

A software tool for your analysis

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Overview

Introduction	The Use Case of Sprout	The Sprout Family	
How Sprout came about	How Sprout can help you in your analysis	Overview of Sprout's features	



Introduction

- Sprout **Y**: a little C++ project that grew from ROOT
- My background: Hyperon data analysis for Panda@HADES
- Data from proton-proton beam-time at HADES collected in 2022
- **Goal:** Measure *cross sections* for various Sigma0 production channels
- Current focus: Careful studies of Σ⁰ reconstruction efficiencies for different channels







Introduction

The data we're looking for:

Reconstruct final-state particle four-vectors based on detector response to learn something about short-lived particles







The Analysis Workflow





Preselection

Look for presence of final-state particles that could originate from a Σ^0 event

- Assess the data quality:
 - Various qualifiers for detector track quality
 - Use of different triggers
- Determine particle type from detector information
- Check for presence of relevant final-state particles





Preselection

Difficulties:

- Many different parameters and combinations to test and keep track of
- Often changing code to alter parameters
- Easy to lose track and prone to human error





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SproutParam

Offers a simple way to store and access parameters throughout your analysis





Assess the likelihood that the final state particles actually originated from a Σ^0 event

- Need to filter true Σ^0 events from *background*
 - Falsely identified particles
 - Particles from secondary reactions
 - Combinatorics
- Use *cuts* to decrease background







Assess the likelihood that the final state particles actually originated from a Σ^0 event

- Identify suitable cut variables to separate signal from background
- Use simulations to define a suitable cut value,
 e.g. by optimizing the signal yield
- Must ensure that simulations accurately describe the data in each cut variable







Difficulties:

- An analysis might contain *many* different cuts
 - Many variables to check for agreement between data and simulations
 - Each cut value tuned by trying out ~ 100 positions to optimize yield
 - <u>A lot</u> of quality assurance plots to generate and compare





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Event

Reconstruction

Preselection

- Even after cuts, some background usually remains
- Easily seen from invariant mass histograms
- Commonly use fits to subtract background contributions from the signal yield





Difficulties:

- Each fit needs visual quality assurance. Does it fit the data accurately?
- Especially background distributions may differ over different kinematic regions, no one-model fits all
- Each fit may require different start parameters to not fail
- Some analyses may require ~ 100 fits





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Inspired by Rafał Lalik's (Jagiellonian University) software package <u>HelloFitty</u> 🂭

Perform a measurement on yield to extract physics of interest

Caveat: selection cuts may impact the measurement.

Potential source of systematic error!



- Must accurately determine, minimize and evaluate sources of systematic errors
- Common (bad) practice: alter cut values and quote systematic uncertainty based on cumulation of observed variations in the relevant measurement
- May result in:
 - Penalised diligence: Ο
 - many checks = larger systematics
 - Larger systematics = overestimated agreement, Ο fewer citations





Better way: Determine *significant* sources of systematic errors with a *consistency check*

- Check passed \checkmark : do not quote as systematic uncertainty
- Check failed X: remove source of the error. If impossible, evaluate and quote systematic uncertainty
- R. Barlow suggests a check based on the uncorrelated uncertainty between data sets (e.g. two different cut values) Extract
- Measurement Data sets containing the same events are highly statistically correlated Event set a Reconstruction set b Preselection See Roger Barlow, 2002, Systematic Errors: Facts and Fictions 18

Difficulties:

- Many cuts throughout analysis for which a consistency check must be made
- Important to visually inspect how a measurement is impacted by each cut value
 - Look for trends a critical region might be present even if check is passed





SITET Figure from: *Viktor Thorén, Hadron Physics in a Polarized World, 2022*

Difficulties:

- Many cuts throughout analysis for which a consistency check must be made
- Important to visually inspect how a measurement is impacted by each cut value
 - Look for trends and critical regions

SproutCut - features to come: Help you perform consistency checks and generate quality assurance plots to detect systematic errors





See Roger Barlow, 2002, <u>Systematic Errors: Facts and Fictions</u>

The Sprout 🌱 Family

- Data analysis relies heavily on ROOT
- ROOT is great, but:
 - Takes up many lines of code
 - Find myself writing a lot of similar code many times over
- General aims of Sprout **?**:
 - Automate the boilerplate
 - Make it easier to keep manageable and modular





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The Use Case of Spro

SproutPlot

An analysis involves hundreds of histograms

- SproutPlot keeps a collection of all of them
- Automatically style, draw and write all histograms at once...
- ... but keep the ability to modify individual histograms as usual when needed
- Contains many more features you might find useful!





SproutPlot

SproutPlot myPlots; // create and fill histogr

```
TFile myFile("myFile.root", "recreate");
myFile.cd()
myPlots.writeHist();
```

// create and fill histograms

TFile myFile("myFile.root", "recreate");
myFile.cd();
h1.Write();
h2.Write();
h3.Write();
h4.Write();
h5.Write();
h6.Write();
h6.Write();
h8.Write();
h9.Write();
h10.Write();



SproutFit

- Fits user-specified signal+background distributions to histograms contained in a SproutPlot.
- Signal and background models specified by the user in a .txt file
- Automatically finds suitable start parameters
- Easily modify parameters of bad fits

pol2 gaus -3.11928 4.99871 -2 2 28.7394 7.78128 pol2 gaus -3.11928 4.99871 -2 2 41.237 9,95509 pol2 gaus -3.11928 4.99871 -2 2 19.3014 7.24632 pol2 gaus -3.11928 4.99871 -2 2 29.7916 8.59777 pol2 gaus -3.11928 4.99871 -2 2 36.1238 8.90835 pol2 gaus -3.11928 4.99871 -2 2 72.5726 14.8853 pol2 gaus -3.11928 4.99871 -2 2 37.4449 9.37459 pol2 gaus -3.11928 4.99871 -2 2 34.4013 9.57495 pol2 gaus -3.11928 4.99871 -2 2 75.3767 14.8835 pol2 gaus -3.11928 4.99871 -2 2 22.0282 7.48923 pol2 gaus -3.11928 4.99871 -2 2 23.6093 7.68782 pol2 gaus -3.11928 4.99871 -2 2 24.4318 8.17412 pol2 gaus -3.11928 4.99871 -2 2 10.4908 6.32788 pol2 gaus -3.11928 4.99871 -2 2 29.1846 7.97549 pol2 gaus -3.11928 4.99871 -2 2 37.3783 9.18131 pol2 gaus -3.11928 4.99871 -2 2 22.4391 7,93826 pol2 gaus -3.11928 4.99871 -2 2 78.2404 14.4209 pol2 gaus -3.11928 4.99871 -2 2 24.6451 7.74814 pol2 gaus -3.11928 4.99871 -2 2 36.5364 10.5195 pol2 gaus -3.11928 4.99871 -2 2 34.7326 8.65592 pol2 gaus -3.11928 4.99871 -2 2 33.2054 9.41413 pol2 gaus -3.11928 4.99871 -2 2 32.8913 8.84999 pol2 gaus -3.11928 4.99871 -2 2 28.615 8.67445 pol2 gaus -3.11928 4.99871 -2 2 11.157 6.63691 pol2 gaus -3.11928 4.99871 -2 2 22.894 8.08729



SproutFit

Perform many fits and produce quality assurance plots with ease:

SproutPlot myPlots;
// Add and fill histograms to myPlots
SproutFit fitter;
fitter.fit(myPlots,"myFits.png");





Access Sprout

Who can use Sprout?



- You use C++/ROOT as part of your analysis
- Sprout can be used from both executables and ROOT-macros

Download the package and access guides for set-up and usage at:

https://github.com/malle-b/Sprout









Backup



SproutParam

- Store parameters by name in .txt file and easily retrieve them from your analysis code
- Retrieval algorithm is O(1) on average
- Stored parameters can be changed easily without needing to recompile
- Easily define and run different analysis cases with different sets of parameters for comparison





SproutCut

- Prepare cuts by specifying name (for later access) and cut value
- SproutCut keeps a collection of all created cuts
- Upon applying a cut, automatically generates histogram for quality assurance
- Easily write all generated histograms at once
- See full documentation on GitHub for more and future features



