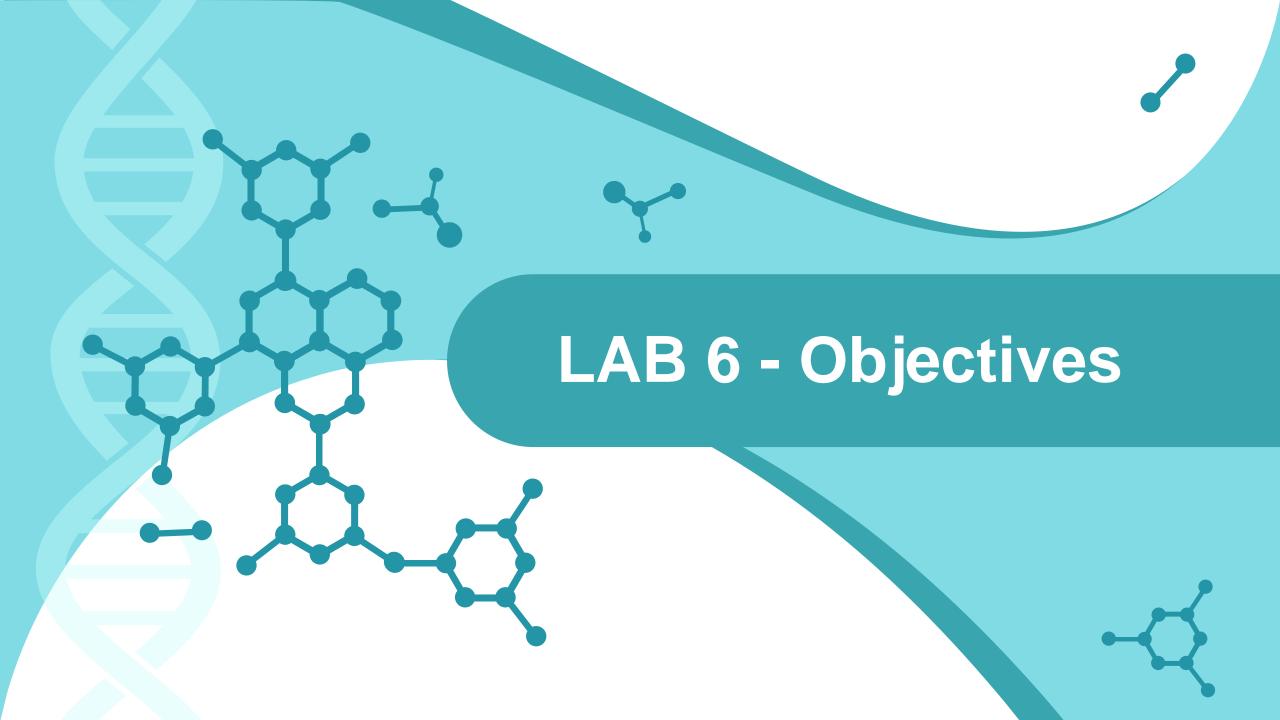


# Bioinformatics LAB 6 ML basics



Prof.ssa Elisa Ficarra
Prof.ssa Santa Di Cataldo
Eng. Marta Lovino
Eng. Alessio Mascolini

Politecnico di Torino DAUIN Dept. of Control and Computer Engineering

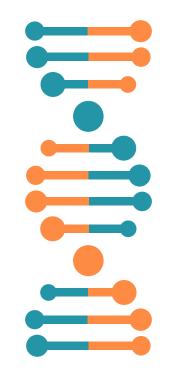


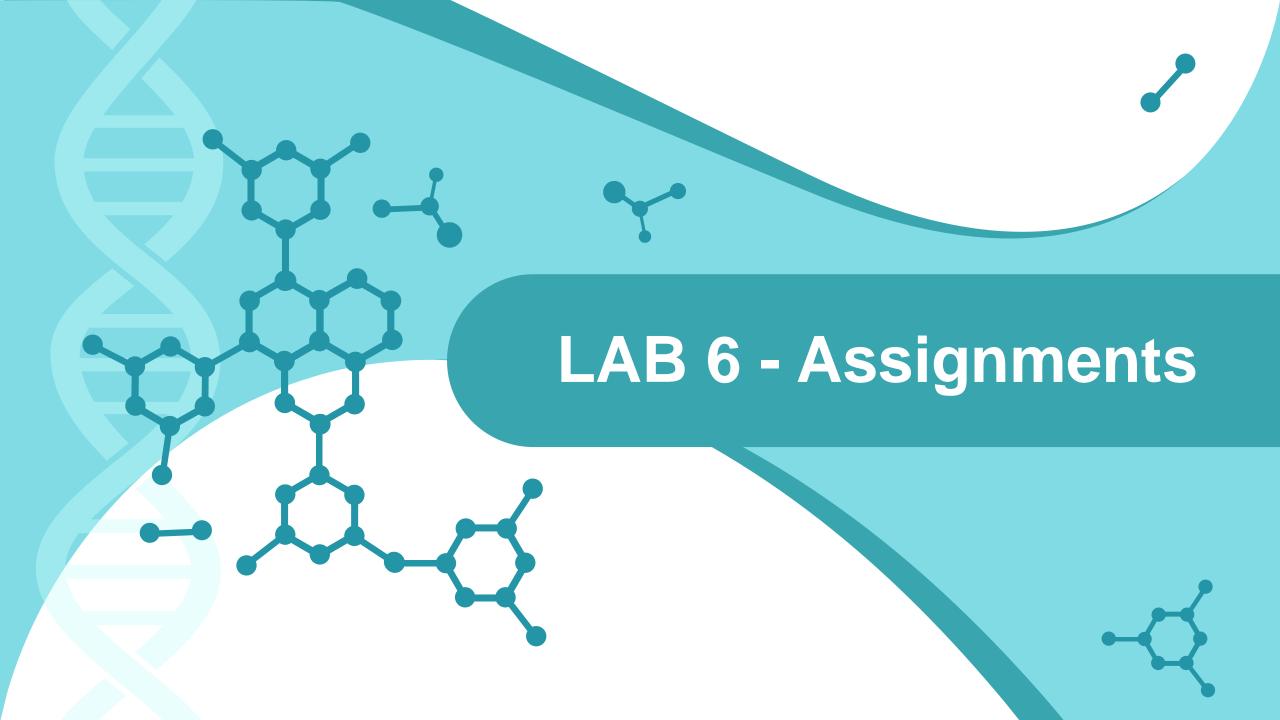
## Objectives

- Build gene fusion sequence from breakpoints
- Bayesian Convolutional Neural Networks on gene fusions sequences



# Gene fusion sequences



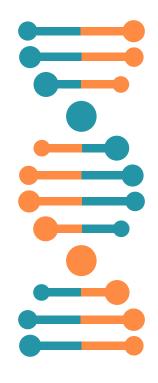


### Assignment 1: Implement a B-CNN

Download training\_set.csv and test\_set\_1.csv from the Teaching Portal and adapt the steps provided you during the previous lesson in order to implement a 1D Bayesian Convolutional Neural Network.

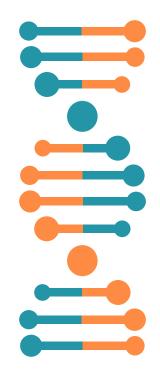
The network should distinguish between oncogenic and not oncogenic gene fusions exploiting the gene fusion sequence.

In details, analyze and explore the certainty classification value for the samples which are correctly classified.



### LAB6 – Take home message

 The oncogenic property is complex: most likely you need additional information to succeed in the classification task.





Questions?

# Remember: no question is stupid