



# MIAD Analysis 5 *status and news*

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## Overview of MadAnalysis 5 v1.1.9

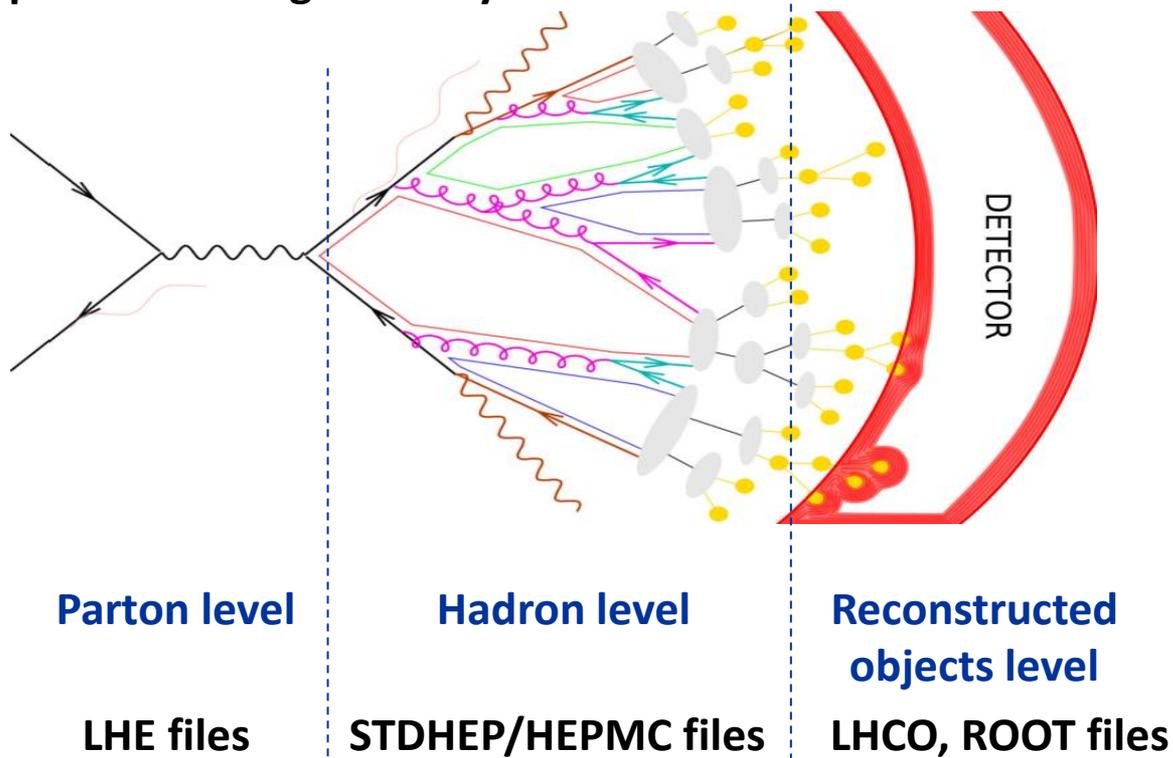
The normal mode and its metalanguage

Fast-simulation packages

Summary & perspectives

## Starting points of the project:

Several levels of sophistication for phenomenological analyses



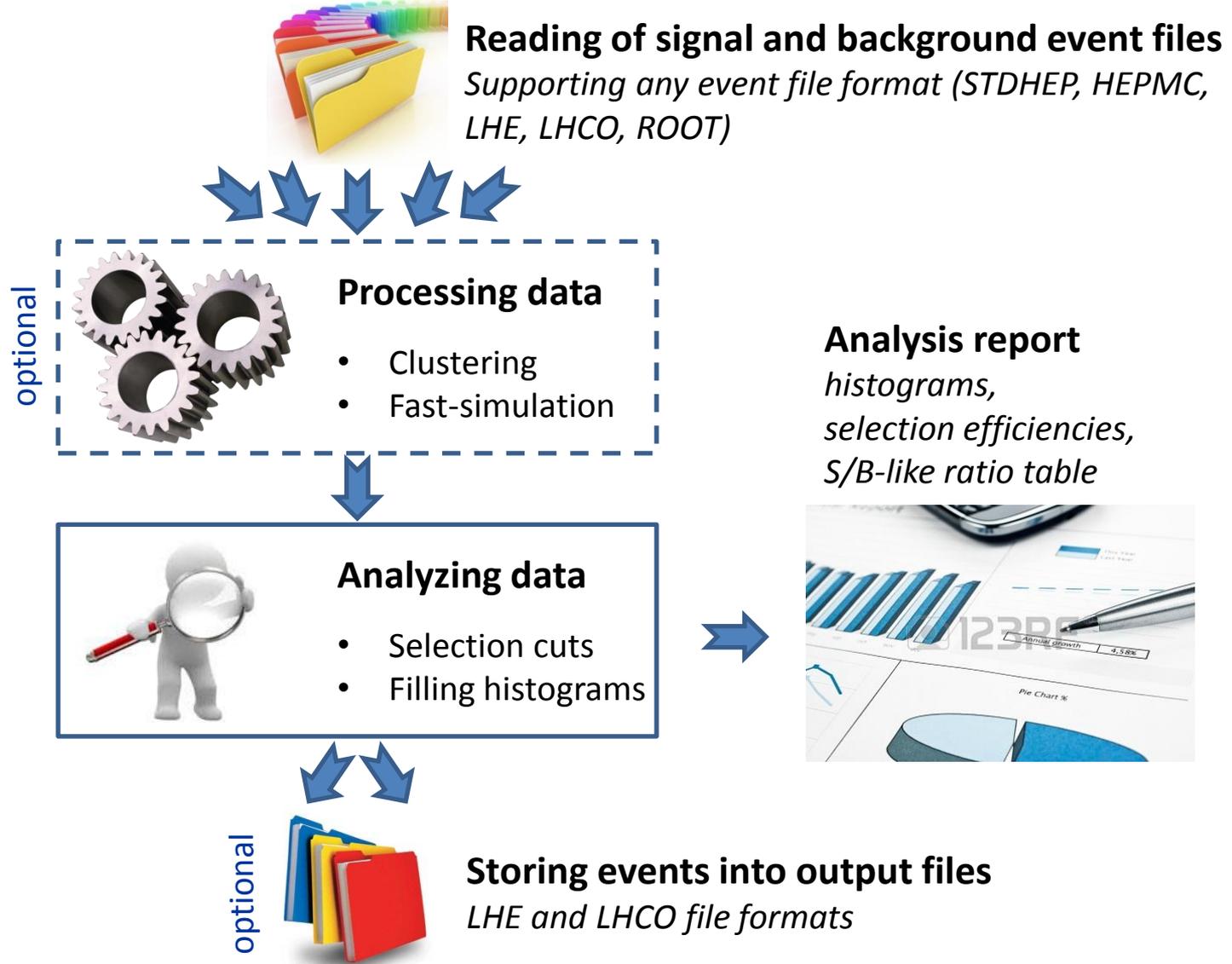
### Relevant features of design

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

**A unique framework : MadAnalysis 5**

## Scope:

Defining an analysis with a dedicated metalanguage



## 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
	<b>Delphes</b> 3.0

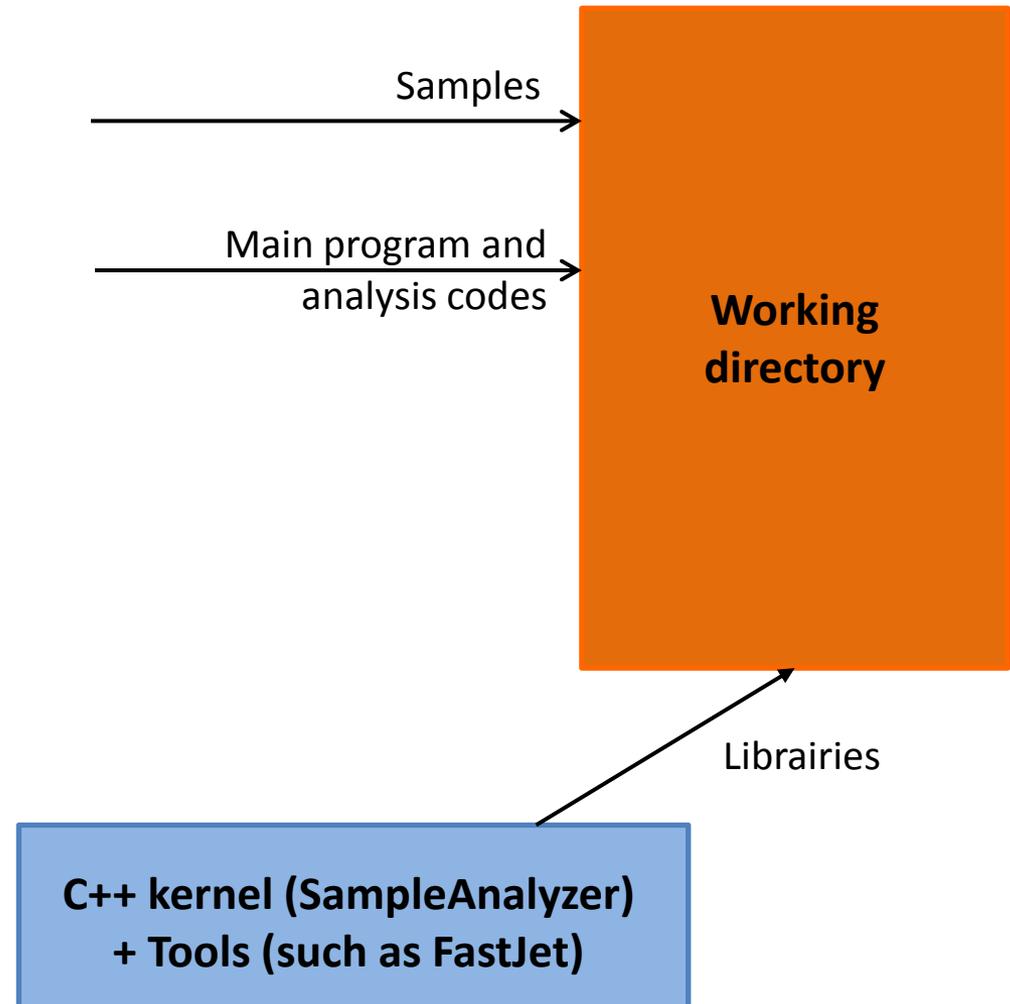
## Where MadAnalysis can be downloaded ?

- From the official website <https://launchpad.net/madanalysis5> (new address)
- From MadGraph 5 interface (available soon)



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

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



## Validation suite:

- Series of automated tests targeting to check all the functionalities of the program. Last ~ 3 hours.
- Launched for each compilation configuration of the kernel.
- One bug report → creation of a dedicated automated test.



## Documentation:

- **Manual** published in January 2013:  
[Comput. Phys. Commun. 184 \(2013\) 222](#)
- **First tutorials** are available on the website  
<https://madanalysis.irmp.ucl.ac.be/wiki/tutorials>  
→ Firsts steps with the Expert Mode

## 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 (if necessary) the C++ library.
- Step 3: Importing the list of particles and multiparticles  
(from MadGraph if this program is found on your system).

## 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 an analysis: 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 an analysis: 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 PT(mu+[1])
```

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

```
ma5> plot PT(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

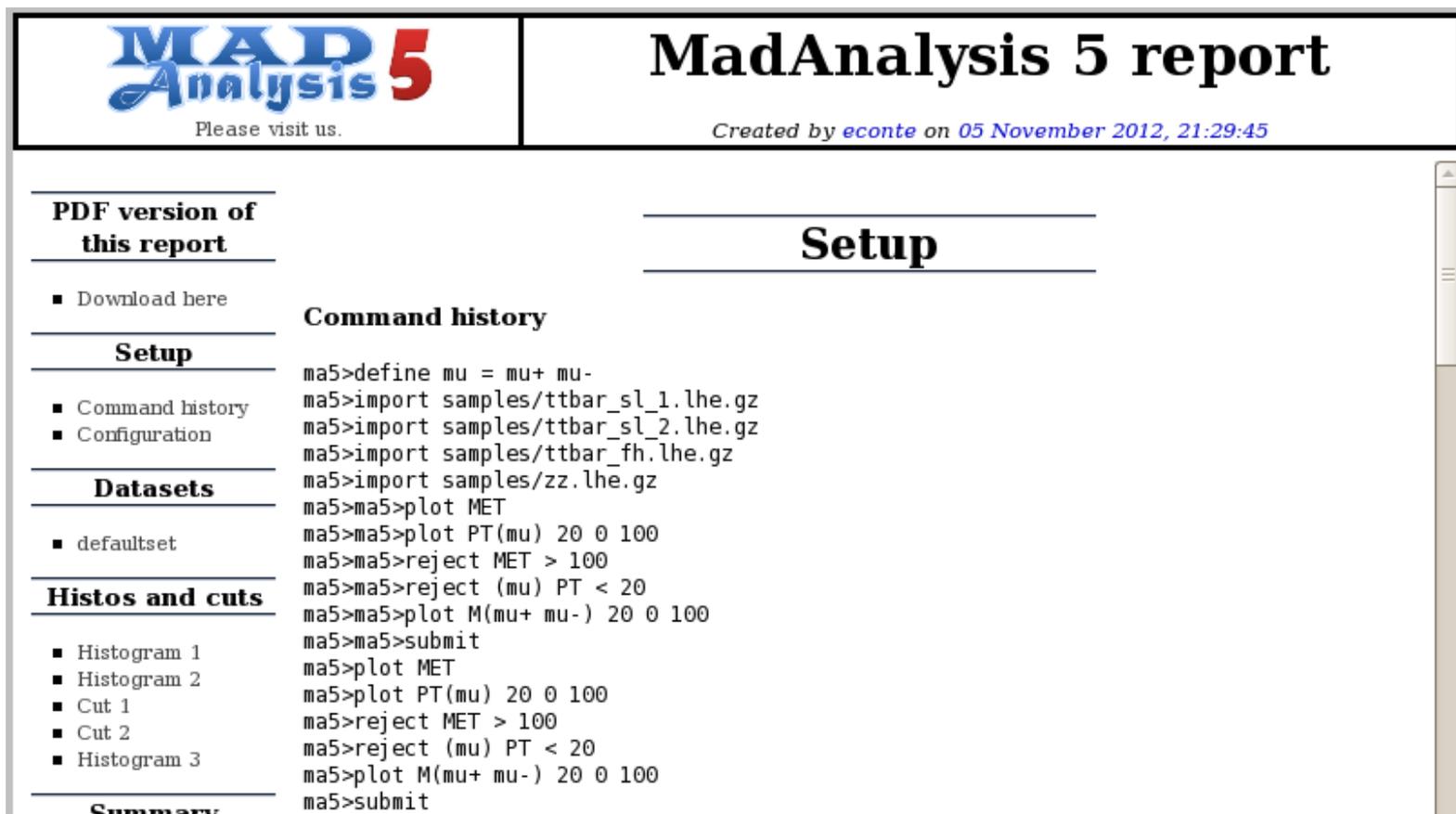
```
* SampleAnalyzer 2.0 for MadAnalysis 5 - Welcome.
* Option choices: selecting analysis = 'MadAnalysis5job'.
* Extracting the following sample files:
* 1/4 ~/samples/ttbar_sl_1.lhe.gz
  => file size : 107.09 Mo
  => sample produced by MadGraph.
  => progress [=====> ]
```

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.

Reports in PDF and DVI format are also available.



The screenshot shows the MadAnalysis 5 report interface. The top header contains the MAD Analysis 5 logo with the text "Please visit us." and the title "MadAnalysis 5 report" with the creation date "Created by econte on 05 November 2012, 21:29:45". The main content area is divided into a left sidebar and a main panel. The sidebar has sections for "PDF version of this report" (with a "Download here" link), "Setup" (with links for "Command history" and "Configuration"), "Datasets" (with a "defaultset" link), "Histos and cuts" (with links for "Histogram 1", "Histogram 2", "Cut 1", "Cut 2", and "Histogram 3"), and "Summary". The main panel is titled "Setup" and displays the "Command history" as a list of commands and their outputs:

```
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
```

## MadAnalysis 5 = multi-purpose interface

External static or shared libraries are linked to the built executable file.

Technical packages

- **ZLIB**: Reading and writing gzip-compressed files.

- **FASTJET**: M. Cacciari, G.P. Salam and G. Soyez, Eur.Phys.J. C72 (2012) 1896 [arXiv:1111.6097]
  - Generation of the ME/PS merging validation plots.
  - Applying an ideal detector simulation (jet clustering algorithm) to hadronic events.

Physics packages

- **DELPHES 3.0.10**: arXiv:1307.6346
  - Reading ROOT files.
  - Applying a realistic detector simulation to hadronic events.

Assisted installation



```
ma5> install fastjet
ma5> install delphes
ma5> install zlib
```

How to go from hadronic events to reconstructed events ?



**MadAnalysis 5 recipe  
based on FastJet**

**Delphes**

**Jet-clustering and identification algorithms  
+ efficiencies on (mis)identification**

- Selecting the fast-simulation package:

```
ma5> set main.fastsim.package =  
delphes          fastjet          none
```

## (Almost) ideal detector simulation based on FastJet

Thanks to  
A. Alloul for  
validation



- Selecting the fast-simulation package called: `fastjet`

```
ma5> set main.fastsim.package = fastjet
```

- Adopting a fast-simulation package → new options:

- **Large selection of jet clustering algorithms**

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

- **algorithm & object-identification** parameters. For instance:

```
ma5> set main.fastsim.algorithm = antikt  
ma5> set main.fastsim.ptmin     = 5  
ma5> set main.fastsim.radius    = 0.5  
  
ma5> set main.fastsim.exclusive = true  
ma5> set main.fastsim.bjet_id.  efficiency = 0.6  
ma5> set main.fastsim.bjet_id.  misid_cjet = 0.  
ma5> set main.fastsim.bjet_id.  misid_ljet = 0.
```

## Realistic detector simulation based on Delphes 3

- Selecting the fast-simulation package called: `delphes`

```
ma5> set main.fastsim.package = delphes
```

- Adopting a fast-simulation package → new options:
  - **Choice of the default Delphes cards**

```
ma5> set main.fastsim.detector =  
      cms atlas
```

- **Pile-up input:**

```
ma5> set main.fastsim.pileup = "MinBias.pileup"
```

- Once the analysis is submitted, MadAnalysis 5 invites the user to edit the configuration card with your favorite text editor (set the `EDITOR` shell variable).

```
ma5> Would you like to edit the configuration card used by  
Delphes ? yes/no [default = no]
```

- Delphes fast-simulation is launched event-by-event. No intermediate file is produced.

# Summary and perspectives



- **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.
- **Interface to physics-relevant tools:**
  - **Fast-Jet** for clustering and for generating ME/PS merging validation.
  - **Delphes 3** for fast-simulation.
  - Expected soon: shower programs.

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<https://launchpad.net/madanalysis5>  
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