

Project Documentation

Classification of Statistical Pearson diagrams

(△ This project is under active development)

About

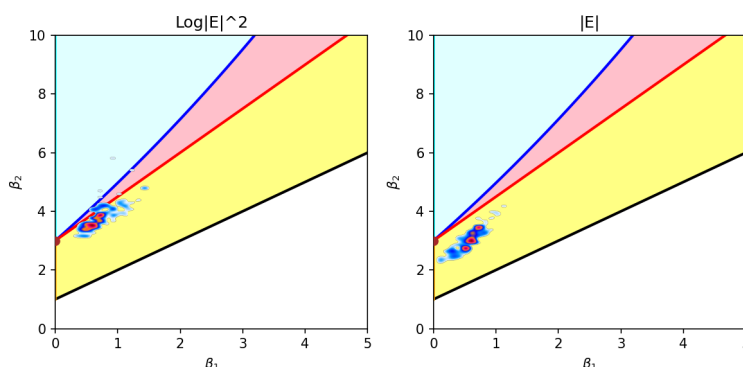
The project is designed for automatic classification of statistical diagrams obtained from plasma turbulence measurement data by Solar Orbiter (RPW-TDS instruments). For details on the physical formulation of the problem, obtaining and analyzing diagrams, see *V.Annenkov, C.Krafft, A.Volokitin, and P.Savoini Statistical properties of beam-driven upper-hybrid wave turbulence in the solar wind*, accepted to **A&A Letters**. The results, including those obtained with the help of this project, are planned to be published in the journal **Journal of Geophysical Research: Machine Learning and Computation**.

What This Project Demonstrates

- Modular project design for ML pipelines
- Deep feature extraction from images
- Unsupervised learning (clustering without labels)
- Data processing and pipeline construction
- Logging, visualization, and result organization

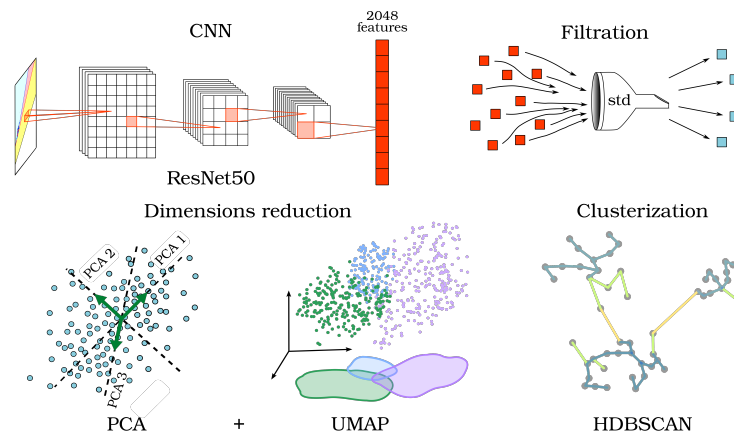
Features of the Project

For an example of typical Pearson charts that need to be classified, see below and in the folder `./images/images_regular_data/`.



The problem: a set of more than 2 thousand statistical diagram images must be classified and analysed. No labeled training data is available.

Image processing: 1. Features extraction from an image by pretrained CNN ResNet50, without classification head; 2. Filtering of low-informative features; 3. Smooth reduction of features space dimension: PCA + UMAP; 4. Clustering and sorting images by HDBSCAN;



The project implements the ability to manually construct a pipeline based on any (reasonable) combination of standardization methods (StandardScaler, Normalizer - sci-kit learn library) and dimension reduction methods (UMAP, PCA). For comparison, clustering methods like DBSCAN and KMeans are also supported. To evaluate unsupervised clustering, Silhouette score and Davies Bouldin score are used. To evaluate cluster stability, Adjusted Rand score is used.

The interactive mode (run `main_interactive.py`) allows you to select a combination of models and their parameters "on the fly".

```

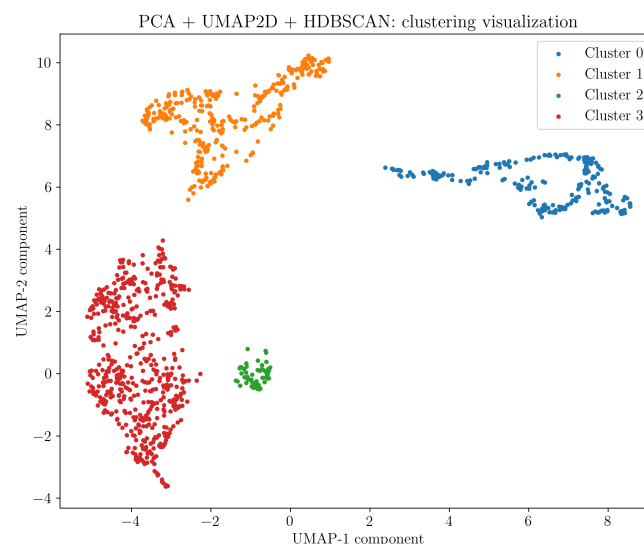
Welcome to the script!
First, put your images in a separate folder (in "/images/") named according to the pattern:
images_(your_specification)
Use "processed" folder for processed images
"data" folder for info metadata
"figures" and "results" for graphic and data results, respectively.
Please enter the name of directory with images > images_example
If folder /mnt/Data/Documents/Projects/pearson-diagram-classification/processed/processed_example already exists would you like to clear its contents? (True or False) >
Enter a path to the file with metadata (or press "Enter" to use default /mnt/Data/Documents/Projects/pearson-diagram-classification/data/SOLO_info_rswf.txt): /data/SOLO_info_rswf_example.txt
To extract features from images located in images_example enter 1.
To load features from a file enter 2.
[1/2] >

```

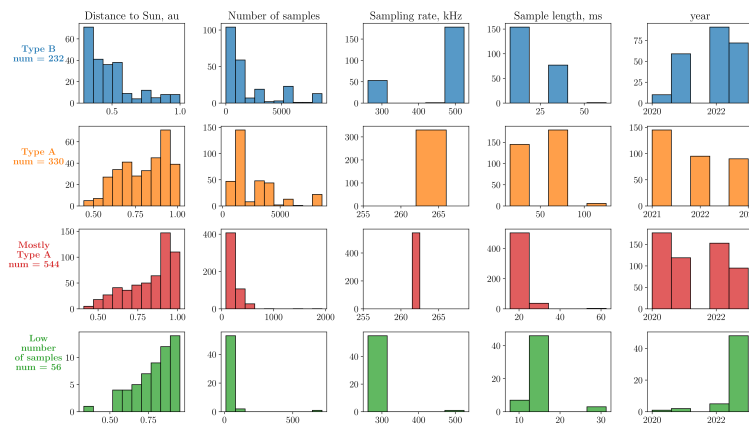
Cluster labels are stored in a resulted DataFrame along with image paths.

Results

The example of clustered data obtained by a standard algorithm: ResNet without a last layer + PCA(with 0.95 dispersion threshold) + UMAP 2D + HDBSCAN, - is below.



Histograms for clusters in the dependence on parameters of measurements:



Structure

Contents

The main part of the project responsible for data processing is packaged in the `mclustering` module. An example of using the module is given in `main.py`. The `manalyse` module is used to visualize the results, an example of using `main_plot.py`.

The `minteractive` module implements interactive (from the console) launch of processing, guiding through the entire process, with the ability to select a specific pipeline. An example of use is presented in `main_interactive.py`, (is in progress)

```
project
├── main.py # basic pipeline
├── main_interactive.py # interactive processing
├── main_plot.py # visualization
├── README.md
├── LICENSE
├── mclustering # the main module for processing
│   ├── __init__.py
│   ├── libfeatures.py # feature extraction
│   ├── libpreprocessing.py # dimension reduction
│   ├── libclustering.py # clustering, clusters visualization and evaluation
│   └── libservice.py # supplementary functions
├── minteractive # interactive module
│   ├── __init__.py
│   ├── libintercative.py # supplementary functions for input/output
│   └── libprocesspipeline.py # automatical launch of the process
├── manalyse
│   ├── __init__.py
│   └── plot_funcs.py # functions for visualization
```

Inputs

`images` folder is for source images to be analysed. The source images are supposed to be placed in a separate subfolder with name starting with `'images_'`.

`data` folder is supposed to contain a metadata-file describing the images.

```
├── images # source images
```

```
| |— images_regular
|— data # for metadata, data description
```

Outputs

Results are supposed to be saved as - .pkl or .json file for a resulted database with labels - results folder - subdirectory in processed folder named as 'processed_{specification as for images folder}'; images are sorted in there in different subdirectories according their label: 'label_0', 'label_1', ... 'noise' - Visual data .pdf/.png should be saved in figures folder.

```
|— results # .pkl/.json resulted database
|— processed
|   |— processed_regular
|   |   |— label_0 # for sorted images with label 0
|   |   |— label_1 # for sorted images with label 1
|— figures # for visual results
```

Other

documentation folder is for .html/.pdf documentation. The documentation is generated semi-automatically with Pdoc.

logs - a folder for full logs per run.

Dependencies

- Python 3.8+
- torch, torchvision
- sklearn
- umap
- hdbscan
- pandas, numpy
- tqdm (progress bar)
- matplotlib, seaborn
- logging
- typing (signatures)
- shutil (copying files)

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- Clustering module: features extraction
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- Interactive module: service functions
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- Visualization