

UNIVERSITÉ CATHOLIQUE DE LOUVAIN

# LPHYS2131 : PARTICLE PHYSICS I

CMS OPENDATA LAB: MEASURING Z AND W BOSONS

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# 1 Introduction

## 1.1 Welcome to the LPHYS2131 wiki!

This project assembles material for the LPHYS2131 lab at UCLouvain. This wiki also documents the three sessions of the laboratory and provides some additional information.

The results that are obtained in this lab can be compared to the published cross-section measurement for the Z and W at the LHC, at 7TeV, by the CMS collaboration: Measurement of the Inclusive W and Z Production Cross Sections in pp Collisions at  $\sqrt{s} = 7$  TeV

### 1.1.1 Credits:

Thanks to Liliya Milenska, Gilles Perez and Martin Michel for their help in preparing this activity during their internship in summer 2015. Thanks to Victor Massart and Julien Touch  que for their help in porting the lab to 2011 data in summer 2016. Thanks to J  r  me de Favereau, Pavel Demin and Michele Selvaggi for their support.

## 2 Day 1 - Understanding the data and simulation workflows

### 2.1 Looking at the 2011 CMS data

Data comes in different formats, from bulky raw data to compact processed and skimmed data. In CMS, it goes like this (for that period):

RAW -> RECO -> AOD -> PAT -> Ntuples

- RAW data is a binary format. It is compact but proprietary and requires a lot of knowledge about the detector electronics to be decoded.
- RECO data contains higher level objects (tracks, calorimeter clusters, electrons, ...). It typically takes a lot of disk space (because low-level data is kept) but can be the starting point of data reprocessing.
- AOD (for Analysis Object Data) is a thinner data format where only the high-level calibrated objects are retained. It is centrally produced. These are the AOD files that CMS made public.
- PAT (for Physics Analysis Toolkit) is an even more simplified dataformat, typically produced by each analysis team. They typically contain only the subset of objects needed for a given study.
- Ntuples are simple data tables generally containing simple float quantities. Very compact, these are the files that we usually use in the interactive analysis.

In this lab, we will inspect AOD files with the event display tool of CMS, then produce a small ntuple. In the second lab, we will analyze larger ntuples produced by our IT team in advance of the lab.

#### 2.1.1 Event visualization

In a fresh terminal, run the following commands. Make here a choice between electrons (first file) and muons (second file).

```
cms-shell
cd CMSSW_5_3_32/src/
. /cvmfs/cms.cern.ch/cmsset_default.sh
cmsenv
cmsShow root://eospublic.cern.ch/eos/opendata/cms/Run2011A/DoubleElectron/AOD/120ct2013-v1/20000/0014CE62-9C3E-E311-8FCC-00261894389F.root
cmsShow root://eospublic.cern.ch/eos/opendata/cms/Run2011A/DoubleMu/AOD/120ct2013-v1/20000/045CCED6-033F-E311-9E93-003048678F74.root
```

1. What kind of events do you see?
2. Do you see electrons or muons?
3. You may try to filter the collection, for example asking for one or more muons... note that it takes time.

#### 2.1.2 Processing data to create a ntuple

The data directory contains a sample of type "PAT". It contains some preprocessed events, similar to the AOD just visualized, but filtered and slimmed down (only the

high-level info). We will process it and create a ntuple with the information relevant for this lab.

Take some time to look at the code in `Labo/WeakBosonsAnalyzer/src/WeakBosonsAnalyzer.cc`. Most of the job is done in the method `WeakBosonsAnalyzer::analyze`. This analysis code accesses the information in the PAT (using the CMS library) and create a simple file that can be read by the ROOT standalone software.

To run the code on few events:

```
cd Labo/WeakBosonsAnalyzer/test
cmsRun weakBosonsAnalyzer_cfg.py
```

By defaults, this runs on one file containing 21000 CMS event passing the double muon trigger. You can change that by editing the configuration file, `weakBosonsAnalyzer_cfg.py` to either point to another file or to run on a subset of events. When this is done, you can close the CMS shell.

We will now run Jupyter to look at the result using the ROOT package. In a new terminal:

```
conda activate cms-analysis
cd LPHYS2131/analysis
jupyter-lab Visualization.ipynb
```

Take some time to get familiar with the various quantities in the ntuple.

## 2.2 Simulating Monte Carlo (MC) events to compare to

To understand what the data contains, a standard approach consist in simulating event using Monte Carlo techniques to reflect the best of our understanding of the theory. By emulating the detector response and the analysis workflow, we obtain files that can be compared to data.

In this case, we will use MadGraph to simulate events of interest ( $pp \rightarrow Z \rightarrow l+l-$  and  $pp \rightarrow W \rightarrow l\nu$ ), pass them through a simplified detector simulation (Delphes) and produce a ntuple with the same format as for data.

### 2.2.1 Running MadGraph

Start MadGraph in a new terminal:

```
cms-shell
cd CMSSW_7_1_1/src/
. /cvmfs/cms.cern.ch/cmsset_default.sh
cmsenv
cd ../../
cd MG5_aMC_v2_4_3/
./bin/mg5_aMC
```

Generate Drell-Yann events:

```
MG5_aMC>generate p p > l+ l-
MG5_aMC>output ppNeutralCurrents
MG5_aMC>open index.html
MG5_aMC>launch ppNeutralCurrents
```

On the first prompt, enable "Pythia".

*PYTHIA is a computer simulation program for particle collisions at very high energies. Some of the features PYTHIA is capable of simulating are Hard and soft interactions, Parton distributions, Initial/final-state parton showers, Multiple interactions, Fragmentation and decay, etc. PYTHIA is used here to perform the hadronization of final state quarks, and to simulate both initial state and final state radiations.*

*Various parameters can be modified in the run card. This goes from the number of events (the default 10k is perfect in our case), some cuts (for example on the minimum  $P_t$  of some particles), or the beam energy.*

Then, repeat the operation to generate W bosons leptonic decays:

```
MG5_aMC>generate p p > l- vl~
MG5_aMC>add process p p > l+ vl
MG5_aMC>output ppChargedCurrents
MG5_aMC>open index.html
MG5_aMC>launch ppChargedCurrents
```

Again, on the first prompt, enable "Pythia".

*By default, and as just described, the simulation is performed at leading order (LO). To perform the simulation at next-to-leading order (NLO) it is enough to add "[QCD]" at the end of the process string. In that case, Pythia is not used, as the hadronization is performed internally by another method adapted to NLO.*

Then, we quit:

```
MG5_aMC>quit
```

Finally, we will unzip the results:

```
gunzip ppChargedCurrents/Events/run_01/tag_1_pythia_events.hep.gz
gunzip ppNeutralCurrents/Events/run_01/tag_1_pythia_events.hep.gz
```

### 2.2.2 Running Delphes

You just generated MC events, with sets of particles as they would emerge from the collisions. The next step is to emulate the detector response. We use for that Delphes, a standalone public tool that emulates a simplified version of the CMS detector.

```
cd ../Delphes-3.3.2
cp -r /LPHYS2131/analysis/delphes_card_CMS_mod.tcl cards/
./DelphesSTDHEP cards/delphes_card_CMS_mod.tcl ppChargedCurrents.root ../MG5_aMC_v2_4_3/ppChargedCurrents/Events/run_01/tag_1_pythia_events.hep
./DelphesSTDHEP cards/delphes_card_CMS_mod.tcl ppNeutralCurrents.root ../MG5_aMC_v2_4_3/ppNeutralCurrents/Events/run_01/tag_1_pythia_events.hep
```

### 2.2.3 Visualizing the events

We can now look at the events as we did for data:

```
root -l examples/EventDisplay.C'("cards/delphes_card_CMS.tcl","ppChargedCurrents.root")'  
root [0] .q
```

What are the similarities and differences between the two types of events?

### 2.2.4 Processing the events to generate a ROOT ntuple

Finally, we will produce a ntuple similar to the one we have for data, using the `CreateTreeFromDelphes` script. You can briefly look at it and then run the conversion:

```
cp ~/LPHYS2131/analysis/CreateTreeFromDelphes.C .  
root -l  
root [0] gSystem->Load("libDelphes.so")  
root [1] .L CreateTreeFromDelphes.C++  
root [2] CreateTreeFromDelphes("ppChargedCurrents.root","delpheAnalysisW.root")  
root [3] CreateTreeFromDelphes("ppNeutralCurrents.root","delpheAnalysisZ.root")  
root [4] .q
```

You can look at the resulting files... how does it compare to what we have for data?

## 3 Day 2 - Measuring the Z boson and determining the luminosity

### 3.1 Goals

In this lab, we want to

- see the Z boson
- measure the mass and width; understand the limitations of the approach
- determine the purity (estimate the background)
- determine the luminosity, from the number of Z events observed
- study the jet multiplicity if these events, and compare to MC predictions

Half of the class will work with electrons, the other half with muons. We will then compare.

### 3.2 Preparation for the lab

We have to copy some files to the work area (more precisely, we will create symbolic links to the files available). When this is done, we can start Jupyter, that will be used for the rest of this lab.

```
conda activate cms-analysis
cd LPHYS2131/analysis
./prepare.sh
jupyter-lab
```

### 3.3 Selection of the Z events

We have ntuples for data and Monte Carlo events (Drell-Yann events produced with MadGraph). Using these files, one can determine the selection criterias needed to get a pure sample, keeping the efficiency high. Monte Carlo files generated by CMS using a detailed simulation of the detector are provided in addition to the sample that was produced in the first lab.

We are looking for Z bosons decaying into either electrons or muons. How would you select such events based on the content on the ntuples produced on day 1?

In Jupyter, open the `visualization.ipynb` in labo2. That notebook contains already the skeleton of the analysis selection. Many more possibilities are available (like 2D plots, etc.) but should not be needed for this activity.

A natural selection will consist in asking two leptons of the same flavor. A cut on the isolation and on the Pt of these leptons might also be a good idea. Where should we cut? You can make up your mind by comparing data to Monte Carlo, interactively. Would you suggest additional cuts? What about jets and missing transverse momentum?

When preparing the cuts, keep in mind that the Monte Carlo simulation is imperfect. Don't blindly cut on quantities there without making sure that this makes also sense on data.



### 3.4 Fit of the data

Even with a good selection, there may be remaining background events. To estimate this, we will do a fit of the data with two contributions:

- the invariant mass Z events, either from the MadGraph Monte Carlo distribution or from an analytical form
- an ad hoc background distribution

This will give us an estimate of the purity, and depending on the fit a measure of the Z boson mass and width.

The fit is easily done with MINUIT. The `Zlineshape.ipynb` notebook will guide you in the fitting procedure. Things that should be adapted in the script are: the input files, the binning, the cuts, and the analytical form of the background contribution.

#### 3.4.1 Determination of the luminosity

Knowing the number of selected events, the purity, the Z cross-section (from MadGraph) and the selection efficiency, it is easy to get the luminosity:

$$N\_Z = \sigma * L * \text{efficiency} = N\_data * \text{Purity}$$

For this, we have of course to assume that the efficiency in data and simulations are the same.

- if you use the sample produced on day 1, the cross-section is given by MadGraph and can be found in the MadGraph folder where you produced the MC sample: `file:///home/cms-opendata/MG5_aMC_v2_4_3/ppNeutralCurrents/crossx.html`. Remember that we produced 10000 events.
- in case you use the official CMS sample, you must know that it has a cross-section of 2475pb, and contains originally 40'000'000 events.
- the efficiency is given by the number of MC events passing your cuts divided by the number of generated events.
- `N_data`, `Purity`, and `N_Z` (the product of the two) are given by the Jupyter notebook. There are different ways to determine these numbers, and the precision will depend on your selection.

Good to know: we run on a subset of the 2011 data. The total luminosity recorded in 2011 was 5.55/fb, but the sample available for this lab (2011A, corresponding to data up to 21/08/2011) represents 2.676/fb.

#### 3.4.2 Study of the Z lineshape

One of the fit that you can perform relies on an analytical description of the Z mass peak. A proper function will allow you to determine the Z boson mass and its width. Does it match your expectations?

Does the result depend on the fit function? How would you take that into account when giving the uncertainty?

### 3.5 Other studies

1. Look at events with one jet. What is the angle between the lepton pair and the jet? How does the Pt of the jet and the Pt of the lepton pair compare. Is that expected? What is the origin of that jet?
2. Look at the jet multiplicity. Does the MC reproduce the data? What if you generate events at NLO? Try to fit with a power law or an exponential. Why does it work?
3. Compare electrons and muons samples. Are the results the same?

## 4 Day 3 - Measuring the W boson and determining its cross section

### 4.1 Goals

In this lab, we want to

- see the W boson
- estimate the background from data using a control region
- determine the W boson cross-section from the event yield and the luminosity determined previously

Half of the class will work with electrons, the other half with muons. We will then compare.

### 4.2 Preparation for the lab

We have to set up the jupyter environment.

```
conda activate cms-analysis
cd LPHYS2131/analysis
jupyter-lab
```

### 4.3 Selection of the W events

We have ntuples for data and Monte Carlo events (W<sup>+</sup> and W<sup>-</sup> events produced with MadGraph). We will use a full-simulation sample from CMS instead of the Delphes sample from day 1 to make sure that the efficiency is under control. We can then compare with the Delphes sample. Using these files, one can determine the selection criteria needed to get a pure sample, keeping the efficiency high.

While we were looking at the invariant mass for the Z, this is not possible here, as the neutrino escapes undetected. Still, by momentum conservation we can estimate the transverse component of its momentum. Using the lepton 4-vector and the missing transverse momentum, one defines the transverse mass. This variable will be used here.

You will notice that it is more difficult to get a pure sample. It is also more difficult to simulate that background, mostly driven by QCD events with leptons. We will therefore measure the background shape in a control region (to be defined) and then do a template fit using the background shape from data on one side and the signal shape from MC on the other side.

We are looking for W bosons decaying into either one electron or one muon, plus one neutrino. How would you select such events based on the content on the ntuples produced on day1?

A natural selection will consist in asking one lepton and some missing transverse momentum. A cut on the isolation and on the Pt of the lepton might again be a good idea. Where should we cut? You can make up your mind by comparing data to Monte

Carlo, interactively in `Labo3/visualization.ipynb`. Would you suggest additional cuts? What are the expected specificity of the kinematic of a W event?

#### 4.4 Choice of the control region where to measure the background

As already discussed, there is an irreducible background that comes from QCD events, with leptons produced in jets or wrongly identified (fakes).

Since the background cannot easily be estimated from simulations, we propose to define a control region where it can be measured. That control region should be

- as similar as possible to the signal region
- as much as possible free of any signal event Especially, as we will use the control region to derive the shape of the background for the transverse mass, the kinematics should be the same in both regions.

Given all of the above, how would you define the control region?

#### 4.5 Determination of the cross-section

The `labo3/Wyield.ipynb` Jupyter notebook will guide you through the fit of the signal and background contributions. Things that should be adapted in the script are:

- the binning (min/max/number of bins)
- the cuts for the signal and control regions (that you just decided)

That script performs the following actions:

- Fills histogram for MC and data for the chosen variable with chosen cuts.
- Fills an histogram with an estimate of the background, obtained from data in a control region.
- Fits MC to data.

Knowing the number of selected events, after background subtraction, the luminosity measured in lab 2 (or the luminosity announced by CMS), and the selection efficiency (from MC), it is easy to get the cross-section:

$$N_W = \sigma * L * \text{efficiency} = N_{\text{data}} * \text{Purity}$$

$$\sigma = N_{\text{data}} * \text{Purity} / (L * \text{efficiency})$$

For this, we have of course to assume that the efficiency in data and simulations are the same.

- `N_data`, `Purity`, and `efficiency` are given by the jupyter notebook,
- the luminosity is the one obtained on day 2 (or the luminosity announced by CMS),
- the MC sample used contains 78'347'691 events, for a cross-section of 2.584E4 pb.

How does the cross-section obtained that way compare to the value announced by the MC? That one can be found on that local link: `file:///home/cms-opendata/MG5_aMC_v2_4_3/ppChargedCurrents/crossx.html`

#### 4.6 Other studies

1. What is the charge ratio between  $W^+$  and  $W^-$ ? Do the results depend on the charge?
2. How could we determine the  $W$  boson mass?
3. Does the jet multiplicity behave as for the  $Z$  boson studied last time? Do the results depend on the jet multiplicity?
4. Look at events with one electron and one muon. Do we expect such events? Ask for two additional jets. What are these events in data?

## 5 The analysis tree

List and meaning of the branches stored in the analysis TTree:

Branch name	Content
nMuons	Number of muons
MuonsPt	Transverse momentum of muons
MuonsEta	Pseudorapidity of muons
MuonsPhi	Azimuthal angle of muons
MuonIsolation	Relative isolation of muons
nElectrons	Number of electrons
ElectronsPt	Transverse momentum of electrons
ElectronsEta	Pseudorapidity of electrons
ElectronsPhi	Azimuthal angle of electrons
ElectronIsolation	Relative isolation of electrons
nJets	Number of jets
JetsPt	Transverse momentum of jets
JetsEta	Pseudorapidity of jets
JetsPhi	Azimutal angle of jets
MET_pt	Missing transverse momentum
MET_phi	Azimuthal angle of the missing transverse momentum
MET_eta	Pseudorapidity of the missing momentum (MC only, should not be used)
invMass	Invariant mass of lepton pairs
transvMass	Transverse mass built of the leading lepton + MET
dileptonPt	Transverse momentum of the dilepton pair
dileptonEta	pseudorapidity of the dilepton pair
dileptonPhi	azimuthal angle of the dilepton pair
dileptondeltaR	distance in eta-phi between the two leading leptons
dileptondeltaPhi	distance in phi between the two leading leptons
Wcharge	charge of the leading lepton

## 6 To save images for the report

The various tools that we use have built-in features allowing to save plots or images. There is no need to take screenshots. In some cases, these tools will even produce files better suited for documents than the version shown on screen.

### 6.0.1 CMS event display

In Fireworks, the best option is to go to **File > Export Images of all views**. To get the best result on a printed document, you can switch to a white background in **Edit > Change Background Color**. We recommend to save pictures in a new folder on the desktop.

### 6.0.2 Jupyter

In Jupyter, just shift-right-click on the picture and select "save as".

### 6.0.3 Delphes event display

In Delphes, plots can be saved as in ROOT. The only subtlety is that the File menu is hidden on top of the plot. To make it appear, first undock the subpicture by selecting **Action > UndockWindow**. Then pass the mouse on top of the picture, which should reveal the menu. Once you are done, you can close the window to dock it again.

## 7 Linux Primer

The machine that we run for this lab (either the lab PCs or the virtual machine) runs a modern version of Linux. Interested in going further with linux? Give it a try with one recent distribution (Fedora, Ubuntu, Mint, or other).

Linux is not much different from any operating system, but we will use a lot the console to navigate through the directories and issue commands. We then run in what is called a shell.

### 7.1 Shell commands

Some basic commands that will be useful:

```
pwd (print current directory);  
cd DIRECTORY (move to directory DIRECTORY)  
ls [-ltrh] DIRECTORY (list files in directory DIRECTORY)  
cp [-r] FILE DIRECTORY (copy file FILE to directory DIRECTORY)  
mv FILE1 FILE2 (rename FILE1 into FILE2)  
rm [-ir] FILE (remove FILE)  
mkdir DIRECTORY (create new directory)
```

When navigating accross directories, remember that "." represents the current directory. ".." represents the parent directory. Compared to the OSes of the MS family, the directory separator is "/" instead of ".".

### 7.2 Editing files

To edit files, you have several options:

- **vi or vim**  
vi runs fully in the terminal. It is available on most of the UNIX-like systems, which makes it the default choice on many systems. MadGraph uses vi to edit the config files. A dedicated page on this wiki is available to help you start with vi.
- **emacs**  
emacs is another very well known editor in the UNIX universe. It is more user-friendly than vi.
- **leafpad**  
leafpad looks like the notepad in Windows. It is definitely the simplest and easiest editor in the list.

All the three can be launched directly from the terminal or from the "start menu" at the bottom left of the screen.

### 7.3 About the copy/paste

If there is one thing easy in Linux, this is the copy paste. Just forget about ctrl-c/ctrl-v, that's not needed anymore. To copy, just highlight the text. It is "in the mouse". To



paste, click on the central button (on the wheel).

## **7.4 More ...**

You will find much more on the web, for example at <https://lifehacker.com/a-command-line-primer-f>