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Master Thesis

Abstract

Master degree in Computer Engineering

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The AEgIS Experiment at the CERN aims to verify the weak interaction principle for antimatter. This document talks about "gAnWeb", a web application designed to simplify the analysis of physical data under the AEgIS experiment. This analysis can be performed using Root Data Analysis Framework by the Linux Terminal, but a graphical interface can ensure a better user experience, ease the user training and improve the productivity. A web application is a smart way to implement the interface because allows users to avoid installations, and centralizes all the eventual modifications. This document explains the choices made during the development of this application related to the goal of the user friendly data analysis, and shows the design process that led to the final product.

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Introduction

First of all it is important to understand at least generically what is the AEgIS experiment at the CERN and what are its goals. The acronym AEgIS stands for "Antimatter Experiment: gravity, Interferometry, Spectroscopy", this research aims to measure weak equivalence principle for antimatter. In the first part of this chapter some particulars are explained about this experiment, in the second part is introduced gAn Web, the main topic of this document, the application that allows the physicists to do data analysis in the AEgIS experiment environment easily, through a web interface.

1.1 The AEgIS experiment

AEgIS

FIGURE 1.1: AEgIS's Logo

The weak equivalence principle, also known as universality of free fall, states that in the same field all bodies fall with the same acceleration, regardless of the mass and the composition. This principle has been thoroughly tested for the matter, but not for the antimatter: the most important goal of AEgIS experiment is to measure the weak equivalence principle for the anti-matter; to test the universality of free fall, AEgIS measures gravitational interaction between matter (the earth) and anti-matter (anti-hydrogen). Anti-hydrogen is the antimatter counterpart of hydrogen, it is the simplest atom

built with antimatter (see below for a description of its components). The first anti-hydrogen in an electromagnetic trap has been produced by the ATHENA experiment in 2012 (http://www.nature.com/nature/journal/v419/n6906/full/419439a.html) with the contribution of a group of the University of Brescia.

The AEgIS experiment if the result of a wide and international scientific collaboration, as visible in the figure 1.2.



FIGURE 1.2: Aegis Collaboration institutes

In the context of neutral antimatter, the gravitational interaction is of high interest, because it can potentially revealing new forces that violate the weak equivalence principle. Thomas Phillips, from Duke University, says: "If antimatter fell down faster, it would mean the discovery of at least one new force, probably two. If it fell up, it would mean our understanding of general relativity is incorrect". In a practical point of view AEgIS tries to mesure the time of flight and the vertical displacement of anti-hydrogen, by a moire deflectometer: this process is quite complex, and it is easier explain it by the following two figures (1.3 and 1.4).

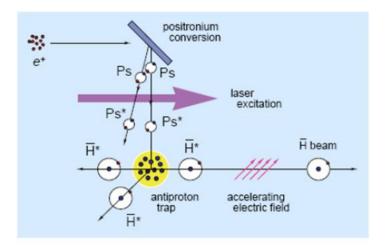


FIGURE 1.3: AEgIS's Scheme, taken from "AEgIS experiment at CERN: measuring antihydrogen free-fall in Earth's gravitational field to test WEP with antimatter" TODO INSERT-bibliographical-reference

In this first figure we can see the process that allows to create some anti-hydrogen. To correctly explain this process it is better start with some definitions:

- 1. Positron: it is the correspondent of the electron in the antimatter. It is an anti-electron, so an electron with positive electrical charge. It is indicated by "e+".
- 2. Positronium: it is an unstable system consisting of an electron and a positron, bound together into an exotic atom. It is indicated with *Ps*.
- 3. Antiporoton: it is the antiparticle of the proton. Antiprotons are stable, but they are typically short-lived since any collision with a proton will cause both particles to be annihilated in a burst of energy. It is indicated with \overline{p} (pronounced pbar).
- 4. Antihydrogen: it is the antimatter counterpart of hydrogen. Whereas the common hydrogen atom is composed of an electron and proton, the antihydrogen atom is made up of a positron and antiproton. It is indicated with \overline{H} (pronounced hbar).
- 5. Antiproton trap: a devices that uses an axial magnetic field to transversely confine charged particles, in this case antiprotons.

The process shown in the figure is the following: a beam of positrons (that comes from a 22 Na radioactive source) is accelerated and driven

to collide against a "positron-positronium converter" (that is a mesoporous silica film). This process creates positronium, that needs to be excited by lasers, to reach the Rydberg State. The positronium in Rydberg state is indicated by Ps*, it has a longer life than the unexcited positronium, and can be driven to fly into an antiproton trap.

Antiprotons are provided in this way: Protons collide with nuclei inside a metal cylinder called "target". About four proton-antiproton pairs are produced in every million collisions, and is possible to separate antiprotons from matter using magnetic fields. The following step is to guide antiprotons toward the AD (Antiproton Decelerator) where they are slowed down (it is easier work with slow antiprotons). To carry out AEgIS experiment antiproton must be trapped and conserved inside an antiproton trap, where magnetic fields force the charged antiparticles to spiral around the magnetic field lines, and electric fields confine them along the magnetic axis.

In the following step Ps* and \overline{p} can combine themselves to generate excited antihydrogen ($\overline{H}*$) and electrons. The antihydrogen beam is accelerated using an electric field towards a Moiré deflectometer, during the travel it decays to ground state.

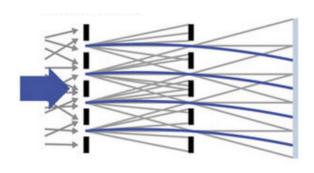


FIGURE 1.4: Moiré Deflectometer's Scheme, taken from "A http://www.nature.com/articles/ncomms5538"
TODO INSERT-bibliographical-reference

In the second figure is visible how does the Moirè deflectometer work. A antihydrogen beam is thrown toward two subsequent gratings that restrict the transmitted particles to well-defined trajectories. The trajectories are inflected by a force (in this case the force related to m * g) and follows a parabolic path. At the final part of the deflectometer there is a detector that shows where the antimatter annihilates, so is possible to compare the expected trajectories without forces with the obtained trajectories, and measure the force.

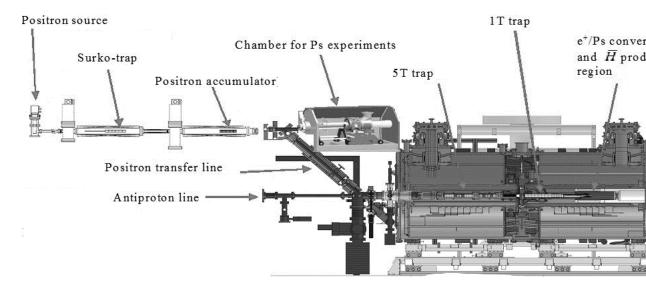


FIGURE 1.5: AEgIS apparatus set up, taken from "AEgIS experiment at CERN: measuring antihydrogen free-fall in Earth's gravitational field to test WEP with antimatter" TODO INSERT-bibliographical-reference

1.2 User friendly Data analysis: gAn Web

GAn Web is a web application, that creates a user friendly web interface, based on the most important human-machine interaction principles, between the users and a pre-existing data analysis application named gAn. This document's most important goal is explain in detail how gAn Web works, how and why it was created, what are the reasons for the choices made.

GAn Web is based on the pre-existent stand-alone program gAn, that allows users to do data analysis using a Linux terminal as interface. In turn, gAn is based on Root Data Analysis Framework, a vast and modular scientific software framework that provides all the functionalities needed to deal with big data processing, statistical analysis, visualisation and storage of physical data. GAn exploits and organizes the functionalities of Root, the resulting software is practical and achieves his goals, but a web interface can improve it in two ways:

1. gAn is a stand-alone program based on Root, installable on the user's machine; the user has to install the correct version of Root to avoid compatibility problems (Root is still not perfectly version independent: different versions can lead to different behaviours). Furthermore, this kind of program is continuously changing, the performed analysis is continuously improved, so the installed version of gAn is not final and unchangeable, and the user musts often update it. Instead, a centralized version installed on a server, with services accessible from a normal browser by the user can avoid (at least reduce) this kind of problems and be more usable.

2. a Linux terminal interface is practical for expert users, but a web based interface can be more attractive for new users, and, if well done, can be easier to use. It is important to notice that the users are physicists, not necessarily specialized in computer science, so, create a friendly and easily learnable interface can avoid them problems and time wasting.

The goal of gAn Web is to allow users to do analysis through a more friendly web interface, without install nothing on their machine. In the following figure there is a schema that shows how this program is organized.

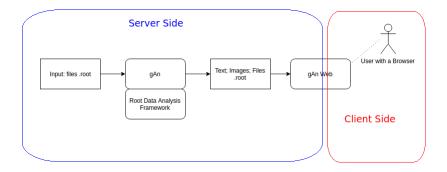


FIGURE 1.6: gAn - gAn Web simple scheme

The input of this system is represented by a set of files .root. These are raw, very big, binary files, incomprehensible to humans, generated by the hardware (mostly by detectors) of the AEgIS experiment, they need the Root Data Analysis Framework to be interpreted. These files contain a lot of information, too much and too disorganized to be helpful for the analysis. GAn can read these Root files, make an ordered and organized analysis, extract the most important information, and produce an output that consists of a text with the most important informations, comments, and eventual error logs, a folder

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with some figures, that can summarise effectively the most important points of the analysis, and some other .root files, with useful informations that allow gAn Web to make further processing. GAn Web can close the cycle acting as intermediary between the users and gAn: gAn Web can receive requests from the users, configure gAn to satisfy these requests, and deliver to the users exactly what they need.

Data Analysis

This chapter aims to explain first of all what is the "big data analysis", and why it is important in the modern world. Subsequently it will be exposed the exact use of the big data analysis technologies in AEgIS experiment, with reference to the used technologies and the choices made.

2.1 What is?

According to the John Tukey's definition data analysis is:

"Procedures for analyzing data, techniques for interpreting the results of such procedures, ways of planning the gathering of data to make its analysis easier, more precise or more accurate, and all the machinery and results of (mathematical) statistics which apply to analyzing data" (http://projecteuclid.org/euclid.aoms/1177704711).

The basic idea is that in the modern world almost each activity can provide a big amount of data, but only a few of them are really useful to gain interesting information. The data analysis is an structured process that allows to select the most important parts of this row data and exploit them to gain information able to answer questions, test hypotheses and approve or disprove theories. In the following image we can see the schema of this process.

Operational Environment Collection Processing Analysis and Exploitation Production

Relationship of Data, Information and Intelligence

FIGURE 2.1: Here is visible a basic schema of data processes and analysis

Data analysis can be divided in some steps:

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- 1. Data collection: data can be collected in a variety of ways. For example they may also be collected from sensors in the environment, such as satellites, recording devices, physical sensors etcetera. they may also be obtained through interviews, downloads from online sources, or reading documentation, so the analysis is feasible with a large variety of kinds of data.
- 2. Data processing: raw data must be well-organized for analysis: for example, placing data into row, columns, vector, etcetera.
- 3. Data cleaning: Once pre-processed and organized, the data may be incomplete, contain duplicates, or contain errors. Data cleaning is the process of correcting these errors, eliminating duplicates and handling incomplete data. Some ways to do this are record matching (confrontation between the records to find if there is something suspicious), validation of data (if there is the sureness that data values has to respect some limits), overall quality of existing data, de-duplication (process of removing of duplications). For particular kinds of input this process is very complex (for example vocal input needs an advanced spell-checker), for others is simple (for instance online-survey interviews made using closed choices)
- 4. Exploratory data analysis: in this step the data is analyzed. There are a variety of techniques referred to as exploratory data analysis to begin understanding the real content of the data.

This process may result in Descriptive statistics, such as the calculation of average or median, or in Data visualization, that allows to examine the data in graphical format, through graphics and other graphical objects.

- 5. Modeling and algorithms: another step is using mathematical models to find relations between different variables, such as causality or correlation. An example is the regression analysis.
- 6. Communication: this is the final step, and it is absolutely not trivial. Is important to find a way to report the obtained information to the user in an understandable format. The communication must be adapted to the different users, and to their requirements, to allow the data analysis to match them.

2.2 What a user can obtain from the data?

An user can benefit from the analysis of big data in various ways, in particular a working data analysis software can perform the subsequent tasks:

- 1. Retrieve Value: the system receives in input some cases (for instance case-A, case-B etcetera) and a set of variables (for instance variable-A, variable-B etcetera), and can shows in output the values of the variables in the set in the data cases (in case-A, case-B, variable-A = x, variable-B = y).
- 2. Filter: the system can show only the subset of the data that respect some conditions.
- 3. Find extremums, ranges and characterize distribution: the system can show the maximums o minimums values in the datasets, and in how large is the range in that the values are distributed. It is also possible to approximate with mathematical functions the statistical distributions of subsets of data.
- 4. Find Anomalies: check the dataset to find if there are some exceptional values, that are source of interest and need an explanation (errors in the data? unknown phenomena?).
- 5. Clustering: the analysis is extremely easier if is possible to group different subset of data with similar characteristics. For example: in a market research, obtaining a correct clustering on the customer allows to understand the different categories of customer with different goals and needs.

2.3 Relevance in the modern market

How and how much can the big data analysis be useful in the modern world? This analysis in recent years he has taken an extraordinary importance, The Economist says: "Big data has increased the demand of information management specialists in that Software AG, Oracle Corporation, IBM, Microsoft, SAP, EMC, HP and Dell have spent more than 15 billion dollars on software firms specializing in data management and analytics. In 2010, this industry was worth more than 100 billion dollars and was growing at almost 10 percent a year: about twice as fast as the software business as a whole." (taken from http://www.economist.com/node/15557443). This fact shows that companies (and governments) have realized the importance of the exploiting of the huge amount of data that the modern world produce. The hugeness of this amount of raw data is visible in the following data:

"There are 4.6 billion mobile-phone worldwide, and 1-2 billion people accessing the internet. In recent years, more than 1 billion people in the world entered the middle class, which means more people become more literate, which in turn leads to information growth. The capacity to exchange information has grown rapidly in recent years, to 667 exabytes annually in 2014." (data taken from: Wikipedia https://en.wikipedia.org/wiki/Big_data#Applications; The Economist http://www.economist.com/node/15557443; martinhilbert.net http://www.martinhilbert.net/WorldInfoCapacity.html/)

An advanced big data analysis offer good opportunities to improve decision-making in critical development, whether in a public or private institution, in numerous fields such as health care, employment, economic productivity, resource management etcetera.

Despite this kind of analysis provides benefits in a wide set of fields, in this document is specially important show the opportunities that data analysis can provide in a scientific environment, in particular in the AEgIS experiment. This is the main topic of the following paragraph.

2.4 Data Analysis at CERN

Some experiments at the CERN represents about 150 million sensors delivering data 40 million times per second. There are nearly 600 million collisions per second. The point is that not all these collisions are scientifically interesting. In the huge amount of collisions discussed previously there are only around 100 collisions of interest per second. This leads a big problem: it is difficult find a rational way to work with flows of data having this dimension, is absolutely necessary take only the part of the data actually required for the scientific analysis but it is also important don't waste nothing interesting in this dataset. Another problem is the speed with which physicists develop and change focus in their experimental work: they are not interested always at the same things, and they cannot predict in advance what they will need in the future, because their needs are related the process of experimentation, and can change continuously. So a system of analysis must be very flexible and dynamic, always ready to answer to new requirements, always able to find exactly what is the "interesting" part of data. This kind of problems are not just CERN's problems, they are common in major research centers. Bob Jones, Project Leader at CERN, says: "CERN is a leader but not alone in having to deal with such high data throughputs. We expect to see similar scales in other sciences (such as next generation genome sequencing as well as the Square Kilometre Array which will primarily be deployed in Australia and South Africa) and various business sectors linked to the growing Internet of Things in the near future." (quote taken from http://www.

cloudwatchhub.eu/what-big-data-really-looks-cern-universe-and-ever Big data analysis can provide an answer to this problems: with an intelligent and parameterizable process of filtering is possible extract only the subset of information scientifically interesting, and delivering them to the users in an organized and structured way, by tables, statistical values, graphical images. In this way is possible improve the productivity of the physicists, exempting them from unnecessary commitments and dynamically meeting their needs.

COMMENT: TODOTODOTODO ASCOLTATI QUESTO VIDEO E VEDI SE CI SONO BUONI SPUNTI https://ieondemand.com/presentations/big-data-analytics-for-improving-the-cern-s-large-hadron-collider-operations

2.5 Root - Physical Data Analysis

There are some software (often free software) specialized in the analysis of big data. Each these software has strengths and weaknesses, and none of them appear to be absolutely better than the others. Following there are some examples:

- 1. MATLAB (matrix laboratory) is a numerical computing environment. Is a proprietary (so, it is not available for free, and it is quite expensive) programming language developed by Math-Works. Matlab allows matrix manipulations, plotting of functions and data, implementation of algorithms, creation of user interfaces, and interfacing with programs written in other languages, including C, C++, Java, Fortran and Python. Some detractors say that the statistical support is incomplete if compared with other solutions (also free).
- 2. R is a programming language and a software environment for statistical computing. It is supported by the R Foundation for Statistical Computing. The R language is widely diffused for developing statistical software and for data analysis. His popularity has increased in recent years, this is due to the fact that R is free and allow to user a good front-end interface. On the other hand some users say that the learning curve is quite hard at the beginning (in a big research center this is not a big problem..).
- 3. SciPy/NumPy/Matplotlib are libraries that work in the field of big data analysis written for the general purpose language Python. It is a quite immature technology, but it is freely available, and uses a general purpose and widely diffused language like Python.
- 4. ROOT is an object-oriented program and library developed by Cern, released the first time in 2003 (the process of development started in 1994 and continuously updated until now). It was originally designed for particle physics data analysis and contains several features specific to this field, but it is also used in other applications such as astronomy and data mining.

For the AEgIS experiment ROOT is the chosen software to carry out the activities of data analysis. The reason is that this software has been specifically tailored to meet the requirements of the analysis applied to particle physics. Another advantage of Root is that there are a lot of libraries created during the years related to the activities of the experiments of the Cern and it is nearly impossible rebuilt them from scratch with another software.

ROOT development was started by René Brun and Fons Rademakers in 1994 (but a more extended and precise list of collaborators is accessible here https://root.cern.ch/root/htmldoc/guides/users-guide/ ROOTUsersGuide.html#preface). The ROOT's user guide start with this prefaction, that explains in detail how ROOT was born.: "In late 1994, we decided to learn and investigate Object Oriented programming and C++ to better judge the suitability of these relatively new techniques for scientific programming. We knew that there is no better way to learn a new programming environment than to use it to write a program that can solve a real problem. After a few weeks, we had our first histogramming package in C++. A few weeks later we had a rewrite of the same package using the, at that time, very new template features of C++. Again, a few weeks later we had another rewrite of the package without templates since we could only compile the version with templates on one single platform using a specific compiler. Finally, after about four months we had a histogramming package that was faster and more efficient than the well-known FORTRAN based HBOOK histogramming package. This gave us enough confidence in the new technologies to decide to continue the development. Thus was born ROOT." (Taken from https://root. cern.ch/root/htmldoc/guides/users-guide/ROOTUsersGuide. html#preface)

This software is partially released under GPL (this means that everyone is allowed to use, redistribute and change the software, but any changes made must also be licensed under the GPL), and partially under LGPL (The LGPL is similar to the GPL, but is more designed for software libraries where you want to allow non-GPL applications to link to your library and utilise it).

ROOT is an object oriented framework that aims to solve problems related to high-energy physics. To better understand what is ROOT is important to start with understanding what is a framework: in IT a framework is a structure that helps the programmers providing them a set of already working utilities and services (for example, I/O, graphics, etcetera) often related to the sector in which the framework aims to work (for example, the services of a web development framework are related to the layout of a web page, to the organization of DBs etcetera). ROOT in particular offer services, functions, and packages related on the world of high-energy physics research, that allow to save much work. It provides, for example the possibility to use a computer's graphics subsystem and operating system

with abstraction, allowing the developer to create a graphical user interface and a GUI builder. Root provides also an abstract platform that allows to run C++ and command line scripts. More precisely Root includes (among others):

- 1. Libraries related to histogramming and graphing, that allows the developer to easily represent graphically statistical distributions. Also 3D visualization is allowed.
- 2. Libraries related to regression analysis.
- 3. Various statistics tools.
- 4. Libraries related to digital image processing and manipulation.
- 5. Libraries aimed to allow Parallel computing, (parallelize data analysis can be really useful to manage the complexity of the calculations).
- 6. The possibility of interfacing with Python and Ruby code in both directions.
- 7. The possibility of interfacing with Javascript, allowing the developer to access Root functionalities by a Browser.

In the following image we can see the structure of the ROOT's libraries:

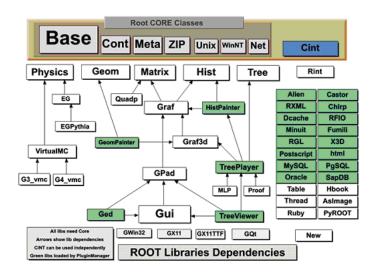


FIGURE 2.2: ROOT structure

Human-Machine Interface

labelChapter3

3.1 Human Machine Interaction principles

todo todo

3.2 Expected users

todo todo

3.3 Validation of gAn Web against HMI principles

todo todo

3.4 Modifications to match HMI principles

todo todo

Used Technologies

todotodo

4.1 Web interface vs Java Fx vs Xojo

todo todo

4.2 Used Technologies and Framworks

todo general

4.2.1 PHP

todo particular

4.2.2 Javascript

todo particular

4.2.3 Bootstrap

todo particular

4.2.4 Sass

todo particular

The resulting software

todotodo

5.1 A tour of the application

todotodo

5.2 Use example

todotodo

5.3 Some Screenshots

todotodo

Conclusions

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6.1 Conclusions