Development project in Machine Learning

TAF MCE

Elsa Dupraz elsa.dupraz@imt-atlantique.fr

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Versioning with GIT

Project description

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- Develop good programming practices
- ► Use standard development tools
- Get used to collaborative work
- ► Work on Machine-Leaning datasets

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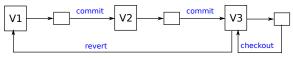
- ► GIT is a versioning system
- It can keep track of the successive code versions
- ► It allows several persons to work on the same files, and can merge the various contributions

Tutorials:

- https://openclassrooms.com/fr/courses/
 1233741-gerez-vos-codes-source-avec-git (Course)
- https://github.com/girliemac/ a-picture-is-worth-a-1000-words/tree/main/git-purr (Pictures)
- https://ohmygit.org/ (Game!)

How GIT works

► Versioning:

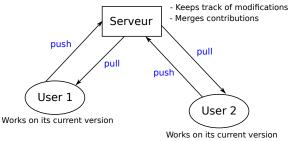


Example: coffee machine

- V1: template (repository organization, readme, etc.)
- ▶ V2: Added function to boil down water
- V3: Added function to pour water
- ▶ Then: Modified function to boil down water, but broke everything!!

How GIT works

► Architecture:

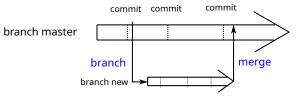


Example: coffee machine

- ▶ User1: works on function to boil down water
- User2: works on function to pour water

How GIT works

► Branches:



Example: coffee machine

- branch master: the machine can only makes coffee
- branch new: we want to add modes to choose between coffee or tea

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Machine Learning Workflow

- 1. Import the dataset
- 2. Clean the data, perform pre-processing
 - ▶ Replace missing values by average or median values
 - Center and normalize the data
- 3. Split the dataset
 - Split between training set and test set
 - ► Split the training set for cross-validation
- 4. Train the model (including feature selection)
- 5. Validate the model

Objective: collaboratively implement this workflow and apply it to different ML datasets

Datasets

The objective of the project is to apply a Machine Learning model for Binary classification onto two different datasets:

- ► Banknote Authentication Dataset: https://archive.ics. uci.edu/ml/datasets/banknote+authentication
- Chronic Kidney Disease: https://www.kaggle.com/mansoordaku/ckdisease

Constitute groups of 3 to 4 students.

What I expect

- Create a git repository for your group: https://gitlab.imt-atlantique.fr/
- ► Test a few different models for classification, and implement feature selection
- Write the Python functions implementing the workflow in one single .py file.
- Write at least one unit test for one of the functions
- Apply the same workflow onto the two datasets, called from a Jupyter Notebook
- ► Think about good programming practices

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Deliverable

Code: **one** .py file for the functions and **one** .ipynb file to run the code included in the Git repository + all what is needed to run the code (datasets, another .py file for unit tests if needed, etc.)

One Git repository per group should be "sent" by e-mail to elsa.dupraz@imt-atlantique.fr before the 20th of November, 11PM.

 \rightarrow please give me the sufficient level of rights to see the repository (e.g. developer)

The .py file must contain...

- ▶ One function for pre-processing
- One function to prepare the dataset for training
- ➤ One function for training (typically applies up to 5 different methods for binary classification)
- One function to display all the results in a convenient form for comparison
- As many sub-functions as needed

The .ipynb file must...

- ▶ Apply the functions **of the .py file** onto the two datasets
- Show and compare the results for each dataset (e.g., numbers, curves, etc.)
- Comment on the results in both cases (which method would you choose at the end?)
- Discuss good programming practices and how they were taken into account into the project

Final comments

- Advice for good programming practices: https://mikecroucher.github.io/reproducible_ML/
- Register your groups before the 14th of October, at https://semestriel.framapad.org/p/ bpvmOuOniv-aagc?lang=fr