Using command line: tester.py file_name n_fold {-d} to get the result of dataset.

And pleas use the readArff.py in my file, since I made some changes to the original one.

- 1. file_name: The arff file that you want to test.
- 2. n_{fold} : The number of chunks that you want to split the data into. For example, if you want to do 5-fold cross validations, then $n_{fold} = 5$. This provide a way to do more cross validation rather than 5-fold.
- 3. -d: It's an optional parameter. If you provide -d, then you will get every result and average result, otherwise you will only get the average result.