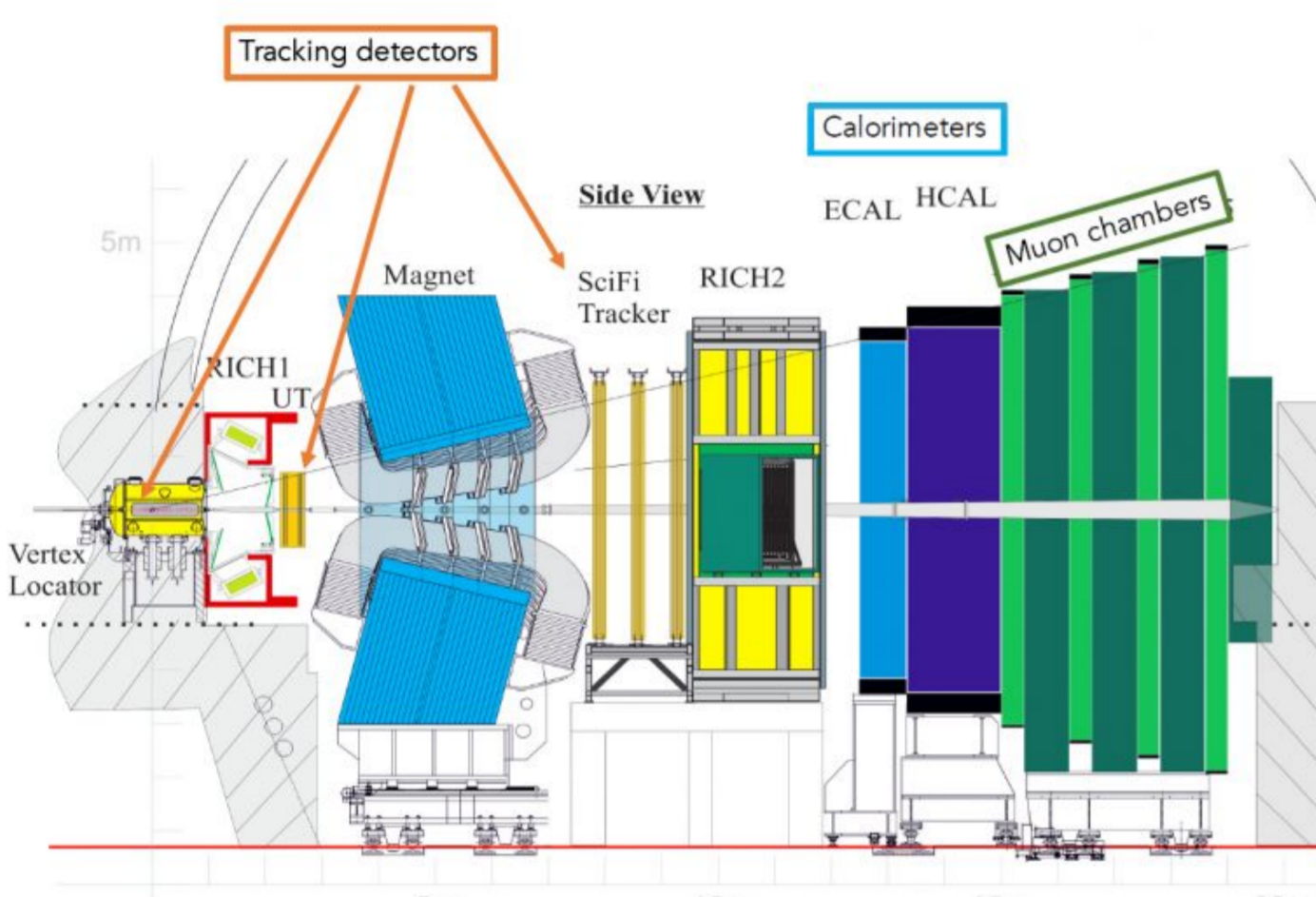


LHCb experiment

- LHCb experiment (b stands for beauty) at CERN specializes in investigating the slight **differences between matter and antimatter** by studying the **beauty quark**
- Whenever measurements show a deviation with respect to the Standard Model (SM) it is crucial to understand if it is due to **systematic uncertainties** or **new physics**:

- Systematic uncertainties:
 - Statistical fluctuation
 - Underestimated theory uncertainties
 - Detector effects

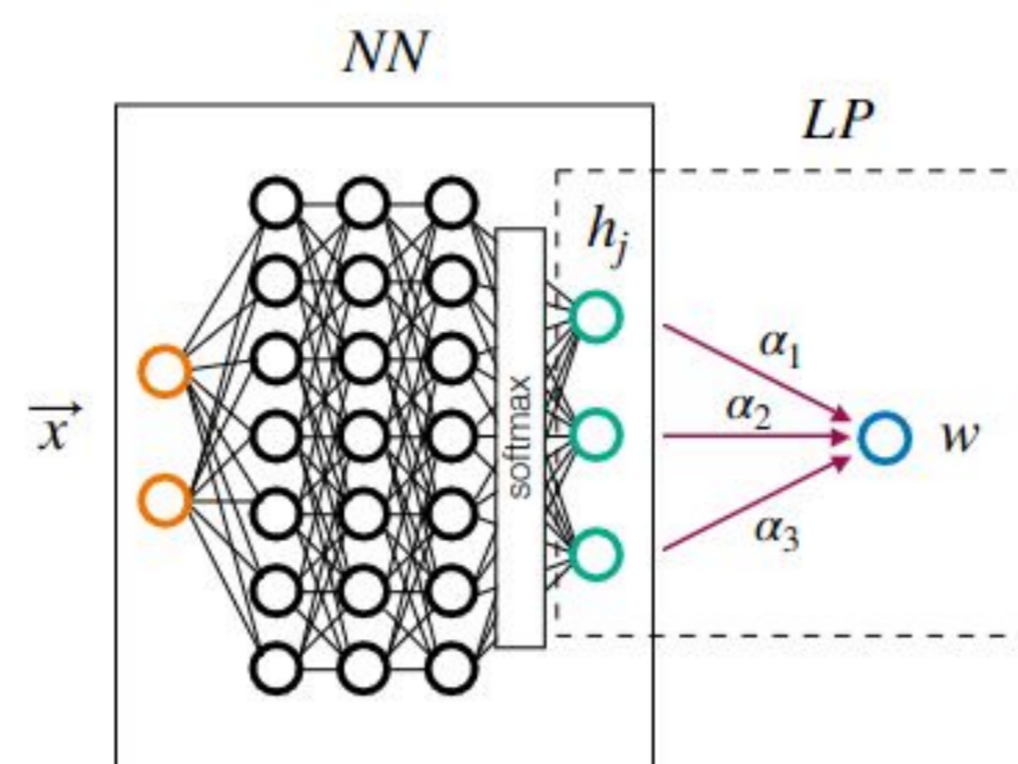
A fifth fundamental interaction?



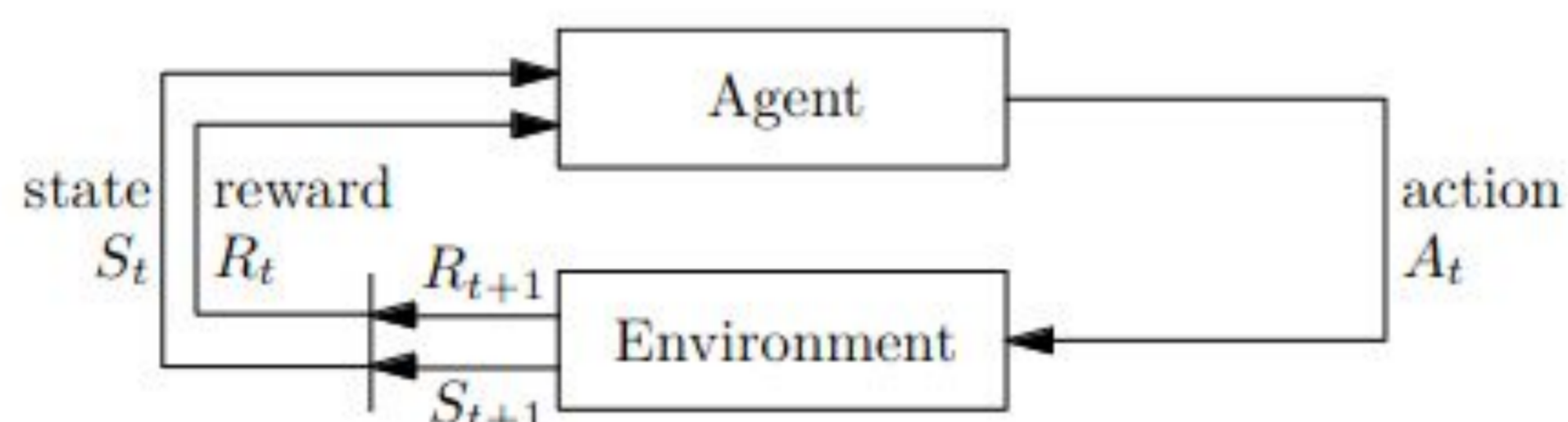
Previous works

Previous works have been made to play the devil's advocate (**DL Advocate project**) using Deep Learning techniques:

- One technique was built by combining a **Neural Network (NN)** with a **Linear Programming (LP)** solver:



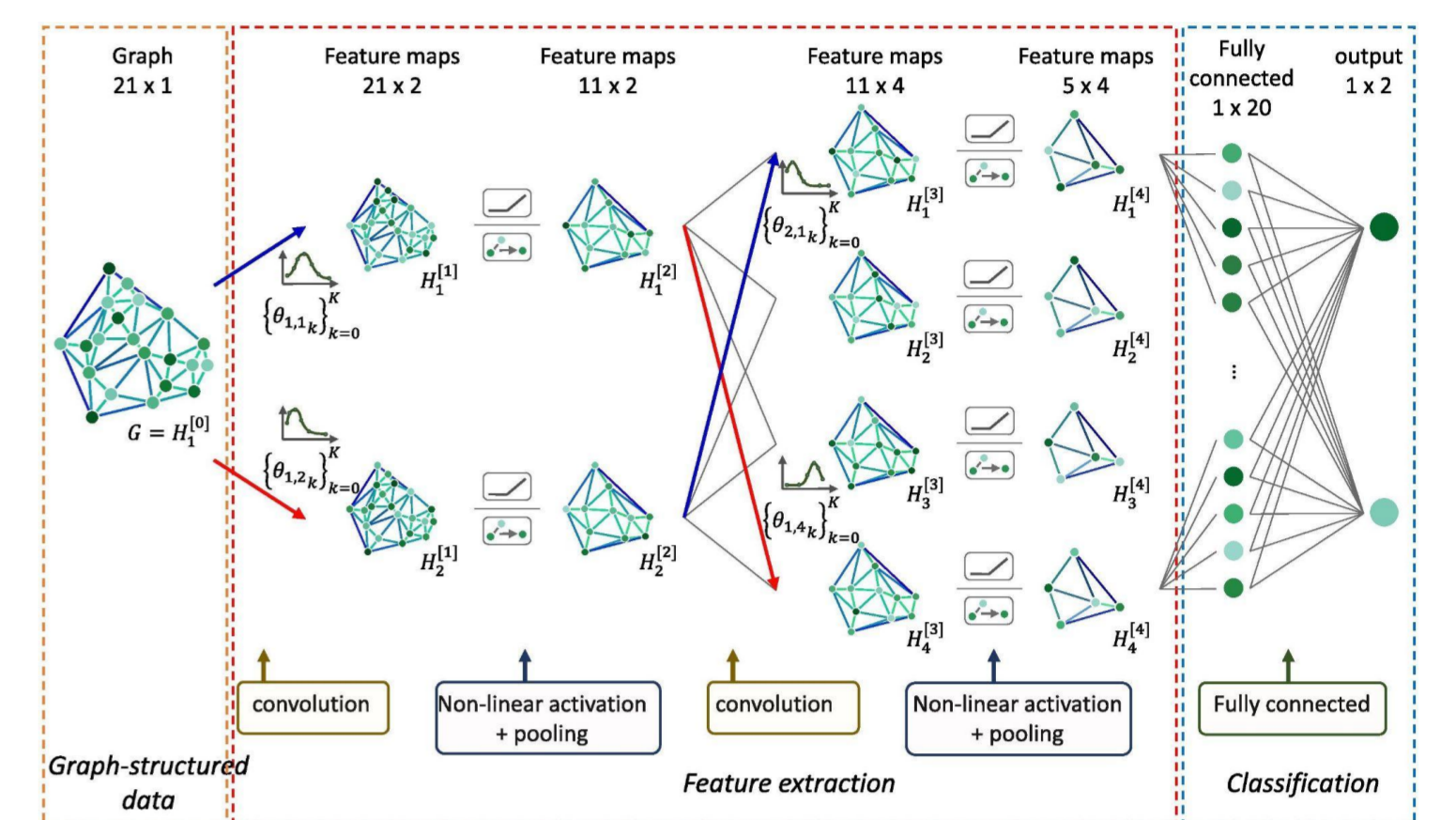
- Another technique was based on a **Reinforcement Learning (RL)** algorithm:



Future works

Future DL Advocate projects will involve:

- **Branching Ratio (BR)** predictions for decays not present in the Particle Data Group (PDG)
- Uncovering of **hidden backgrounds**
- Predictions of relevant **distributions of decays**



Taken from: ISPRS J. Photogramm. Remote Sens. **150**, 259-273 (2019).

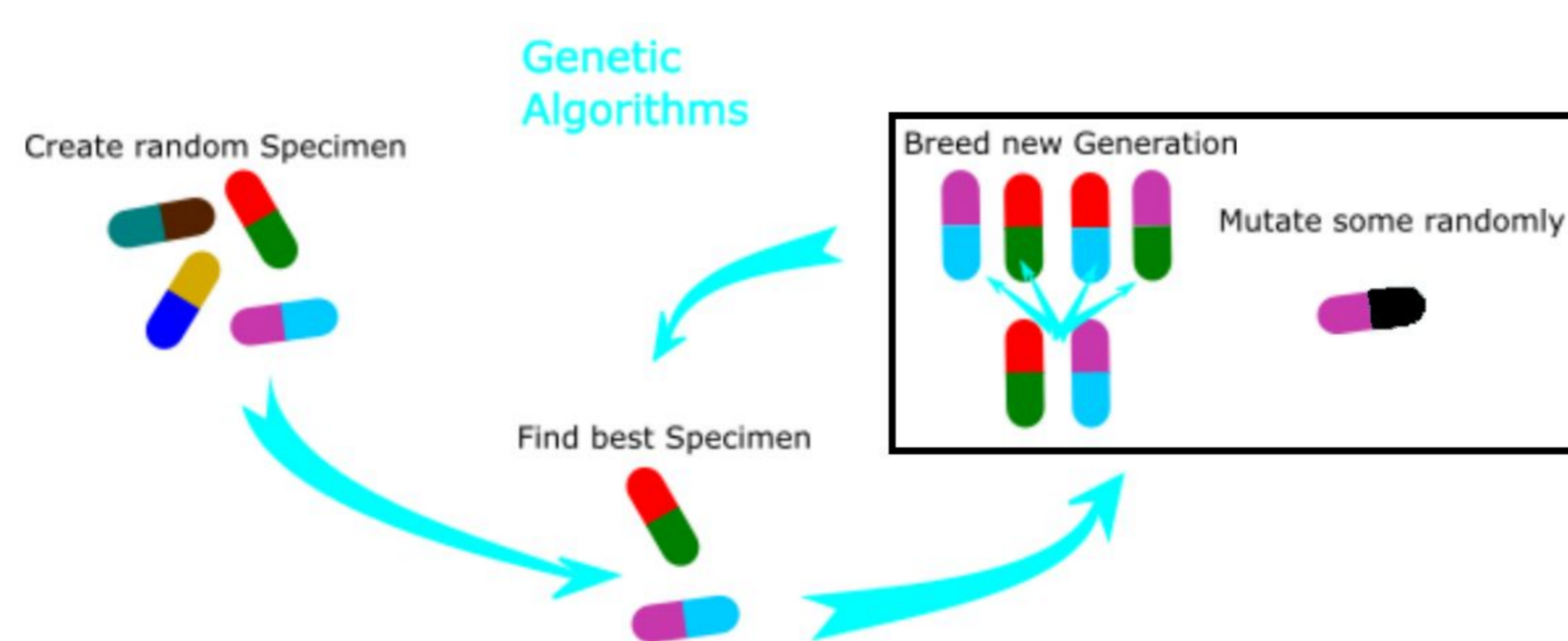
Playing the devil's advocate through deep learning: systematic uncertainties or new physics?

Uncovering hidden backgrounds: 1st approach

- **RL algorithm** where the **state** describes the **particles** in the decay
- The **agent** is described by a **NN** whose input is the state and whose output is an action that will define the next state
 - The NN of the agent is a **Graph Convolutional Network (GCN)**, as this enforces permutation invariance
 - To avoid having a huge number of states, **actions** should be defined to always **transition among states with physical sense** (charge conservation...)
- The **reward** defines how dangerous the background (the final state) is for the signal

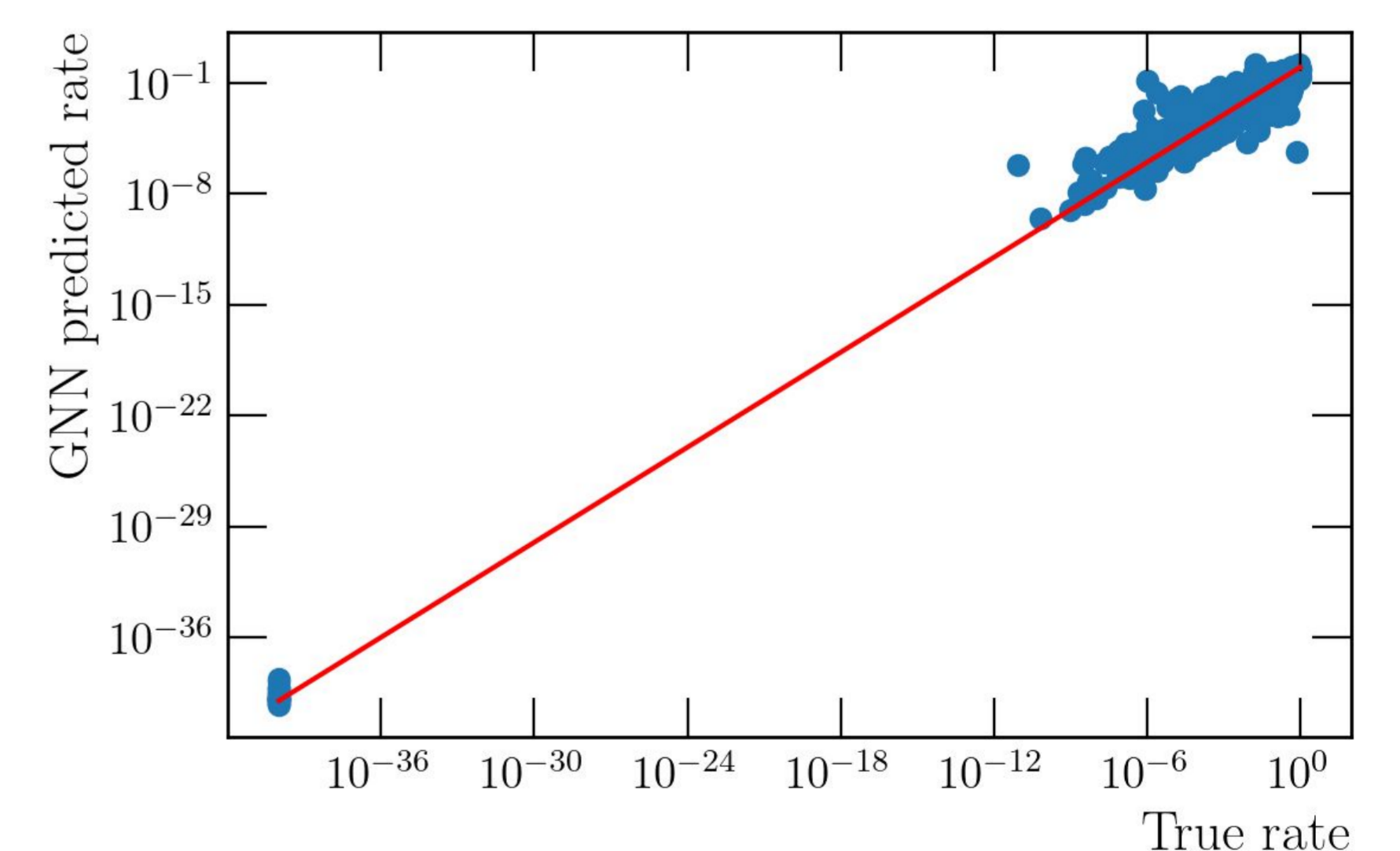
Uncovering hidden backgrounds: 2nd approach

- Use of **Genetic Algorithms (GAs)** to find the most **dangerous backgrounds**
- Each **individual** represents a possible background. The **genes** store the information of the involved particles, the possible intermediate resonances...
- The **fitness function** defines how dangerous each background is



Branching Ratio predictions

- Use of a **GCN** to **predict the BR** of decays not present in the Particle Data Group
- **10-folding** technique to maximize data utilization and obtain a better performance estimation
- PDG API as the source of our data



References:

- [1] <https://hal.science/hal-03777958/document>
- [2] <https://arxiv.org/pdf/2303.15956.pdf>

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