MADANALYSIS 5

A new framework for collider phenomenology

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GDR Terascale @ Clermont-Ferrand

April 25, 2012.



Outline

- Introduction
- 2 Basic concepts
- 3 User-friendly mode
- 4 Expert mode
- Conclusion



- New framework to build complete analyses of Monte Carlo event files.
- Three levels of analysis:

```
partonic anique framework
reconstructed
```

April 25, 2012.

 Introduction
 Basic concepts
 User-friendly mode
 Expert mode
 Conclusion

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 MADANAINSIS 5 overview
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Collider Phenomenology

History

MADANALYSIS 4

- written in FORTRAN
- not very user-friendly
- check samples
- low flexibility

MADGRAPH5 MADANALYSIS 5

- written in PYTHON & C++
- very user-friendly
- uses ROOT
- allows to built sophisticated analysis

MADANALYSIS 5 is going beyond the scheme of the MAD software.

MADANALYSIS 5 is also a **standalone** package.



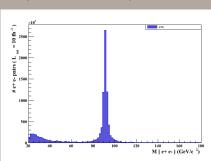
History

What can we do?

• Kinematical variables distributions (MADANALYSIS 4)

Example:

- Generated : p p > e+ e-
- Invariant mass of lepton pair

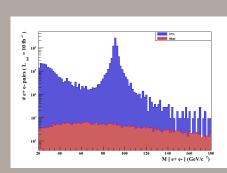


History

What can we do?

- Kinematical variables distributions (MADANALYSIS 4)
- Superposing and stacking

two different datasets

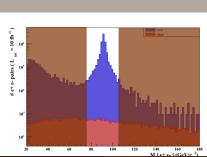


History

What can we do?

- Kinematical variables distributions (MADANALYSIS 4)
- Superposing and stacking
- Applying cuts

• Favored kinematical region



History

What can we do?

- Kinematical variables distributions (MADANALYSIS 4)
- Superposing and stacking
- Applying cuts
- Automated calculation of Signal/Background ratio.

●OOO MADANALYSIS 5 overview

Collider Phenomenology

History

Introduction

What can we do?

- Kinematical variables distributions (MADANALYSIS 4)
- Superposing and stacking
- Applying cuts
- Automated calculation of Signal/Background ratio.

What can we get?

 Results presented in an automatically generated and human readable report.





Wednesday, 25 April 2012 at 08:04:56.

Created by gserret.

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Please visit us.

Command history.
Configuration.
Datasets used..
e+e-(signal).
ttbar (signal).
Histograms / Cuts.
Histogram number 1.

Command history.

maS>inport /home/gserret/MorkingZone/madgraphS/PPEE/Events/run_01/unweighted_events.lhe.gz as e+emaS>inport /home/gserret/MorkingZone/madgraphS/ttbar_dilep/Events/run_01/unweighted_events.lhe.gz as ttbarmaS>pot M(e+ e-) 100 20 180
maS>set selection[1].logY = true
maS>sets relection[1].logY = true
maS>sets relection[1].logY = true

Configuration.



<u>File Edit View History Bookmarks Tools Help</u>	
MadAnalysis 5 HTML report	▼
☐ file:///home/gserret/WorkingZone/madanalysis/trunk/madanalysis-developmer ○ ▼ ⑥ ☑ file:///home/gserret/WorkingZone/madanalysis/trunk/madanalysis-developmer ○ ▼ ⑥	Q 😭
Personnel ▼ ☐Tools ▼ ☐MadAnalysis5 ▼ ☐RPV Pheno ▼ ☐FeynRules ▼ ☐Webmail ▼	

Configuration.

MadAnalysis version 0.5.117 (2012/04/24).

Histograms correspond to an integrated luminosity of 10 fb⁻¹.

Datasets used..

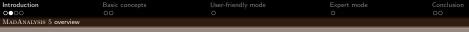
ttbar (signal).

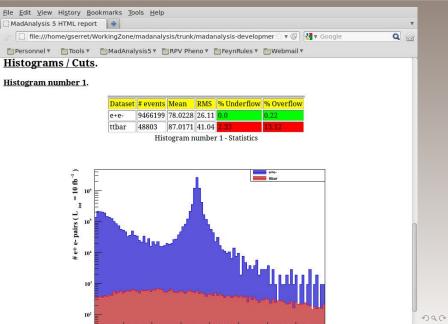
Event files	Number of events	Cross section (pb)
/home/gserret/WorkingZone/madgraph5/ttbar_dilep/Events/run_01 /unweighted_events.lhe.gz	10000	4.8803 +/- 0.0193
FINAL	10000	4.8803 +/- 0.0193

e+e- (signal).

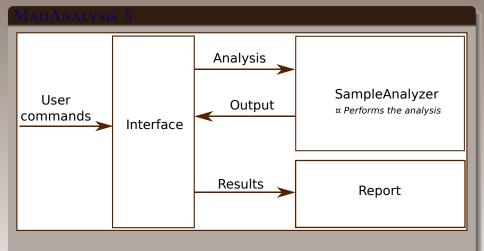
Event files	Number of events	Cross section (pb)
/home/gserret/WorkingZone/madgraph5/PPEE/Events/run_01 /unweighted_events.lhe.gz	10000	946.62 +/- 2.25
FINAL	10000	946.62 +/- 2.25

Histograms / Cuts.

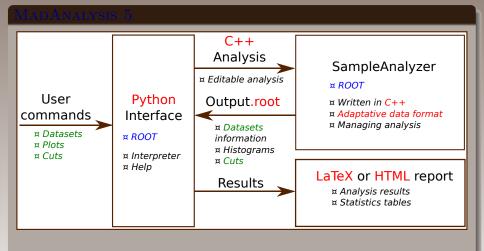




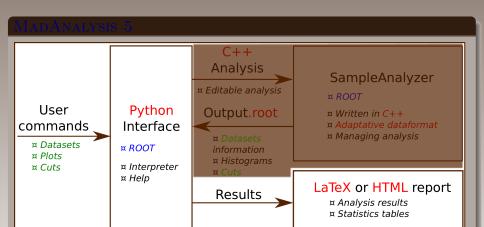
Introduction



Introduction

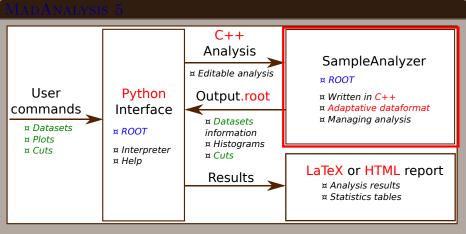


Introduction



• Traditional use: everything but the interface is transparent.

Introduction



- Traditional use: everything but the interface is transparent.
- Advanced use: the user is implementing his analysis in the SampleAnalyzer framework.



Interface

• PYTHON command line interface : ma5>



Interface

• PYTHON command line interface : ma5>

```
[aserret@sbpat588 madanalysis-development]$ ./bin/ma5
Checking ROOT libraries ...
Checking ROOT libraries ...
Checking 9+ libraries ...
Checking 12lb libraries ...
Checking 21lb libraries ...
Checking MadAnalysis library ...
System contiguration mas changed since the last use. Need to repulto the library
Creating (or overwriting) folder 'lib'...
Copying 'SampleAnalyzer' source files...
Creating a 'Makefile'...
Compiling the MadAnalysis library...
Linking the MadAnalysis library...
Checking the MadAnalysis library...
Checking the MadAnalysis library presence...
```

• Auto-completion + Interactive help : ma5>help <command>

$\operatorname{MadAnalysis}\ 5$ lies on simple concepts :

Datasets

Datasets

Regroup sample files that can be treated in the same way during the analysis, as a unique object refered with a label.

Suppose you have three sample files:

```
• Sample 1 : ttbar - 10 000 events
```

- Sample 2 : ttbar 20 000 events
- Sample 3 : Z' 10 000 events

You can

- gather Sample 1 & 2 in the ttbar dataset
- put Sample 3 in the zprime dataset
- ma5>import Sample1.lhe as ttbar
- ma5>import Sample2.lhe as ttbar
- ma5>import Sample3.lhe as zprime

Datasets

Particles/Multiparticles

Particles are identified with a PDG-Id : not very user-friendly.

In Madanalysis 5:

- particle object :
 - denoted with a label,
 - refers to PDG-Id. Example :
 - ma5>define mu+ = -13
- multiparticle object :
 - denoted with a label,
 - refers to several particles

Example:

• ma5>define mu = mu+ mu-

SUSY + SM particles automatically loaded with standard names in MADANALYSIS 5

Datasets

Particles/Multiparticles

Selection: plots & cuts

• plots : displaying information about



Datasets

Particles/Multiparticles

Selection: plots & cuts

- plots : displaying information about
 - global observables with respect to the event
 - missing transverse energy
 - particle content & multiplicity

ma5>plot MET



Datasets

Particles/Multiparticles

Selection: plots & cuts

- plots : displaying information about
 - non-global observables with respect to the particles:
 - kinematical information $(p_T, \theta, \phi, ...)$

ma5>plot PT(mu)

Introduction

MADANALYSIS 5 lies on simple concepts:

Datasets

Particles/Multiparticles

Selection: plots & cuts

- plots : displaying information about
 - non-global observables with respect to the particles:
 - kinematical information $(p_T, \theta, \phi, ...)$
 - multiplicity
 - ma5>plot N(mu)

Datasets

Particles/Multiparticles

Selection: plots & cuts

- plots : displaying information about
 - non-global observables with respect to the particles:
 - kinematical information $(p_T, \theta, \phi, ...)$
 - multiplicity
 - combining particles

•

ma5>plot M(mu+ mu-)

Datasets

Particles/Multiparticles

Selection: plots & cuts

- plots
- cuts : reject events/candidates not fulfiling a certain condition
 - reject event with MET < 50 GeV
 - reject mu candidate with PT > 50 GeV

ma5>reject MET < 50

ma5>select (mu) PT < 50</pre>

Datasets

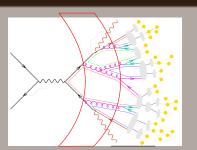
Particles/Multiparticles

Selection: plots & cuts

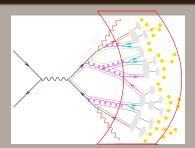
Report

Displaying results in a complete report written in LATEX or HTML

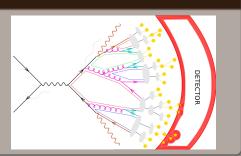
Partonic



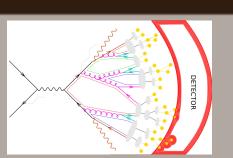
- Partonic
- Hadronic



- Partonic
- Hadronic
- Reconstructed



- Partonic
- Hadronic
- Reconstructed
- Supported files :
 - .lhe,.hepmc,.stdhep,.lhco.



Importing UFO model

Can directly import **particles information** from the model generated by FEYNRULES (through Universal Feynrules Output).

• automatic generation of labels

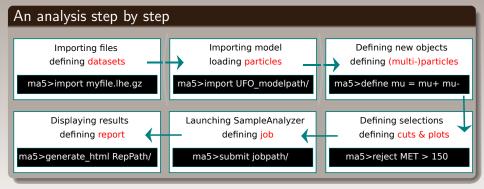


What the user want which concept(s) are involved
How to do it with MA5

buidling an analysis

What the user want which concept(s) are involved

How to do it with MA5

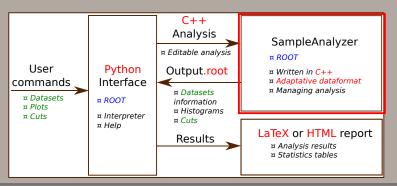


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Advanced analysis

- more sophisticated plots (2D, 3D, ...)
- new observables (asymmetry,...)

Even the expert mode is **developer-friendly**! Just needed to have **good skills** in programing.



Conclusion

Introduction

- New framework based on :
 - C++ kernel.
 - User-friendly PYTHON interface.
- Efficient way of performing an analysis :
 - reading event files,
 - defining analysis.
 - displaying results.
- Expert mode available :
 - open the door to the imagination of the users.
- Many more sophisticated functionalities are implemented ¹
 - particle ordering,
 - event history,
 - displaying options,
 - ...





Thanks for your attention

- Public version available soon.
- Webpage
 - https://server06.fynu.ucl.ac.be/projects/madanalysis/wiki/
- β -version distributed yesterday
- New β -testers are **welcome**, please ask!
- Remarks, suggestions, requests, anything :

ma5team@iphc.cnrs.fr

