**Bioinformatics – Practice 0: Classifications**

You have a dataset with protein descriptors for proteins involved in two types of cancers: HCC and HBC. Using Practise/CancerDataSet.csv database and Weka, find the best classification model that can predict if a protein is related with HCC or HBC. You should use 10-fold cross validation, at least one method from *BayesNet*, *Decision Table*, and *Random Forest*. Order the performance of the models using TP Rate.