



# MIAD Analysis 5 *status and news*

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GDR Terascale @ Palaiseau  
2-4 June 2014

*Thank you*

to the whole MadAnalysis user community

especially

A. ALLOUL, J. ANDREA, L. BASSO , S. BEIN, G. CHALONS,

K. DE CAUSMAECKER, U. GOERLACH, S. KRAML, S. KULKARNI,

K. MATAWARI, L. MITZKA, D. SENGUPTA

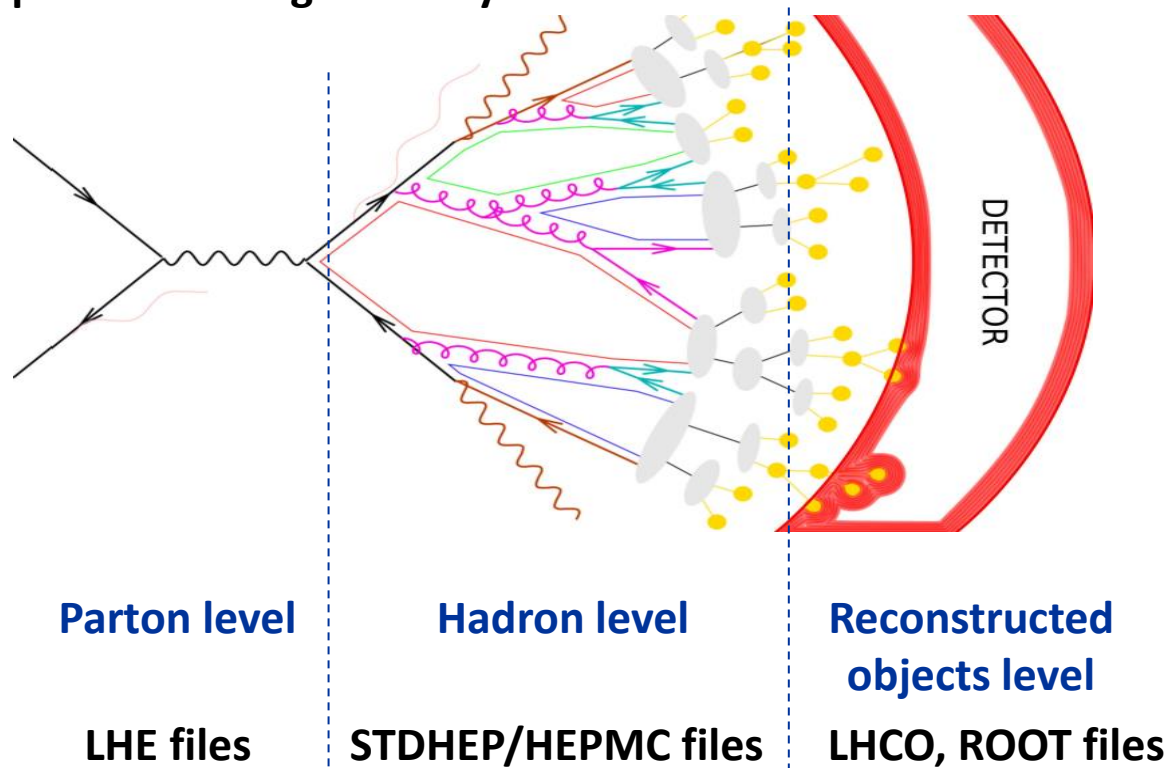
*Based on MadAnalysis 5  
v1.1.10 and v1.1.11beta*

- 1. Reminder : what is MadAnalysis?**
- 2. The normal mode and its metalanguage**
- 3. Fast-simulation packages**
- 4. The expert mode and recasting an analysis**
- 5. The new validation suite**
- 6. Summary & perspectives**

# 1. Reminder

## Starting points of the project:

Several levels of sophistication for phenomenological analyses



### Relevant features of design

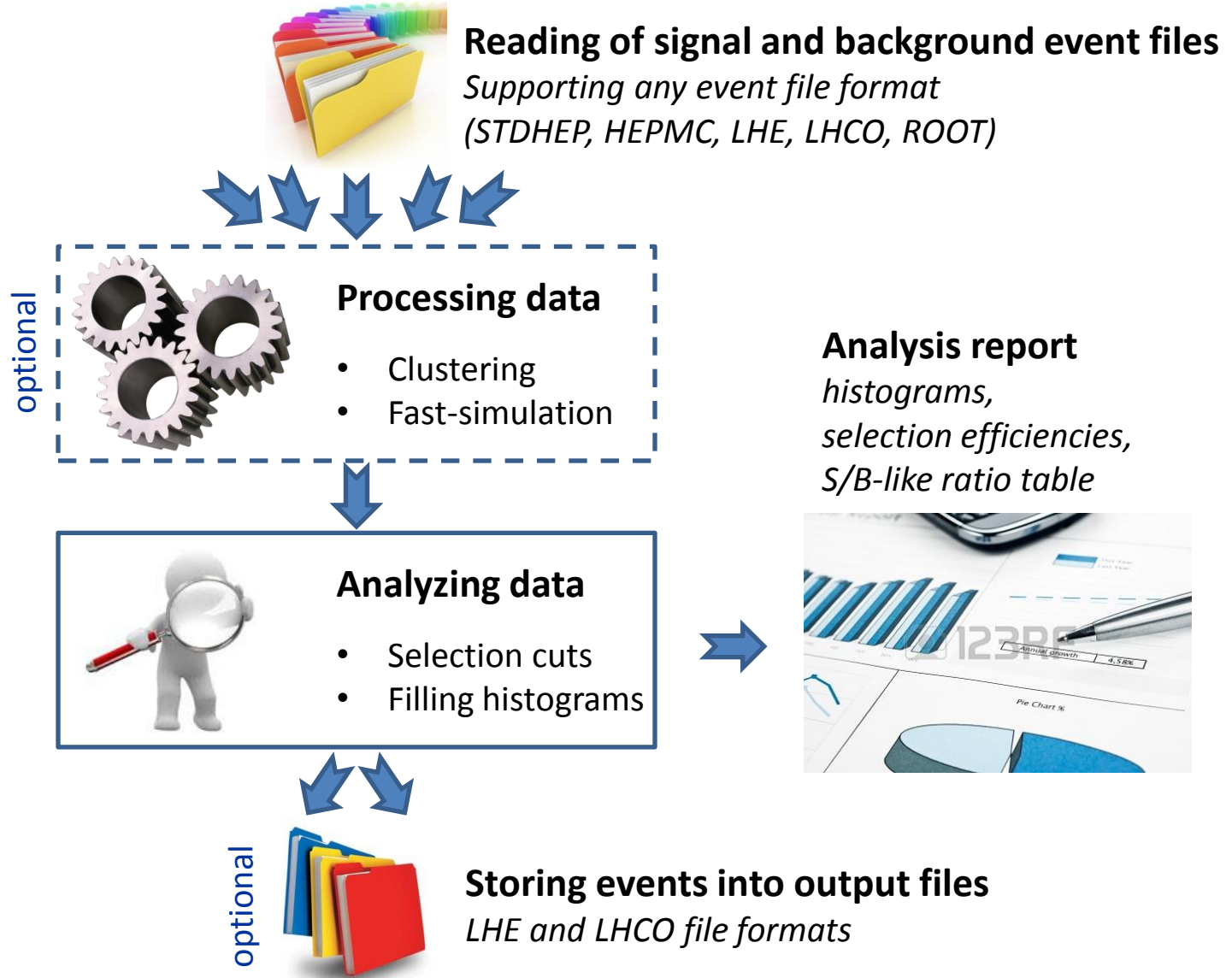
- User-friendly
- Flexible
- Efficient
- Easy to maintain

A unique framework : MadAnalysis 5

# 1. Reminder

## Scope:

Defining an analysis with a dedicated metalanguage





# 2. Normal mode

## User-friendly console with PYTHON



Just type:




`./bin/ma5`

```
Platform: Linux 2.6.18-348.12.1.el5 [Linux mode]
Reading user settings ...
Checking mandatory packages:
  - python [OK]
  - python library: numpy [OK]
  - g++ [OK]
  - GNU Make [OK]
  - Root [OK]
  - PyRoot libraries [OK]
Checking optional packages:
  - pdflatex [OK]
  - latex [OK]
  - dvipdf [OK]
  - zlib [OK]
  - FastJet [OK]
  - Delphes [OK]
  - Delphes-MA5tune [OK]
Checking the MadAnalysis library:
=> MadAnalysis libraries found.
=> MadAnalysis test program works.
*****
MadGraph 5 NOT found:
=> Particle labels from input/particles_name_default.txt
=> 87 particles successfully exported.
=> Multiparticle labels from
madanalysis/input/multiparticles_default.txt
=> Creation of the label 'invisible' (-> missing energy).
=> Creation of the label 'hadronic' (-> jet energy).
=> 8 multiparticles successfully exported.

ma5>_
```

# 2. Normal mode

## Compilation and other software tricks behind the scene = physicist-friendly

- **Inspection of your system:**
  - Autodetection of the required packages (g++, root)
  - Autodetection of the optional packages ( zlib, delphes, fastjet, ...)
  - Autodetection of MadGraph if it is installed.
- First time you used MadAnalysis: compilation of the core libraries  
**New architecture and recipe: more OS-independent** 
- **Debug mode could also used to investigate in case of difficulties.**   
→ `./bin/ma5 --debug`
- **Users could bypass or force a step of the MadAnalysis recipe**   
→ configuration file `madanalysis/inputs/user_configuration.dat`
- Optional libraries could be installed quickly from the Python interface with only one instruction.



# 2. Normal mode

## A metalanguage for describing your analysis

- Designed to be short and intuitive
- Tab completion is very useful!

Example through a tutorial

### Step 1: Defining new particles and multiparticles

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

### Step 2: Importing datasets

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

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

### Step 3: Defining plots

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

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

Plethora of observables: **N, E, ET, M, MT, P, PT, PX, PY, PZ, THETA, ETA, ..., ALPHAT**

New observables: **MT2, MT2W**



# 2. Normal mode

## A metalanguage for describing your analysis

- Designed to be short and intuitive
- Tab completion is very useful!

### Step 4: Defining cuts

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

selecting / rejecting events

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

selecting / rejecting a particle or a combination

### Step 5: Applying the analysis to the samples

```
ma5> submit
```

### Step 6: Displaying results


```
ma5> open
```

Example through a tutorial

# 2. Normal mode

## Reports

All results are gathered in a single report, available in HTML, PDF or DVI format.



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

---

### 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
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
```

# 3. Fast-simulation packages

MadAnalysis is interfaced to several fast-simulation detector packages

## Home made simulation based on FastJet

- Large choice of jet clustering algorithms
- Lepton isolation
- b-tagging (and c-tagging) (mis)efficiency



- Simple analysis.
- Study the impact of one detector effect on a result.

## Delphes

- Full functionalities of the last release of Delphes 3:
- ATLAS & CMS simulation
  - Possibility to include pile-up events



## Delphes MA5tune

- Special tuning of the Delphes package provided by MadAnalysis 5 for the CMS detector simulation



Analysis requiring realistic detector simulation



# 3. Fast-simulation packages



What's new in Delphes MA5 tune package?

- **Lepton isolation**

### Delphes3 strategy

- Determining what is a isolated and a non-isolated lepton during the simulation processing.
- Using this information in the particle flow algorithm.

### MA5tune strategy = Delphes2 strategy

- Let the user deciding what is an isolated lepton → optimization according to the analysis.
- Let the user remove double counting.

- **More realistic parametrization of the b-tagging(mis-)efficiency**

- Using official CMS results
- New track counting algorithm of Delphes is not used.

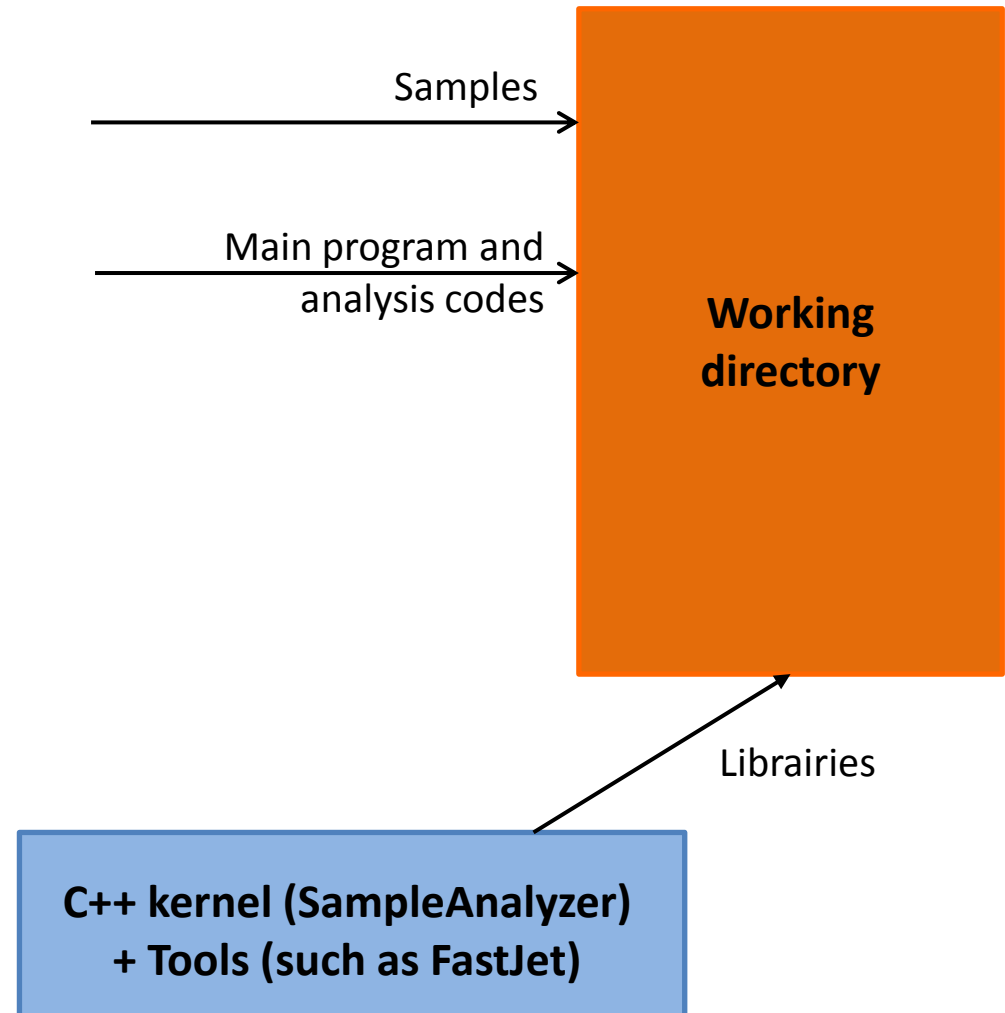
- **Optimization of the Delphes ROOT content** in order to decrease the size of the files. Relevant for heavy analysis requiring a lot of samples and statistics.

# 4. Expert mode

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

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

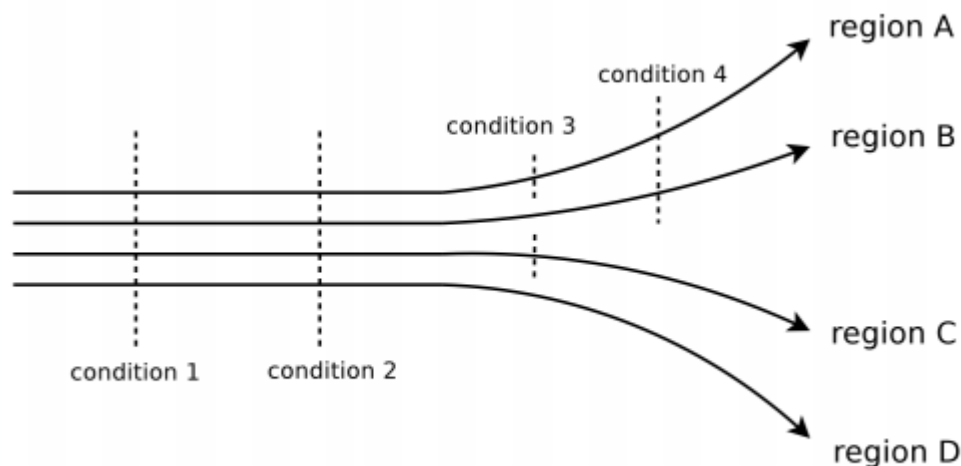
```
./bin/ma5 -e
```



# 4. Expert mode

## Extension of the expert mode for recasting a physics analysis:

- Access for all the information from the Delphes MA5 tune package.
- Support for multiple sub-analyses (signal and control regions)
  - cut-flow chart and plots for each region



E. Conte, B. Dumont, B. Fuks,  
C. Wymant, arXiv:1405.3982

See Guillaume Chalons talk for an application

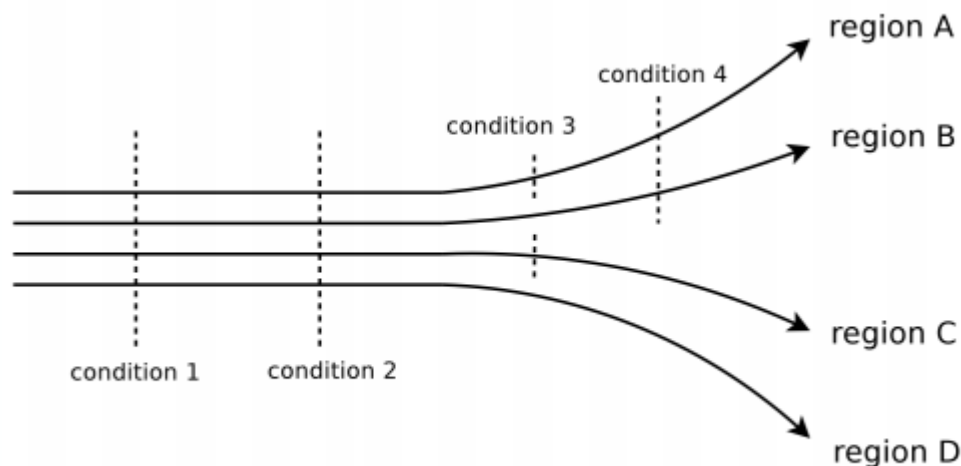
### Naive approach

```
if (condition 1)
{
  ...
}
if (condition 3)
{
  ...
  if (condition 4)
  {
    ...
  }
}
if (condition 4)
{
  ...
}
```

# 4. Expert mode

## Extension of the expert mode for recasting a physics analysis:

- Access for all the information from the Delphes MA5 tune package.
- Support for multiple sub-analyses (signal and control regions)
  - cut-flow chart and plots for each region



### MadAnalysis approach

```
if (condition1, region A+B+C+D)
{
  ...
}

if (condition 3, region A+C)
{
  ...
}

if (condition 4, region A+B)
{
  ...
}
```

**E. Conte, B. Dumont, B. Fuks,  
C. Wymant, arXiv:1405.3982**

See Guillaume Chalons talk for an application



# 5. Validation suite 2.0



**Validation is mandatory when a new version of MadAnalysis must be released.**

**The validation suite has been significantly improved:**

- Several MadAnalysis configurations are tested (presence of optional packages + release of ROOT).
- A huge collection of tests is launched on each configuration. Corresponding MadAnalysis output (reports or samples) are stored in a private database.
- Validation is performed by comparing test-by-test the results between the new and the previous MadAnalysis results.
  - Comparing numbers
  - Comparing plots



# 5. Validation suite 2.0



Validation success is summarized in a web page.

## Validation MadAnalysis 5

Date of HTML creation: 02/06/2014

Author: Eric Conte

Versions: root-5.34.10/v1.1.10beta\_v1.1.11beta/no\_options/plots

Number	Scripts' description	Matching (%)	Comparison log files	Scripts v1.1.10beta	Scripts v1.1.11beta	MA5 output log v1.1.10beta	MA5 output log v1.1.11beta
0	testing all observables with Plot and 0 particle	100	<a href="#">plot_0part.log</a>	<a href="#">plot_0part.ma5</a>	<a href="#">plot_0part.ma5</a>	<a href="#">plot_0part.ma5.log</a>	<a href="#">plot_0part.ma5.log</a>
1	testing all observables with Plot and 0 particle	100	<a href="#">plot_0part_H.log</a>	<a href="#">plot_0part_H.ma5</a>	<a href="#">plot_0part_H.ma5</a>	<a href="#">plot_0part_H.ma5.log</a>	<a href="#">plot_0part_H.ma5.log</a>
2	testing all observables with Plot and 0 particle	100	<a href="#">plot_0part_R.log</a>	<a href="#">plot_0part_R.ma5</a>	<a href="#">plot_0part_R.ma5</a>	<a href="#">plot_0part_R.ma5.log</a>	<a href="#">plot_0part_R.ma5.log</a>
3	testing all observables with Plot and 1 particle	100	<a href="#">plot_1part.log</a>	<a href="#">plot_1part.ma5</a>	<a href="#">plot_1part.ma5</a>	<a href="#">plot_1part.ma5.log</a>	<a href="#">plot_1part.ma5.log</a>
4	testing all observables with Plot and 1 particle	100	<a href="#">plot_1part_H.log</a>	<a href="#">plot_1part_H.ma5</a>	<a href="#">plot_1part_H.ma5</a>	<a href="#">plot_1part_H.ma5.log</a>	<a href="#">plot_1part_H.ma5.log</a>
5	testing all observables with Plot and 1 particle	100	<a href="#">plot_1part_R.log</a>	<a href="#">plot_1part_R.ma5</a>	<a href="#">plot_1part_R.ma5</a>	<a href="#">plot_1part_R.ma5.log</a>	<a href="#">plot_1part_R.ma5.log</a>
6	testing all observables with Plot and 1 particle	100	<a href="#">plot_2parts.log</a>	<a href="#">plot_2parts.ma5</a>	<a href="#">plot_2parts.ma5</a>	<a href="#">plot_2parts.ma5.log</a>	<a href="#">plot_2parts.ma5.log</a>
7	testing all observables with Plot and 1 particle	79	<a href="#">plot_2parts_H.log</a>	<a href="#">plot_2parts_H.ma5</a>	<a href="#">plot_2parts_H.ma5</a>	<a href="#">plot_2parts_H.ma5.log</a>	<a href="#">plot_2parts_H.ma5.log</a>
8	testing all observables with Plot and 1 particle	100	<a href="#">plot_2parts_R.log</a>	<a href="#">plot_2parts_R.ma5</a>	<a href="#">plot_2parts_R.ma5</a>	<a href="#">plot_2parts_R.ma5.log</a>	<a href="#">plot_2parts_R.ma5.log</a>
9	testing all observables with Plot and 3 particles	100	<a href="#">plot_3parts.log</a>	<a href="#">plot_3parts.ma5</a>	<a href="#">plot_3parts.ma5</a>	<a href="#">plot_3parts.ma5.log</a>	<a href="#">plot_3parts.ma5.log</a>
10	testing all observables with Plot and 3 particles (HADRON mode)	100	<a href="#">plot_3parts_H.log</a>	<a href="#">plot_3parts_H.ma5</a>	<a href="#">plot_3parts_H.ma5</a>	<a href="#">plot_3parts_H.ma5.log</a>	<a href="#">plot_3parts_H.ma5.log</a>
11	testing all observables with Plot and 3 particles	100	<a href="#">plot_3parts_R.log</a>	<a href="#">plot_3parts_R.ma5</a>	<a href="#">plot_3parts_R.ma5</a>	<a href="#">plot_3parts_R.ma5.log</a>	<a href="#">plot_3parts_R.ma5.log</a>

- **MadAnalysis 5 = a unique framework with two ways to use it:**
  - **Normal mode:** python interface with intuitive commands.
  - **Expert mode:** requiring programming skills (C++, ROOT).
- **Relevant features of MadAnalysis 5 design:**
  - **User-friendly** → professional analyses in a simple way.
  - **Flexible:** no limit on the analysis complexity.
  - **Easy** to maintain and to validate.
- **Main last developments:**
  - Delphes-MA5tune package for fast-simulation.
  - Support for recasting physics analysis.
  - A new validation suite.



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<https://launchpad.net/madanalysis5>  
Comput. Phys. Commun. 184 (2013) 222