

ROOT Tutorial



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**Practical session for bachelor project
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Contents

- **1st part**

- Create a TGraph
- Create a TH1D
- Write simple code
- Pointers
- Writing a TTree
- Reading a TTree

- **2nd part**

- Read simulation files
- Write a function
- Fitting of data

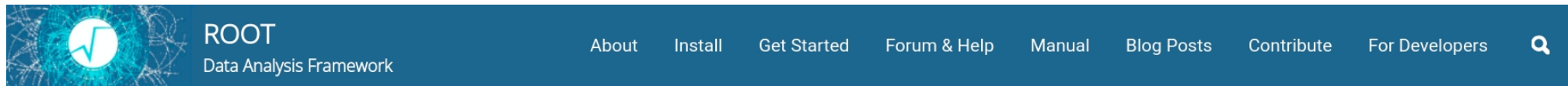
Nomenclature:

- Blue: you type it

- Red: indicate in what environment you are

Macros and slides are here https://github.com/andresgd17/Bachelor_project.git

Google: ROOT CERN

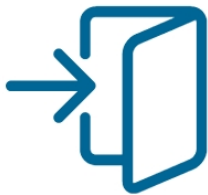


ROOT: analyzing petabytes of data, scientifically.

An open-source data analysis framework used by high energy physics and others.

 [Learn more](#)

 [Install v6.26/02](#)



[Start](#)



[Reference](#)



[Forum](#)



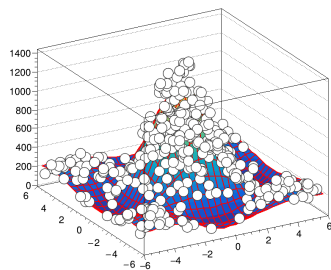
[Gallery](#)

ROOT in a nutshell

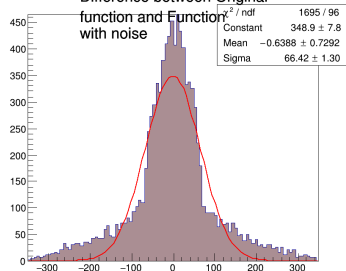
- **ROOT is a large Object-Oriented data handling and analysis framework**
 - C++ interpreter
 - Efficient object store scaling from kB's to PB's
- **Extensive set of multi-dimensional histogramming, data fitting, modeling and analysis methods.**

ROOT: Graphics

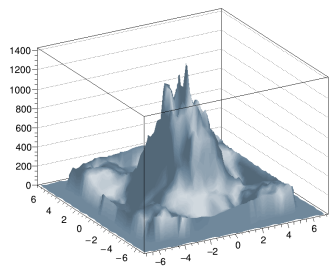
Original function with Graph2D points on top



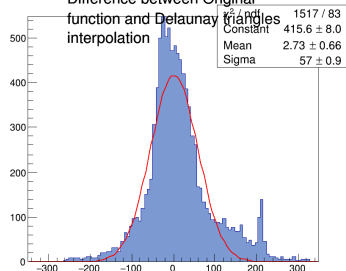
Difference between Original function and Function with noise



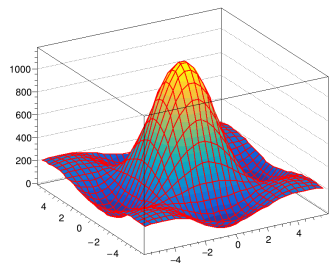
Histogram produced with Delaunay interpolation



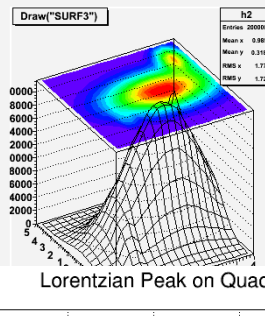
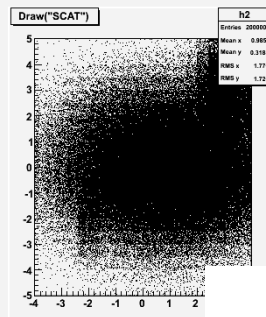
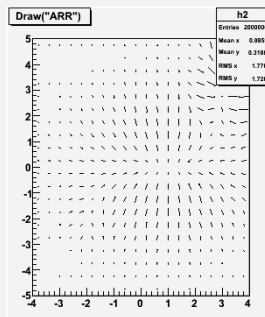
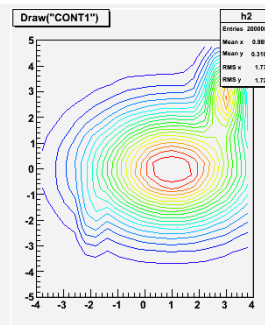
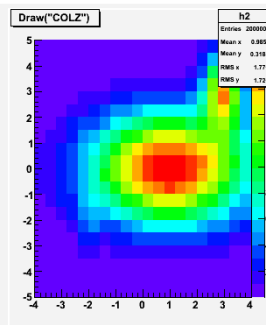
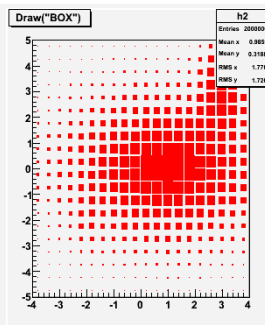
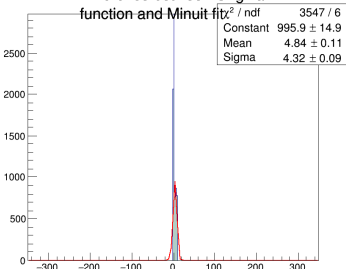
Difference between Original function and Delaunay triangles interpolation



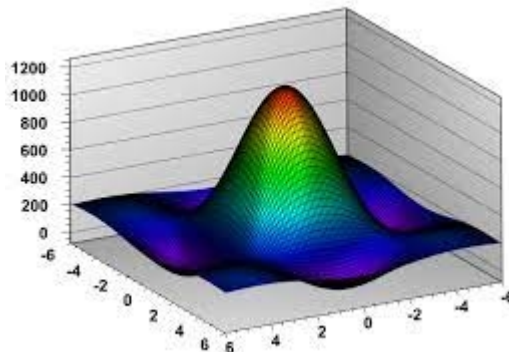
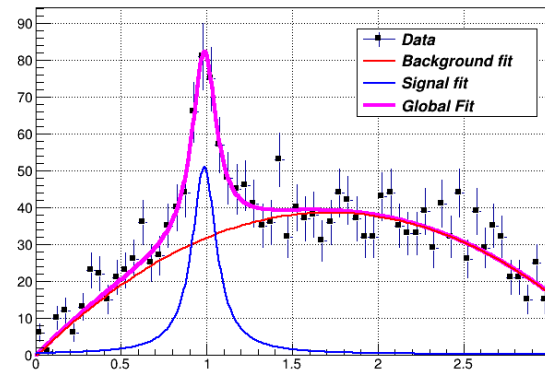
Minuit fit result on the Graph2D points



Difference between Original function and Minuit fit



Lorentzian Peak on Quadratic Background



ROOT in a nutshell

- **The user can interact with ROOT using:**
 - graphical user interface, (probably don't work in rug cluster)
 - the command line, or
 - scripts.
- **The command and scripting language is C++**

ROOT in the RUG cluster

- **Connect to cluster:**
 - ssh <snumber>@peregrine.hpc.rug.nl
 - https://wiki.hpc.rug.nl/_media/peregrine/additional_information/peregrine_course.pdf
- **Load ROOT module in the cluster**
 - \$ module spider ROOT
 - \$ module load ROOT

ROOT prompt

- **Starting ROOT**

- `$ root`

- **In the ROOT prompt you can type:**

- `root [] 5+4` `root [] int i = 3`

- `root [] TMath::Pi()` `root [] cout << i << endl;`

- `root [] float x[5] = {1, 2, 3, 4, 5}`

- `root [] float y[] = {1, 4, 9, 16, 25}`

TGraph

- **Creating a Graph with TGraph in the ROOT prompt:**
 - `root [] TGraph gr(5,x,y)`
 - `root [] gr.Draw("alp")`
 - `root [] TGraph gr2`
 - `root [] gr2.SetPoint(0,1,1)`
 - `root [] gr2.SetPoint(1,2,4)`
 - `root [] gr2.Draw("alp")`
 -

TH1D

- **Creating a histogram of 1 dimension:**
 - `root [] TH1D histo("name", "title", 10, 0, 100)`
 - `root [] histo.Fill(2)`
 - `root [] histo.Draw()`
 - `root [] histo.Fill(50, 4)`
 - `root [] histo.Draw("histo")`
 -

Creating a macro.C

- **macro.C** is the script contains your code
- **To execute your script macro.C:**
 - System prompt: `$ root macro.C`
 - ROOT prompt : `root [] .x macro.C`
- **Useful options (root --help or root.help):**
 - System prompt: `$ root -b -q macro.C`
 - ROOT prompt : `root [] .L macro.C`

Pointers

- A pointer is a variable that stores address of another variable, .i.e., point to the address of the memory location.
- A pointer declaration use `*` and has the form: `data_type * name;`
 - Variable declaration: `int i = 8;`
 - Pointer declaration: `int * i = new int(8);` (To print the value type: `*i`)
- Allocate memory with operator `new` just for pointers:
- We access to members of pointers using: `->`
- You can get the address of variable using: `&`

Pointers

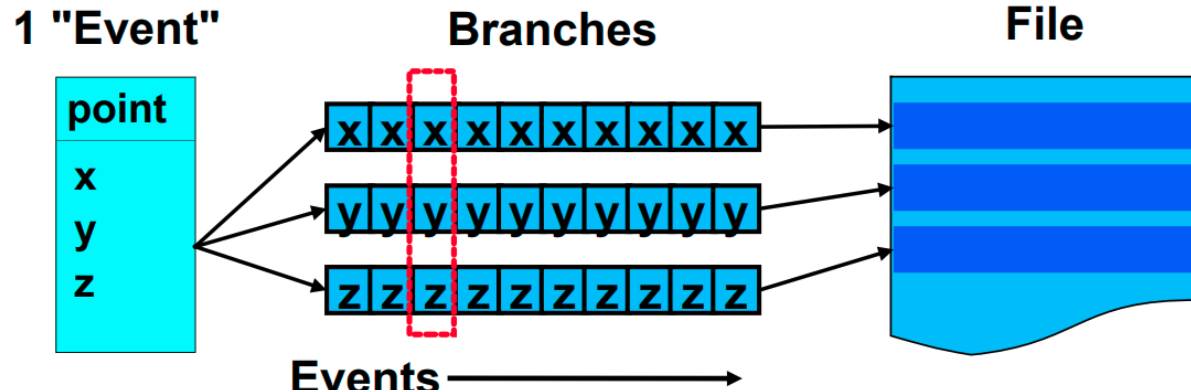
- **Create macro_pointers.C by modifying:**
 - TGraph gr(5,x,y) → TGraph *gr = new TGraph(5,x,y)
 - gr.Draw("alp") → gr->Draw("alp")

Break

- **Everything is ok?**
- **Any questions?**

TTree

- Trees have been designed to support very large collections of objects.
- Trees allow direct and random access to any entry
- Trees can be browsed via TBrowser
- The class TTree is the main container for data storage
 - It can store any class and basic types (e.g. Float_t)
 - When reading a tree, certain branches can be switched off → speed up of analysis when not all data is needed



Ttree - writing

- **Root files have extension <filename>.root**
 - `root [] TFile *file = new TFile("myRootFile.root", "recreate")`
 - `root [] TTree *tree = new TTree("tree_name", "tree_title")`
 - `root [] float energy`
 - `root [] tree->Branch("Energy", &energy)`
 - `root [] tree->Print()`
 - `root [] energy = 50`
 - `root [] tree->Fill()`
 - ...

Ttree - writing

- ...
- root [] tree->Print()
- root [] tree->Scan("Energy")
- root [] energy = 100
- root [] tree->Fill()
- root [] tree->Print()
- root [] tree->Scan("Energy")
- root [] file->Write()
- root [] file->Close()



Don't forget to
write all objects
in the file!!

Ttree - reading

- **Reading myRootFile.root:**
 - `root [] TFile *file = new TFile("myRootFile.root", "read")`
 - `root [] .ls`
 - `root [] tree_name->Print()`
 - `root [] tree_name->Scan("Energy")`
 - `root [] tree_name → Draw("Energy")`
- **Check readtree.C:** Script to read a tree
 - `$ root readtree.C`

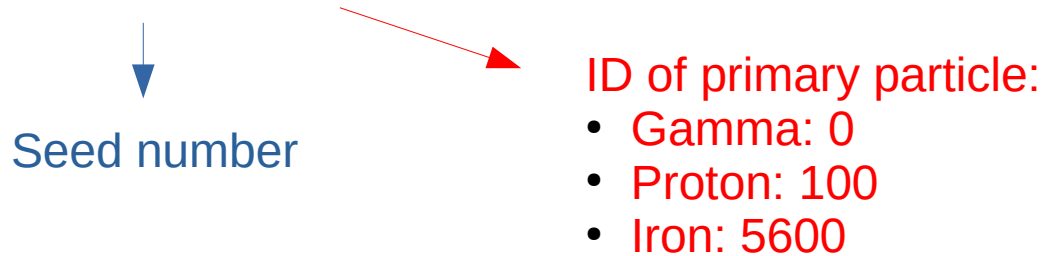
Small assignment

- **Try to type by yourself all the codes again to familiarize with them.**
- **Create a script called writetree.C to create a Ttree:**
 - You can use the same code from Ttree – writing section :)
 - Try to create another branches: i.e. velocity, momentum, etc, and fill them with random numbers :)

2nd part of tutorial

- **Read simulations files:**

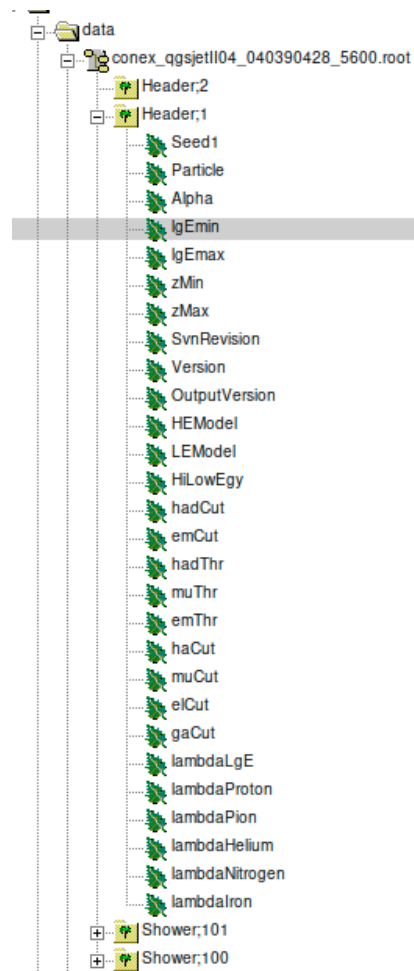
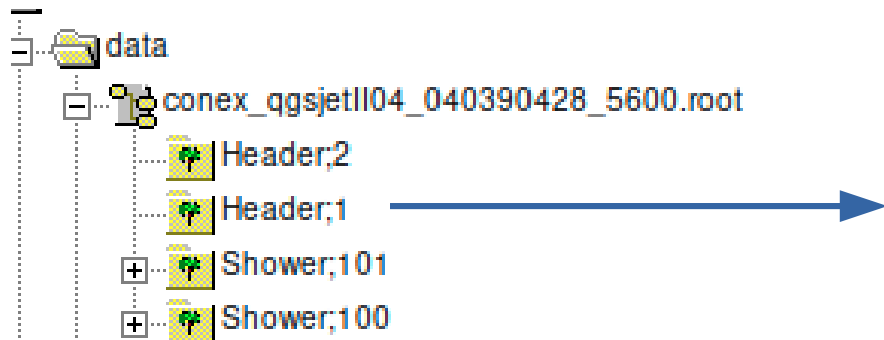
- conex_qgsjetll04_040390428_5600.root



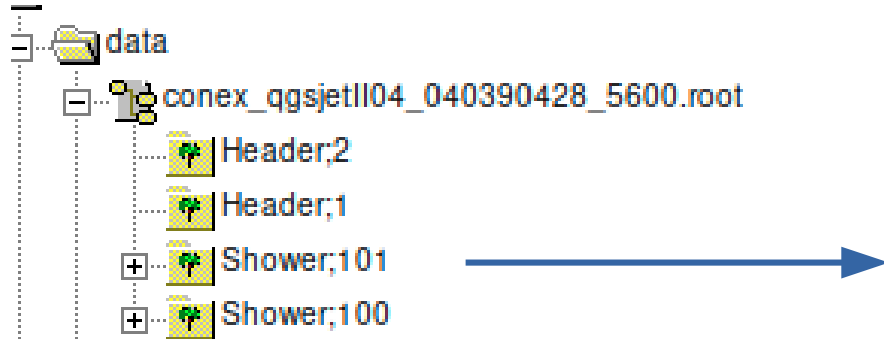
- 12 files in total:

- 4 files for each kind of primary particle: gamma, proton and iron
 - Each of the 4 files for each energy: 10 TeV, 50 TeV, 100 TeV, 300 TeV

Data files



Data files



Important information

- **Tree:** Shower;101
 - **Branches:**
 - **LgE:** $\log_{10}(\text{Energy / eV})$ e.g: $\lg E = 14 \rightarrow \text{Energy} = 10^{14} \text{ eV}$
 - **nX:** number of points of "X", "N",
 - **X:** slant depth array (g/cm^2)
 - **N:** array of number of charged particles
- **For more information check README file in data folder**

Plot profiles

- Use `plotProfile.C` script to plot TGraph of charged particles profiles with legend.
- Let's plot one profile:
 - ROOT prompt:
 - `$ root`
 - `root [] .x plotProfile.C("../data/conex_qgsjet1104_040390428_5600.root",4)`
 - System prompt:
 - `$ root "plotProfile.C(\"../data/conex_qgsjet1104_040390428_5600.root\",4)"`

Assignment

- Open the `conex_xxx.root` files and explore the contents.
- Plot the profile of charged particles, electrons, hadrons and muons (in the same TGraph) for each file: 12 plots in total
- Fix the Y-axis title in the profile plot (search how to do on root fórum, google, ...)

Write a function

- Check plotFitFunction.C
 - Define second-order polynomial function
 - Fit profile around the maximum
- Assignment:
 - Get the X_{max} from the second-order polynomial fitting
 - Plot the distribution of the X_{max} in a histogram TH1D

Backup

Tree

- Branches

- root[] Shower->Print()
- root[] Shower->Show(5)
- root[] Shower->Scan(X)
- root[] Shower->Scan(N)
- root[] Shower → Draw("N:X", "X < 500", "*", 1, 5)
-

Constraints on
data

First shower to
consider

of showers

Variables you
want to draw

Useful

- You can open files when starting ROOT by:
 - `$ root file.root`
- Suppose a macro `myscript.C` with a function `myscript(string file, int i, float x)`
 - `$ root "plot.C(\"filename.root\",1,1)"`
- Type `.*?` to see meta commands in the ROOT prompt

Need help?

- ROOT fórum (<https://root-forum.cern.ch/>)
- Youtube: search for root cern <something> (could work)
- Send me an e-mail! :)