True, saves image windows as .npy files (True). |
| together | If True, saves windows per image in a single file (True). |
| save_jpg | If True, saves full STM images as JPGs (False). |
| collate | If True, flattens directory structure into one folder. (False). |
| save_window_jpgs | If True, saves image windows as JPGs. (False) |
| ${f Autoencoder}$ | |
| model_name | Label used to save and load the trained autoencoder model. |
| batch_size | Number of windows per training batch. |
| buffer_size | Size of shuffle buffer. |
| learning_rate | Learning rate for the optimizer. |
| epochs | Number of training epochs. |
| Clustering | |
| cluster_model_name | Name used when saving the clustering model. |
| cluster_batch_size | Number of latent vectors per clustering batch. |
| cluster_buffer_size | Size of buffer for clustering shuffle. |
| num_clusters | Number of clusters to form using KMeans. |
| n_init | Number of initializations for KMeans. |
| max_iter | Max iterations for KMeans convergence. |
| reassignment_ratio | Fraction of centroids reassigned each step. |
| Image prediction | |
| <pre>predict_window_pitch</pre> | Window spacing during prediction step. |
| mtrx_train_data_limit | Max number of training MTRX files to use. |
| mtrx_test_data_limit | Max number of validation MTRX files to use. |
| train_data_limit | Limit on number of training windows. |
| test_data_limit | Limit on number of validation windows. |

Step 1: Converting MATRIX STM Data Files

The initial step in utilizing the SRSML24 module involves converting raw STM data files from the Scienta Omicron MATRIX format into a standardized format suitable for machine learning analysis. This is accomplished using the process_mtrx_files function.

Function Overview

process_mtrx_files(mtrx_paths, save_data_path, **kwargs) is designed to batch process a list of MATRIX (.mtrx) files, performing the following operations:

- Loading Data: Reads each .mtrx file and extracts image data along with associated metadata.
- **Preprocessing:** Applies flattening methods to correct for background variations and rescales images to a consistent pixel density.
- Window Extraction: Divides images into smaller windows of specified size and pitch, facilitating training of machine learning models.
- Saving Outputs: Stores processed windows and optional JPEG representations in a structured directory hierarchy under save_data_path.

This function ensures that STM data is preprocessed consistently, facilitating reliable training and evaluation of machine learning models within the SRSML24 framework. It will create a new directory called "windows" and subfolders under that with the names corresponding to the folders the matrix data was stored in (e.g., "training" or "testing"). Unless the "collate" variable is set, the windowed data will retain the full directory structure (dates, days, etc) of the matrix data. The individual windows for a given folder of matrix data are all stored within a single .npy file, rather than separate .npy files for each window, since this is much more efficient for data saving and retrieving. Two text files are also saved, one has the meta data (STM bias voltage, current, etc.). The other has the coordinates for each window in the dataset. This is useful since these are not uniform at two edges of the original image for the general case where the full image is not perfectly divided by the dimensions of the individual windows.

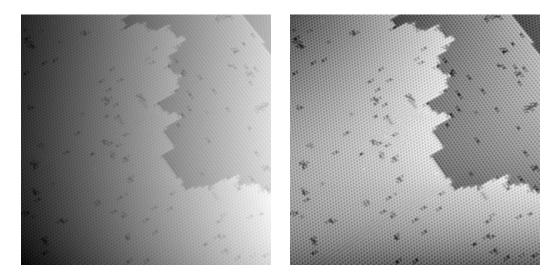


Figure 1: Typical scanning tunnelling microscopy (STM) image. (Left) raw data. (Right) After poly_xy background subtraction.



Figure 2: A sequence of 30×30 pixel windows extracted from the STM image in Fig. 1 with an 8 pixel pitch.