



# MIAD Analysis 5

*A tool devoted to  
phenomenological investigations*

*Eric CONTE, Benjamin FUKS, Manuel KRAFT, Guillaume SERRET*

CMS France @ IPHC Strasbourg  
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## Overview

Writing an analysis step-by-step in the  
normal mode

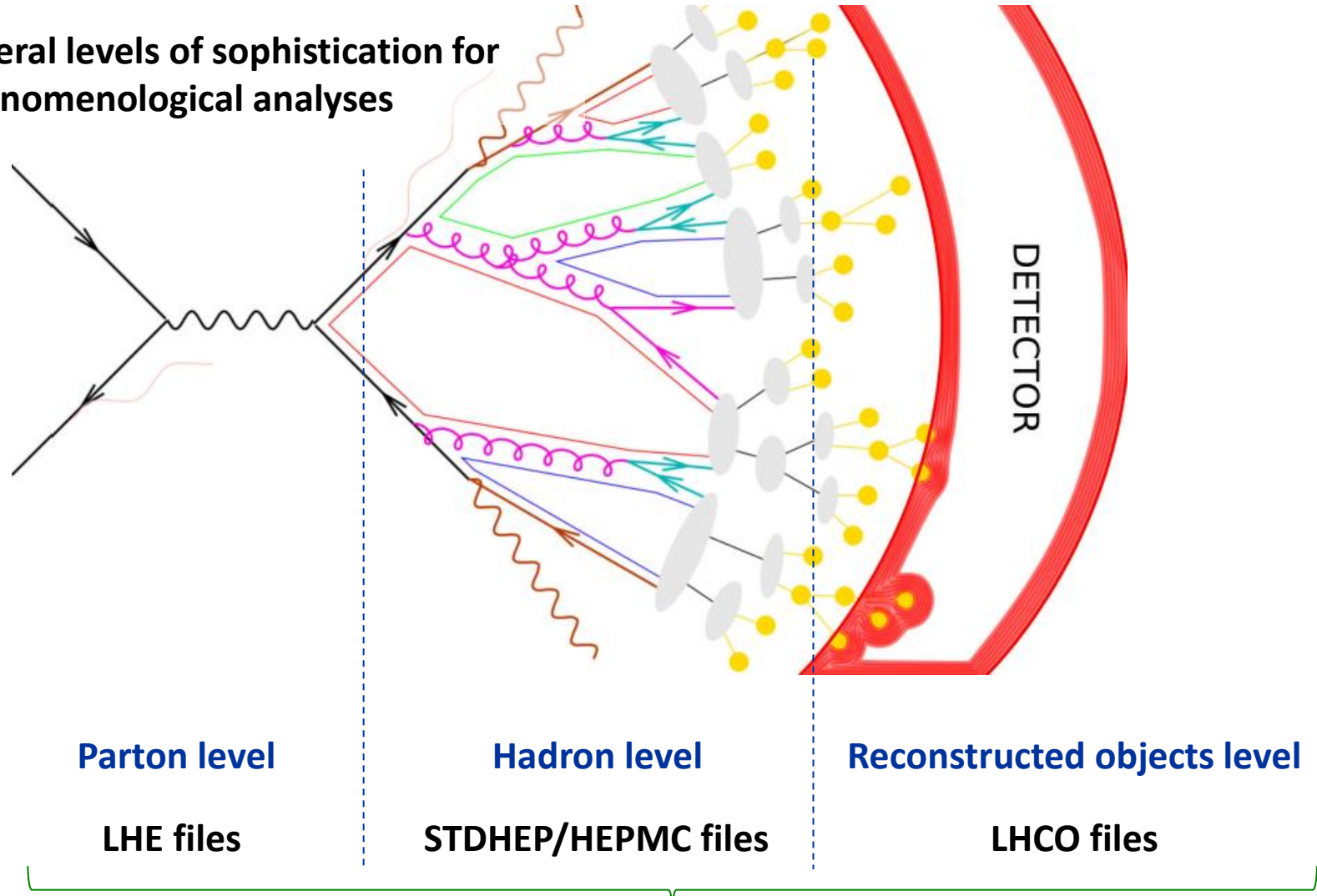
Interface to the FastJet package

Spoilers: what's new in the next releases ?

## Summary

# Overview

Several levels of sophistication for phenomenological analyses



**A unique framework : MadAnalysis 5**

## Scope:

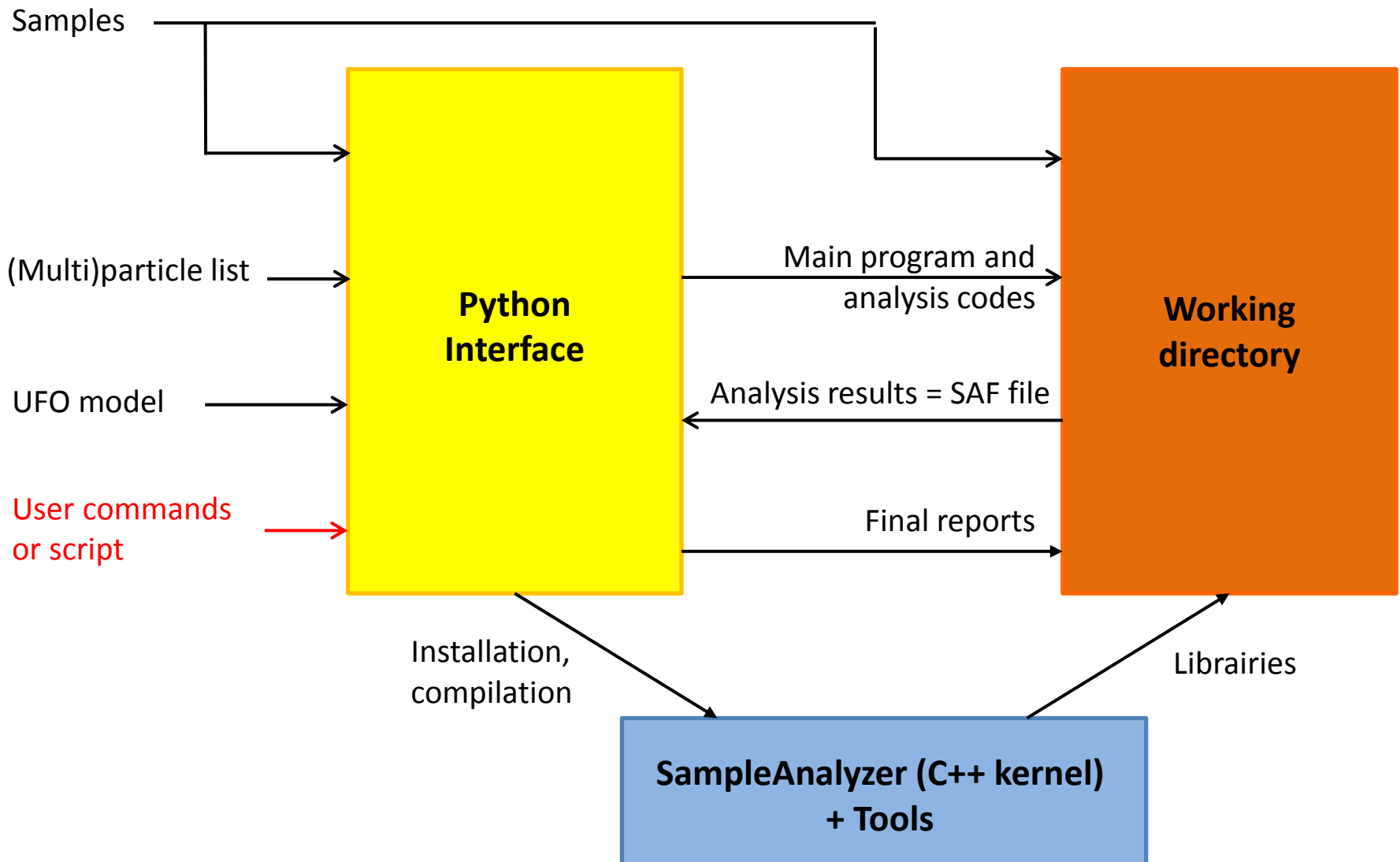
- Reading of signal and background event files
- Definition of various selection cuts on the input samples.
- Production of histograms for different distributions.
- Results of the analysis summed up by a S/B-like ratio table.

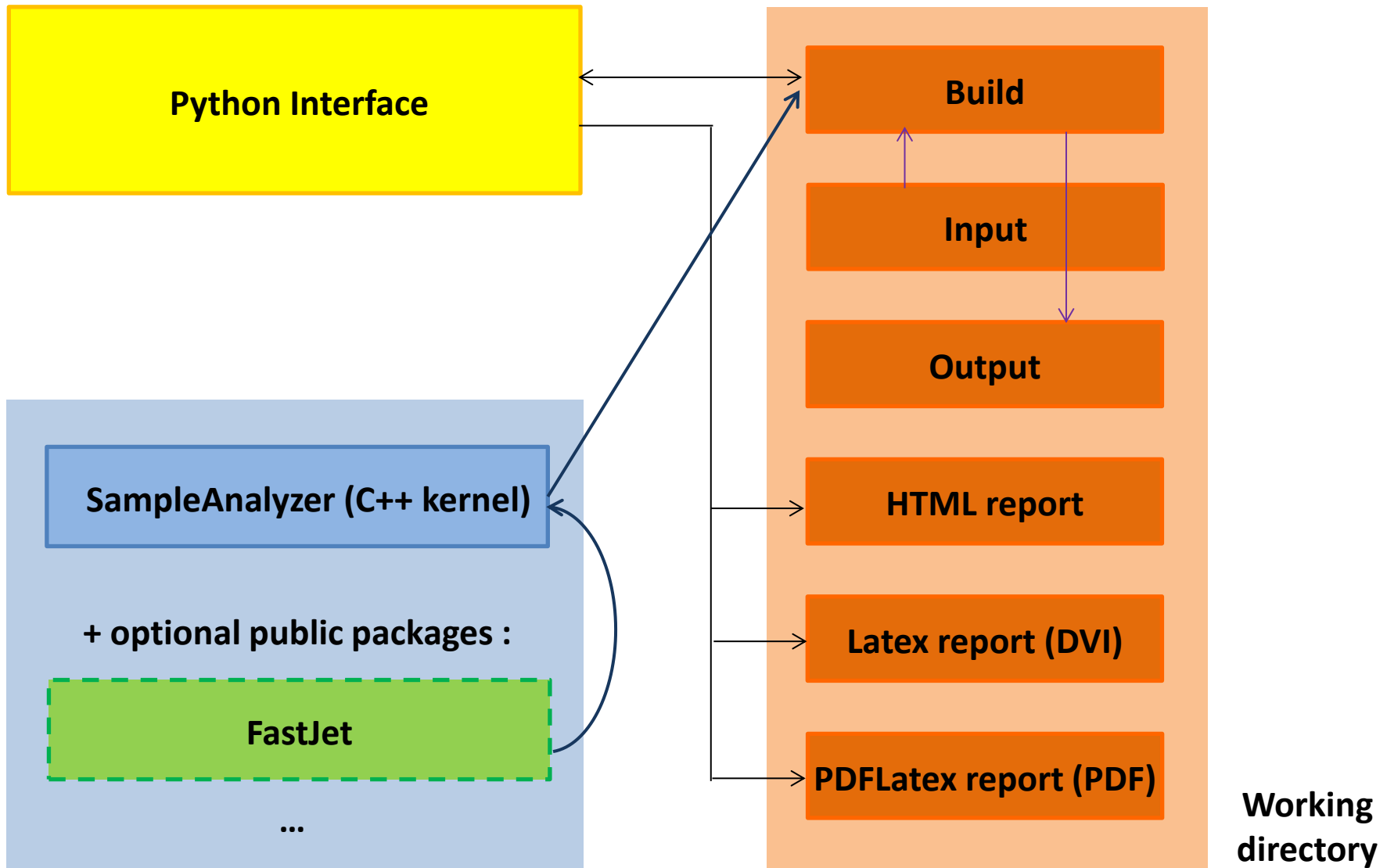


## Relevant features of MadAnalysis 5 design:

- Study at any sophistication level (parton, hadron, reconstructed)
- Supporting any event file format (STDHEP, HEPMC, LHE, ...)
- **User-friendly** → professional analyses in a simple way
- **Flexible**: no limit on the analysis complexity
- **Easy** to maintain and to validate



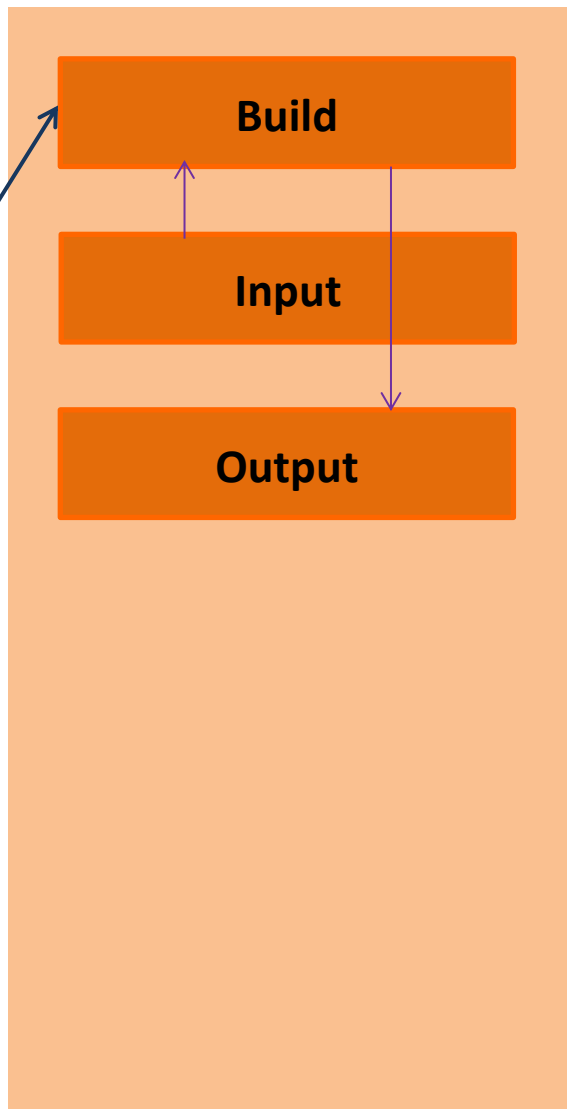
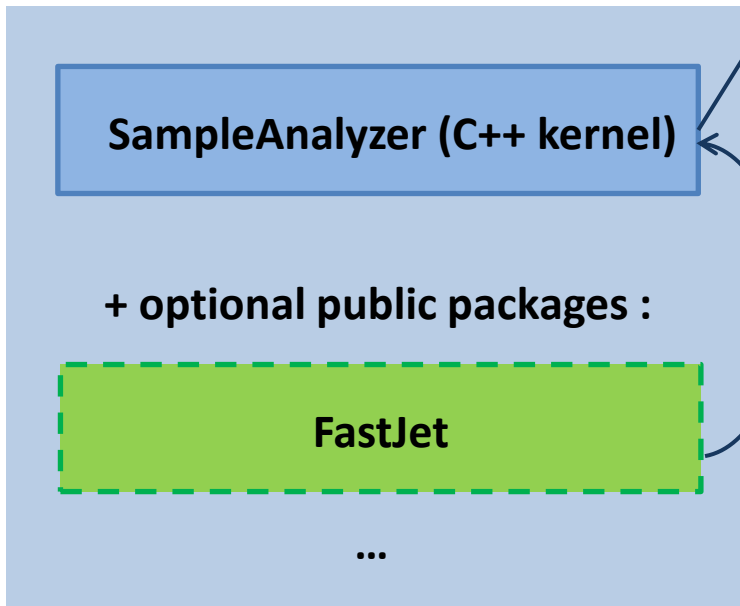




Working directory

MadAnalysis has an **expert mode** (developer-friendly) :

- C++ programming within the SampleAnalyzer framework.
- The Python interface creates a blank analysis as a starting point.



Working directory

## Installation step

- Requirements

Mandatory	Optional
<b>Python</b> 2.6 or a more recent version (but not the 3.X series)	<b>zlib</b>
<b>GNU GCC</b> compiler	<b>Latex</b> / PDFLatex
<b>ROOT</b> 5.27 or a more recent version	<b>FastJet</b> 3.0 or a more recent version

- Downloading MadAnalysis 5:

- From the official website <http://madanalysis.irmp.ucl.ac.be> (tarball to untar)
- From MadGraph 5 interface (**available soon**)



## First start of MadAnalysis 5:

- Execution

Parton level	Hadron level	Reconstructed objects level
bin/ma5 or bin/ma5 -P	bin/ma5 -H	bin/ma5 -R

- Initial sequence:

- Step 1: Testing all dependencies
- Step 2: Compiling the static library of SampleAnalyzer
- Step 3: Locating MadGraph and importing the list of particles and multiparticles

## Defining new particles and multiparticles

- Particles are defined by **labels**, which could point to one or several **PDG-id**.
- SM and MSSM labels are automatically loaded at the starting of MadAnalysis.
- The user can define his own labels :

```
ma5> define mu = mu+ mu-
```

- All labels defined in a UFO model can be loaded too.

## Importing datasets

- For MadAnalysis, a **dataset** is a collection of samples which will be merged.
- All sample files are stored in a dataset.

```
ma5> import tt*.lhe
```

```
ma5> import tt*.lhe as ttbar  
ma5> import Wj*.lhe as Wjets
```

- Possibility to tag datasets as **signal** or **background**.

## Defining a selection : plots and/or cuts

- **Histograms**

- Observable can be related to the event or the properties of a particle
- Plethora of observables: N, E, ET, M, MT, P, PT, PX, PY, PZ, THETA, ETA, ..., **ALPHAT**
- Combining particles

```
ma5> plot MET
ma5> plot PT(mu)
```

```
ma5> plot M(mu+ mu-)
```

- **Cuts : selecting / rejecting events**

```
ma5> reject MHT < 50
ma5> select N(mu) >= 2
```

- **Cuts : selecting / rejecting a particle or a combination**

```
ma5> select (mu) PT > 50
ma5> select 80 < M (mu+ mu-) < 100
```

## Defining a selection : plots and/or cuts

*Several options or syntaxes allow to extend the potential of MadAnalysis.  
Some examples:*

- By default, a combination is interpreted as the vector sum of momenta. This interpretation can be changed by adding a prefix to the observable label. For instance : `vPT`, `sPT`, `dsPT`, `dvPT`, `rPT`
- List of observables specific to the reconstructed object level : `ISOL`, `HE_EE`, `NTRACKS`, ...
- Selecting a particle **according to its rank in energy** (or to other observables)

```
ma5> plot(mu+[1])
```

- Selecting a particle **according to its history** (requirements on mother, grand-mother ...)

```
ma5> plot(mu+ < w+ < t~)
```

## Launching the analysis:

This can be done by the command **submit**

- Creating a working directory (with a default name if no name is specified)
- Compiling the C++ job
- Launching the analysis over the different samples contained in the datasets

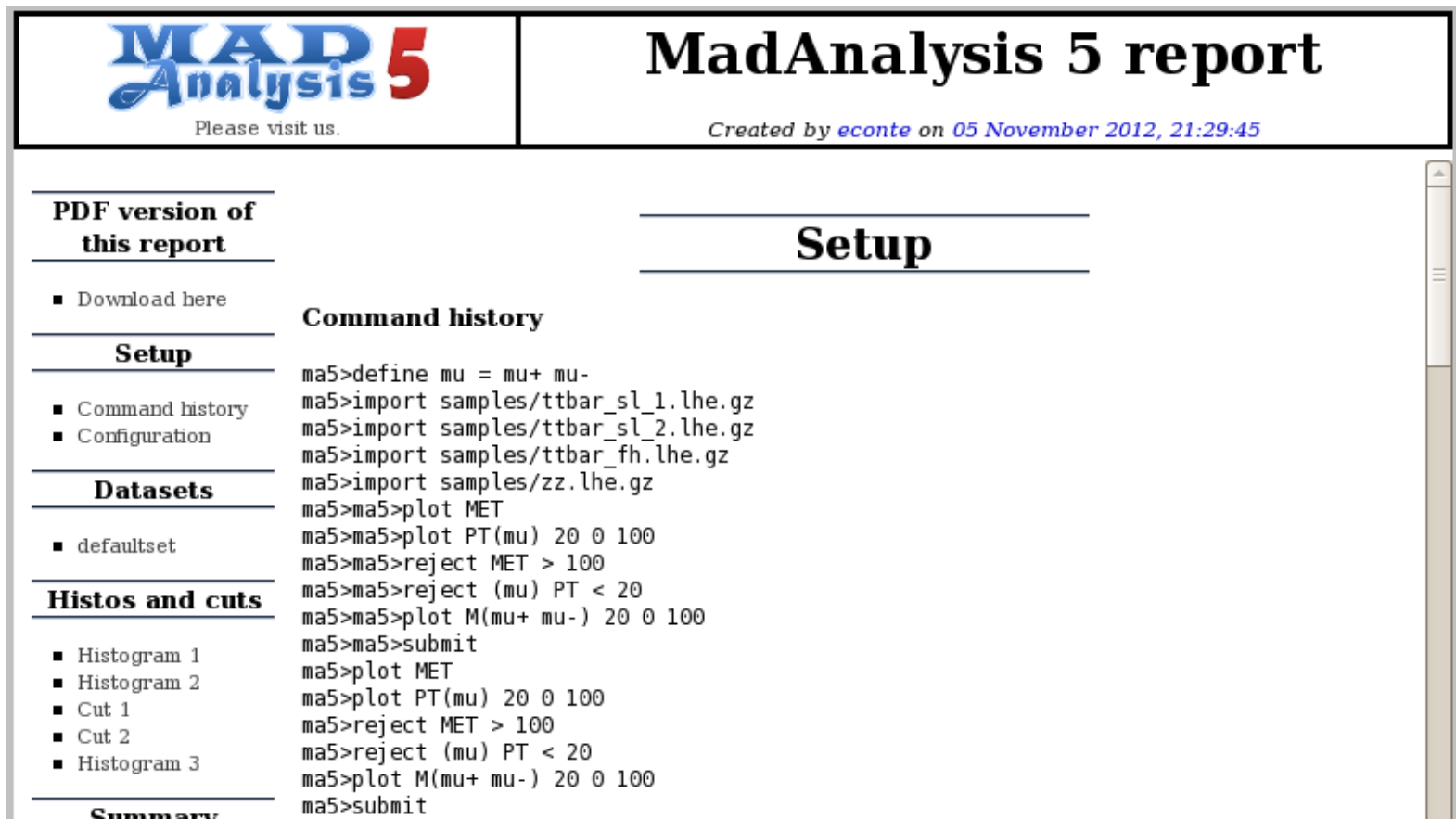
```
* SampleAnalyzer 2.0 for MadAnalysis 5 - Welcome.  
* Option choices: selecting analysis = 'MadAnalysis5job'.  
* Extracting the following sample files:  
* 1/4 ~/samples/ttbar_sl_1.lhe.gz  
* => sample produced by MadGraph.  
* => Number of processed events: 1000.  
* . . .
```

If you modify, after the submission, the analysis or the layout of the plots , the results can be updated in **an optimized way** by the command **resubmit**.

## Opening a generated report:

The command **open** displays the HTML report of the last job created.

Other reports can be opened by: **open workindir/PDF** or **open workingdir/DVI**



**MAD Analysis 5**  
Please visit us.

**MadAnalysis 5 report**  
Created by *econte* on 05 November 2012, 21:29:45

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**PDF version of this report**

- Download here

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**Setup**

- Command history
- Configuration

---

**Datasets**

- defaultset

---

**Histos and cuts**

- Histogram 1
- Histogram 2
- Cut 1
- Cut 2
- Histogram 3

---

**Summary**

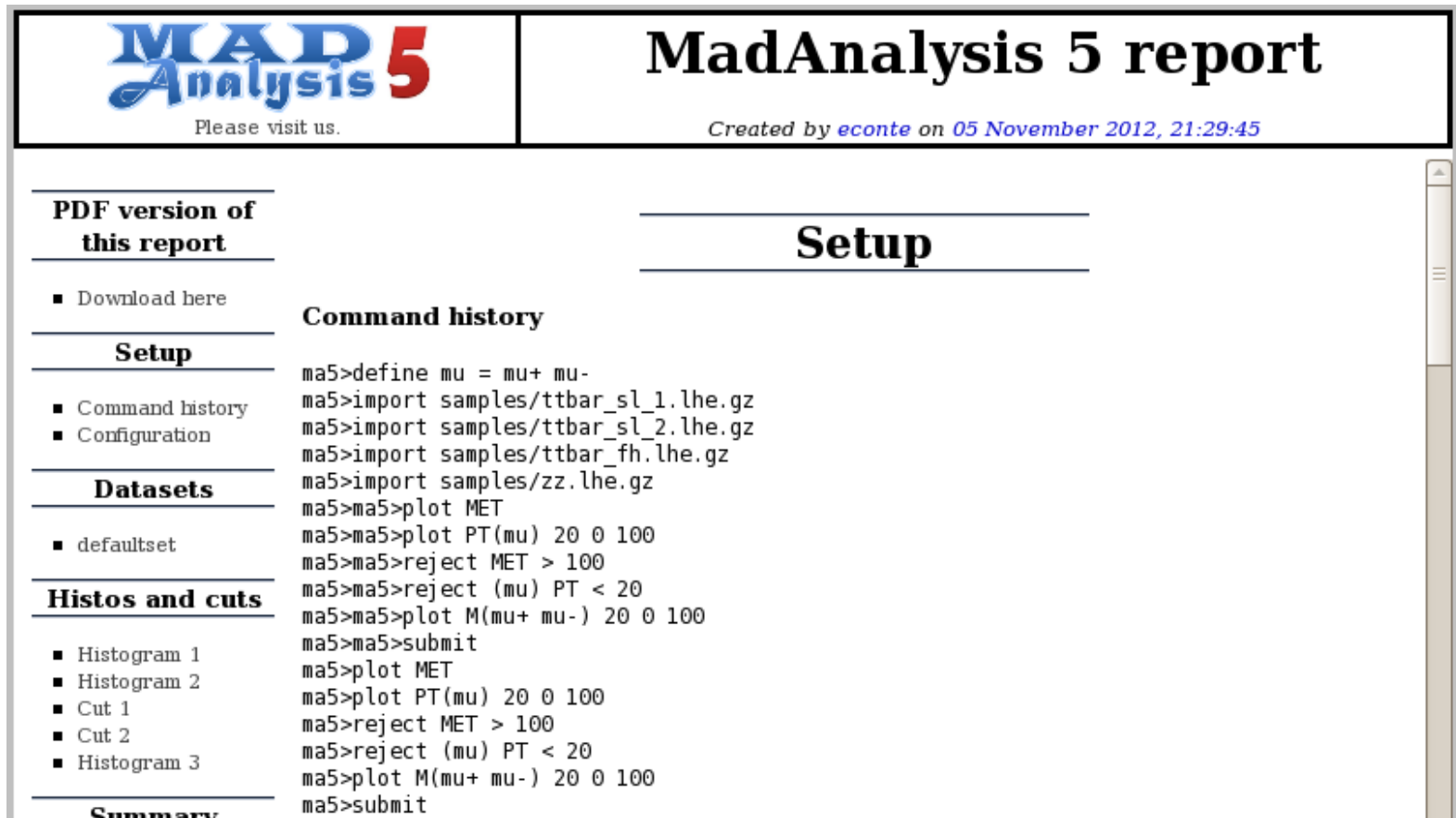
**Command history**

```
ma5>define mu = mu+ mu-
ma5>import samples/ttbar_sl_1.lhe.gz
ma5>import samples/ttbar_sl_2.lhe.gz
ma5>import samples/ttbar_fh.lhe.gz
ma5>import samples/zz.lhe.gz
ma5>ma5>plot MET
ma5>ma5>plot PT(mu) 20 0 100
ma5>ma5>reject MET > 100
ma5>ma5>reject (mu) PT < 20
ma5>ma5>plot M(mu+ mu-) 20 0 100
ma5>ma5>submit
ma5>plot MET
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**Summary**

**Command history**

```
ma5>define mu = mu+ mu-
ma5>import samples/ttbar_sl_1.lhe.gz
ma5>import samples/ttbar_sl_2.lhe.gz
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ma5>plot PT(mu) 20 0 100
ma5>reject MET > 100
ma5>reject (mu) PT < 20
ma5>plot M(mu+ mu-) 20 0 100
ma5>submit
```

## Opening a generated report:

Details on sample information

Path to the event file	Nr. of events	Cross section (pb)	Negative wgts (%)
mg5_merged.hep.gz	5116	162.0	0.0

Path to the event file	Nr. of events	Cross section (pb)	Negative wgts (%)
amcatnlo.hw.hep.gz	9993	313	8.4

- By default, MadAnalysis5 takes into account the **event-weights** contained in the samples. If no event-weights are available, there are set to one.
- MadAnalysis5 handles **negative event-weights** produced by NLO generators and propagates them properly into the uncertainties calculation (**whatever the ROOT version**).



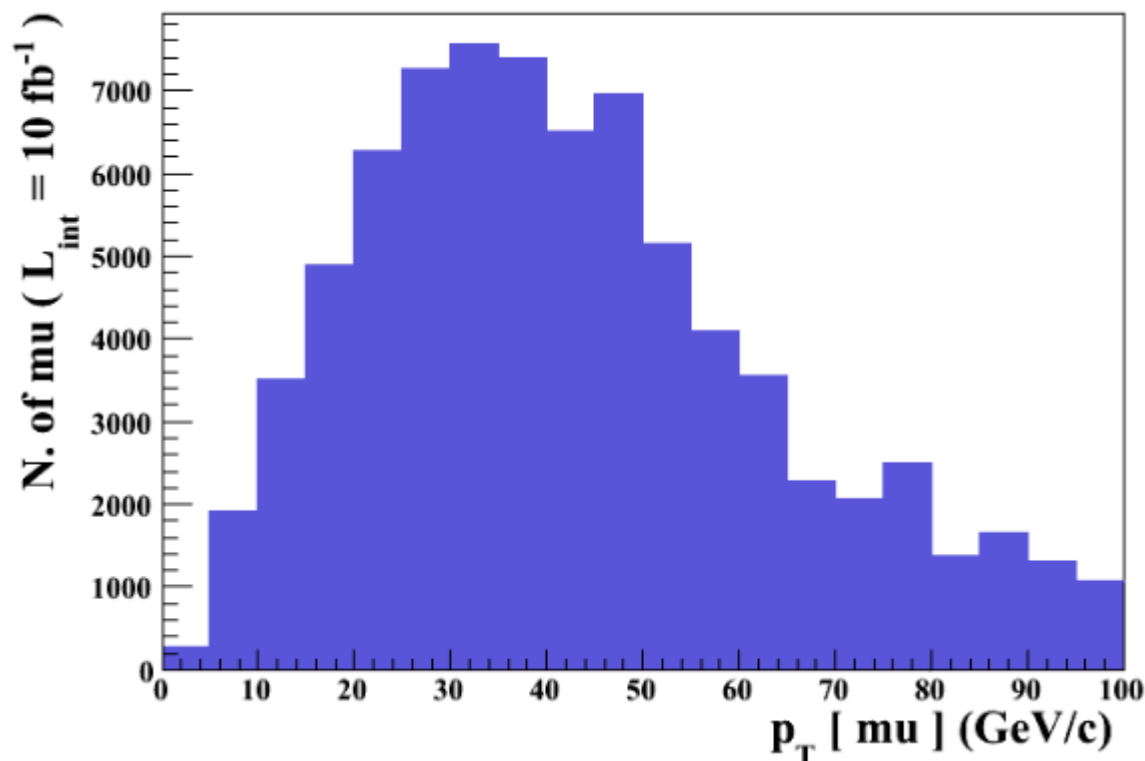
# Writing an analysis step-by-step

## Opening a generated report:

## Details on histogramming

Dataset	Integral	Entries / events	Mean	RMS	Underflow	Overflow
defaultset	82747	0.752	42.8177	21.36	0.0	1.296

Statistics table



- the cross section of the sample is automatically extracted from the sample
- Integrated luminosity is by default  $10 \text{ pb}^{-1}$ . This value can be set by the user:

```
ma5> set main.lumi =
```

## Jet clustering algorithms:

This can be done with  
MadAnalysis 5 in reco mode  
**bin/ma5 -R**

- Need to install **FastJet** and interface it to MadAnalysis  
→ Just one command line from the Python interface !

```
ma5> install fastjet
```

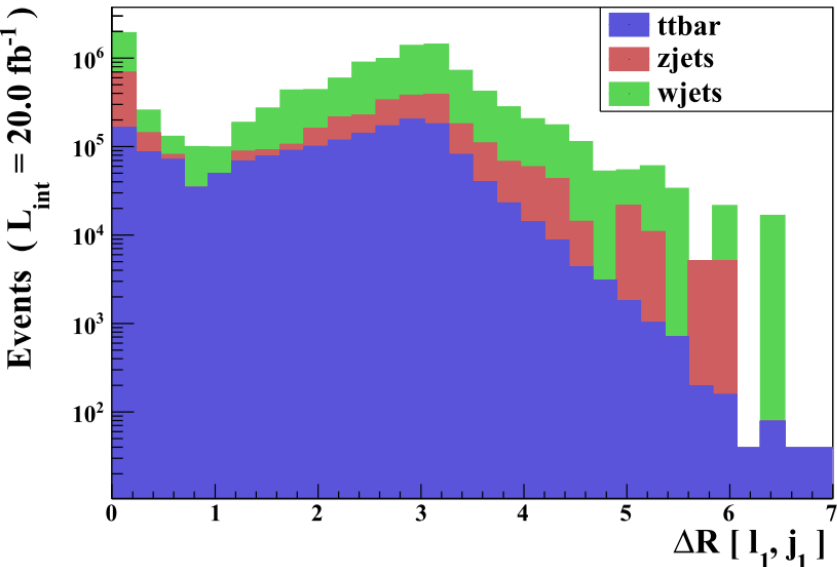
- Large selection of jet algorithms

```
ma5> set main.clustering.algorithm =  
antikt          cdfjetclu          genkt   kt          siscone  
cambridge      cdfmidpoint        gridjet none
```

- Adopting a jet algorithm → new options (the **algorithm parameters**)

```
ma5> set main.clustering.algorithm = antikt  
ma5> set main.clustering.ptmin    = 5  
ma5> set main.clustering.radius   = 1
```

## Jet clustering algorithms:



```
ma5> set main.clustering.algorithm = antikt
ma5> set main.clustering.ptmin = 5
ma5> set main.clustering.radius = 1
```

```
ma5> import ttbar*.hep.gz as ttbar
ma5> import wjets.hep.gz as wjets
ma5> import zjets.hep.gz as zjets
```

```
ma5> set ttbar.xsection = 139.6
ma5> set wjets.xsection = 35678
ma5> set zjets.xsection = 10319
ma5> set main.lumi = 20
```

```
ma5> select (l) PT > 20
ma5> reject (j) PT < 50
ma5> reject THT < 200
ma5> plot DELTAR(l[1],j[1]) 30 0 7 [logY]
```

```
ma5> submit
ma5> open
```

## Simplified LHE output:

- After the jet clustering procedure, events could be saved in a simplified LHE.
- Enabling this option can be done with one command line in your analysis:

```
ma5> set main.outputfile = "eric.lhe.gz"
```

- The conventions used in the simplified LHE format are :

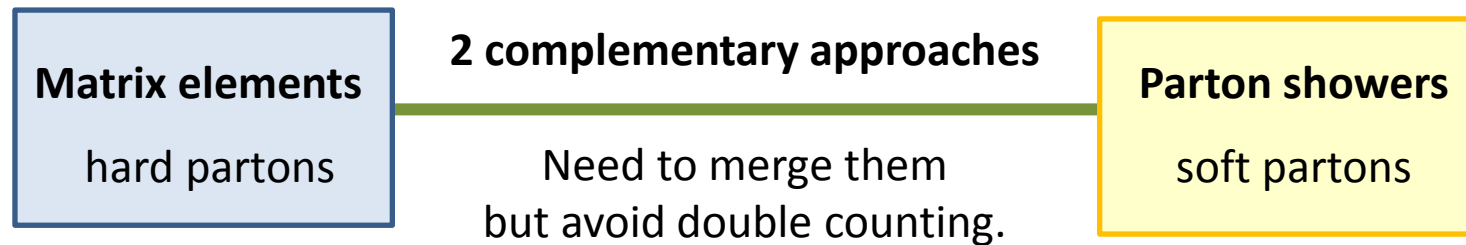
Light jets	PDG id = 21
c jets	PDG id = 4
b jets	PDG id = 5
MET	PDG id = 12

Electrons	PDG id = 11
Muons	PDG id = 13
Hadronic taus	PDG id = 15

This functionality replaces the old module called "HEP2LHE", provided in the MadGraph package

Thanks to Adam Alloul for validation

## Jet-merging validation plots:



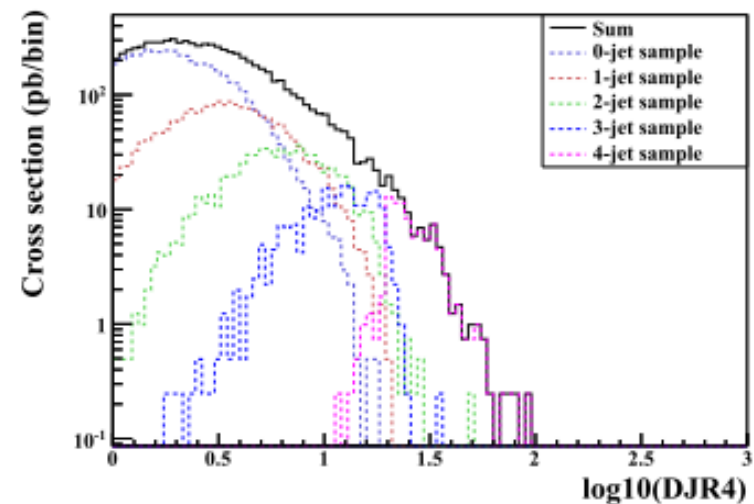
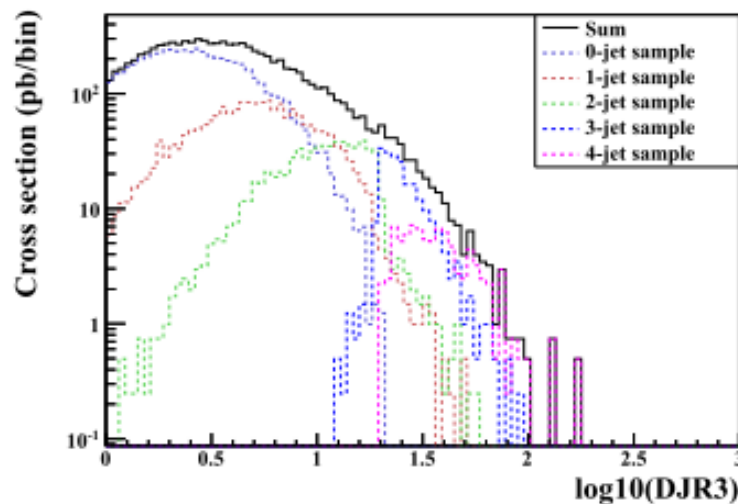
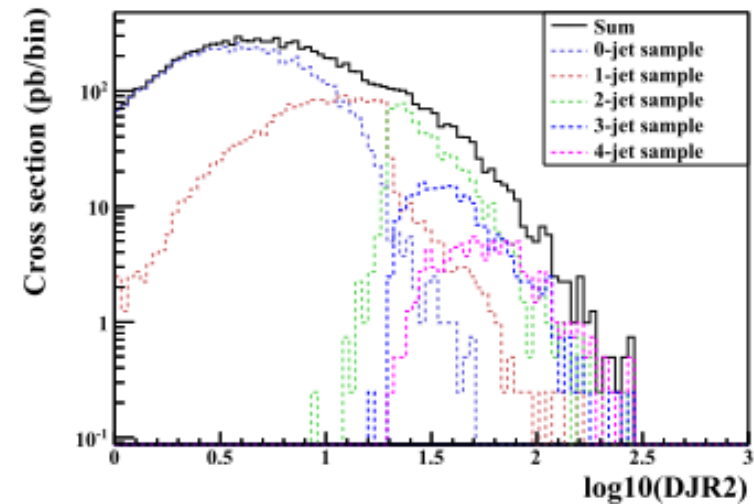
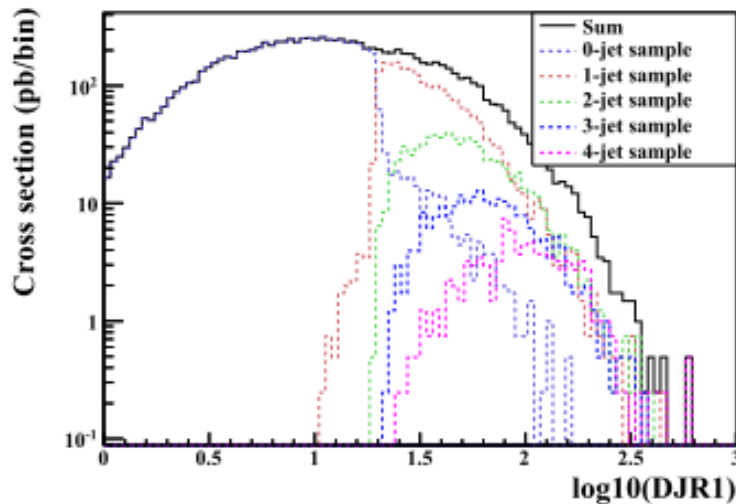
- **Merging matrix-elements with 0, 1, 2, 3, .... extra jets**
  - Study of the smoothness of the differential jet rate (DJR) distributions.
    - The scale for which an event goes from a  $N \rightarrow N+1$  jet configuration.
    - Extremely sensible to the merging procedure.
  - This validates the choices for the merging parameters.
- **Running MadAnalysis 5 in hadron-level mode: `bin/ma5 -H`**

```
ma5> set main.merging.check = true
ma5> set main.merging.njets = 4
```

# Interface to the FastJet package

## Jet-merging validation plots:

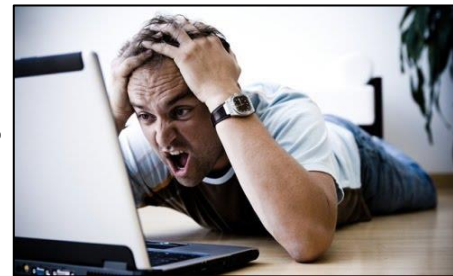
Example of Z production with 0, 1, 2, 3, 4 extra jets



# What's new in the next releases ?

## In the next release (soon):

- **New internal structure → the expert mode is enriched and more flexible:**
  - Plot and cut data are saved as a simplified XML file (**SAF format**).
  - The main loop of the analysis can be modified by the user. Possibility to add several analyses, filters or jet clustering algorithms in a same job.
- **Removing ROOT libraries from python interface: adopting a new graphical style for plots.**
- **Fixing minor bugs.**
- **Completing the user manual by tutorials.**



# What's new in the next releases ?



## In the longer-term: the main milestones

- Ready to perform phenomenological investigations @ NLO:
  - Using information associated to events produced by **aMC@NLO**, in particular reweighting parameters.
  - Interfacing showering/hadronization package (**Pythia 6/8, Herwig6/++**) and configuring it automatically wrt the generation conditions.
- Interfacing the existing detector fast-simulation packages (**Delphes**) and developing new ones related to top physics (in priority: *boosted top reconstruction*).
- Pipelining the data-flow “showering-fastsim-analysis”: the full processing chain is applied event by event (avoiding intermediate sample files).
- Optimizing the processing time by paralleling jobs.





- **MadAnalysis 5 = a unique framework for different levels of analysis:**  
Parton level , Hadron level and Reconstructed objects level
- Designed to be **fast** and **user-friendly**.
- Two ways of using the program
  - **Normal mode:** python interface with intuitive commands.
  - **Expert mode:** requiring programming skills (C++, ROOT).
- **Interfaced to FastJet**, MadAnalysis 5 can:
  - launch a specified **jet clustering** sequence to hadronic events.
  - display the **plots dedicated to check the jet merging procedure**.
- Exepected for 2013: interfaces to **fast-simulation** and **showering** packages.



ma5team@iphc.cnrs.fr  
<http://madanalysis.irmp.ucl.ac.be>  
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