

Profile Likelihood Analysis

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Done

- 1. Installed: Blueice, pax and laidbax(Xenon1T likelihood fitting code)
- 2. Basic understanding of hypothesis test

Problems

- For pax, I followed the steps in Readme to test if the installation is working, it failed:
`pkg_resources.DistributionNotFound: The 'python-snappy'`
distribution was not found and is required by pax

Hypothesis Testing

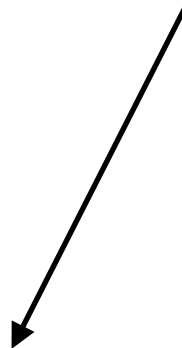
- The basic idea is to accept or reject a hypothesis by formulating two opposing hypotheses: the null hypothesis and alternative hypothesis.
- For discovering a new signal, define two hypothesis: H_0 (only known process/background) and H_1 (background+signal).
- To accept or reject H_0/H_1 , calculate the p-value, the probability under that assumption.
- In particle physics, p-value $\rightarrow Z$ (significance)

H0 (B)

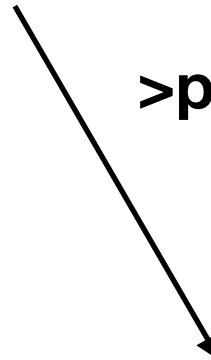


P-value

< p*



> p*



“Discovery”

Nothing special

To Do

- 1. Profile Likelihood Ratio
- 2. Look through the codes and play with them?