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Outline.

- Quick overview.
- 2 Initialization and displaying information.
- Sevent files.
- 4 Performing an analysis.
- Current to-do list.

Scope.

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- * Analysis of event files produced by Monte Carlo generators.
- * Definition of various selection cuts on the input samples.
- * Production of histograms of different distributions.
- Programming languages used by MADANALYSIS.
 - * User interface written in Python and ROOT
 - * LHEANALYZER: core program written in C++, using ROOT, BOOST.
 - * PYTHON version: 2.6 and higher, no 3.x.
 - * ROOT version: 5.28 and higher.
- Additional features
 - * Output in ROOT, HTML, LATEX.
 - * To be used within MadGraph 5 or as a standalone package.
- Wiki page: https://server06.fynu.ucl.ac.be/projects/madanalysis
 - * Demo version related to this talk: http://server06.fynu.ucl.ac.be/downloads/madanalysis-current.tar.gz
 - Please send us your comments and suggestions (tickets on the wiki).

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- The program is far from being achieved.
 - * Please be patient.
 - * Quality before speed!
- Is your favorite feature missing?
 - * Ask.
 - * Wait (our to-do list is still huge).
 - * You will be advertised when it will be implemented.
- No manual presently.

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Initialization.

Download the tar-ball.

http://server06.fynu.ucl.ac.be/downloads/madanalysis-current.tar.gz

- Two different ways of usage.
 - * In the context of MadGraph.
 - ► Unpack the tar-ball within your MADGRAPH directory.
 - * As a standalone Python module.
 - ► Unpack the tar-ball anywhere.
- Please check that ROOT is installed.
 - * The directory root/bin must be included in the \$PATH variable.
 - \Rightarrow Then, users do not have to worry about ROOT anymore.
- Lauching the program.
 - ./bin/ma5

Initialization.



- Detect MadGraph or not.
- Default (multi)particle names.
 - * MG5: MADGRAPH defaults.
 - Alone: own defaults.
- The multiparticle hadronic.
 - * Related to jet energy. (variables such as H_T).
- The multiparticle invisible.
 - * Related to missing energy.

- Help available for any command (type help <command>).
- Command line completion ().
 - * Show available commands/options from what has been typed.
 - * Works with an empty command line.
 - * If a command has been typed \Rightarrow shows the available options.

```
▶help di 🔙
   ma5>help di
   display
                            display_datasets
   display_multiparticles
                            display_particles
   ma5>help display
►help display Enter
   ma5>help display
      Syntax: display <object name>
      Displays the definition and the properties of an object.
   ma5>
```

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```
ma5>display
    a
           cr~
                el-
                                   n2
                                          svm
                                                 ta-
                                                         u~
                                                              x1+
          c~
    b
                er+
                             1+
                                   n3
                                          svm~
                                                 ta1+
                                                         ve
                                                              x1-
    b1
           d
                er-
                             1-
                                   n4
                                          svt
                                                 ta1-
                                                         ve~
                                                              x2+
    b1~
           dl
                             main
                                          svt~
                                                 ta2+
                                                         vl
                                                              x2-
                g
                                   р
    b2
           dl~
                             mu+
                                          ຣ້
                                                 ta2-
                                                         vl~
                                                              z
                go
    b2~
                                                 t~
           dr
                h+
                                   s٦
                                          t.
                             mu-
                                                         vm
    h~
           dr~
                h-
                             m117+
                                   sl~
                                          t1
                                                         vm~
                                                 11
    С
           ď~
                h2
                                          t1~
                             mul-
                                   sr
                                                 ul
                                                         vt
    c1
           e+
                h3
                             mur+
                                   sr~
                                          t.2
                                                 ul~
                                                         vt.~
    cl~
                hadronic
                                          t2~
           e-
                             mur-
                                   sve
                                                 ur
                                                         w+
    cr
           el+
                invisible
                             n1
                                   sve~
                                          ta+
                                                 ur~
                                                         w-
    ma5>
```

- Help available for any command (type help <command>).
- Command line completion ().
 - * Show available commands/options from what has been typed.
 - * Works with an empty command line.
 - * If a command has been typed \Rightarrow shows the available options.

```
ma5>display main.
main.SBratio main.currentdir main.lumi
main.normalize main.uncertainty
ma5>
```

Displaying information.

- The display command (syntax: display <object_name>).
 - * Show the definition and the properties of an object.
- The display_particles and display_multiparticles commands.
 (syntax: display_particles and display_multiparticles).
 - * Show the list of all (pre)defined (multi)particles.
- The display_datasets command (syntax: display_datasets).
 - * Show the list of all imported datasets (i.e., all event files). [see below.]

Creating and destroying particles and multiparticles.

- The define command allows to define new (multi)particles.
 - * Example: collecting all muons (μ^+ and μ^-).
 - * Works as in MADGRAPH5.
- The remove command allows to remove existing (multi)particles.

```
▶define mu = mu+ mu- Enter
 display mu Enter
   ma5>define mu = mu+ mu-
   ma5>display mu
      The multiparticle 'mu' is defined by the PDG-ids -13 13 .
   ma5>
▶remove mu | Enter
 display mu Enter
   ma5>remove mu
   ma5>display mu
        ** ERROR: no object called 'mu' found.
   ma5>
```

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- This proceeds through the command import.
 - * PYTHON-like syntax.
 - * Both .Ihe and .Ihe.gz files are supported.
 - * Wildcards and \sim are allowed \Rightarrow import of several event files in one shot.
- Storage.
 - * Imported files are stored as datasets.
 - * Default set name: defaultset; otherwise specified by the user.
- ▶import demo/ttbar.lhe.gz Enter

ma5>import demo/ttbar.lhe.gz

-> Storing the file '/home/bfuks/Tools/madanalysis/trunk/madan alysis-development/demo/ttbar.lhe.gz' in the dataset 'defaultset'. ma5>

▶import demo/ww.lhe as ww Enter

ma5>import demo/ww.lhe.gz as ww

-> Storing the file '/home/bfuks/Tools/madanalysis/trunk/madan alysis-development/demo/ww.lhe' in the dataset 'ww'. ma5>

Importing event files.

- This proceeds through the command import.
 - * PYTHON-like syntax.
 - * Both .Ihe and .Ihe.gz files are supported.
 - * Wildcards and \sim are allowed \Rightarrow import of several event files in one shot.
- Storage.
 - * Imported files are stored as datasets.
 - * Default set name: defaultset ; otherwise specified by the user.
- ▶import demo/* Enter
 - ma5>import demo/*
 - -> Storing the file '/home/bfuks/Tools/madanalysis/trunk/madan alysis-development/demo/ww.lhe' in the dataset 'defaultset'.
 - -> Storing the file '/home/bfuks/Tools/madanalysis/trunk/madan alysis-development/demo/ttbar.lhe.gz' in the dataset 'defaultset'.
 - -> Storing the file '/home/bfuks/Tools/madanalysis/trunk/madan alysis-development/demo/dy.lhe.gz' in the dataset 'defaultset'. ma5>

- Two type of datasets: signal and background.
 - * To be developed in a not too far future.
 - In the prospects of comparing signal and background distributions.
 - * Default: the type of a dataset is signal-like.
 - * The type of an event sample can be modified through the command set.

► Setting the type of a dataset.

```
ma5>set defaultset.type = signal
ma5>set ww.type = background
ma5>
```

naying dataset information

- The command display_datasets.
 - * Gives the list of all imported datasets.
- The command display also works with datasets.
 - * Cross sections, number of events, line color in histograms, ...
 - * The properties can be changed through the command set.
- ► New MadAnalysis session: defining three datasets.

ma5>import demo/ttbar.lhe.gz as ttbar

- -> Storing the file '/home/bfuks/Tools/madanalysis/trunk/madan alysis-development/demo/ttbar.lhe.gz' in the dataset 'ttbar'. ma5>import demo/ww.lhe as ww
- -> Storing the file '/home/bfuks/Tools/madanalysis/trunk/madanalysis-development/demo/ww.lhe' in the dataset 'ww'. ma5>import demo/dy.lhe.gz as dy
- -> Storing the file '/home/bfuks/Tools/madanalysis/trunk/madan alysis-development/demo/dy.lhe.gz' in the dataset 'dy'. ma5>

- The command display_datasets.
 - * Gives the list of all imported datasets.
- The command display also works with datasets.
 - * Cross sections, number of events, line color in histograms, ...
 - * The properties can be changed through the command set.
- **▶** Displaying the list of imported datasets.

- The command display_datasets.
 - * Gives the list of all imported datasets.
- The command display also works with datasets.
 - * Cross sections, number of events, line color in histograms, ...
 - * The properties can be changed through the command set.
- ▶ Displaying information related to the (still not processed) www dataset. ma5>display ww ************** Name of the dataset = ww (signal). User-imposed cross section = 0.0. User-imposed weight of the set = 1.0. Line color in histograms = black. Background color in histograms = invisible. List of LHE files included in this dataset: - /home/bfuks/Tools/madanalysis/trunk/madanalysis-development/dem o/ww.lhe Cross section = (0.0 + /- 0.0) pb. Total number of events = 0. **************

- The properties of a dataset can be changed through the command set.
- A dataset can be removed through the command remove.

► Setting properties.

```
ma5>set ww.type = signal
ma5>set ttbar.backcolor = red
ma5>set dy.linecolor = blue
ma5>
```

- The properties of a dataset can be changed through the command set.
- A dataset can be removed through the command remove.

```
►remove ww Enter .
   ma5>display_datasets
      ****** List of defined datasets ******
      dy (signal)
     ttbar (signal)
      ww (signal)
      **************
   ma5>remove ww
   ma5>display_datasets
      ****** List of defined datasets ******
      dy (signal)
     ttbar (signal)
```

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- The command plot allows to define the distributions to be investigated.
 - * Requires as argument the observable to be analyzed.
 - * Requires also binning information (number of bins, min and max values).
 - * Future developments: automatic binning information from the event file.
- Cuts: being developed.
- Distributions without any argument (plot <obs> nbin xmin xmax).
 - * Total and missing transverse energy (TET and MET).
 - * Total and missing transverse hadronic energy (THT and MHT).
 - * Final particles present in the samples (NPID).
 - * Linked to the hadronic and invisible multiparticles.

Distributions with one or several argument(s).

```
(plot <obs>[arg1 arg2 ...] nbin xmin xmax).
```

- The arguments are (multi)particles.
 - \diamond plot PT[mu+]: the p_T of the antimuon.
 - \diamond plot PT[mu+ mu-]: the p_T of a muon anti-muon pair.
- * Multiplicities of the final particles present in the samples (N).
- * Kinematical variables (energy, mass, momentum, angles, rapidity, ...). ⇔ E, M, P, ET, MT, PT, PX, PY, PZ, RHO, THETA, PHI, ETA, Y, GAMMA, BETA.
- * Scalar and vector sums/differences/ratios. (sPT, vPT, dsPT, dvPT, rPT, . . .).
- * Remark: differences and ratios only defined for two particles.
- The or and all keywords can be used.
 - * Example-1: plot M[mu+ mu+ or mu- mu-] 201 0 200.
 - * Example-2: plot PT[all mu] 201 0 200 (several muons in the events).
 - * Remark: the keyword all is not fully implemented yet.

- The command display allows to show the analysis to be performed.
 - ► Displaying the list of imported datasets.

▶ Defining the analysis to be performed.

```
ma5>define mu = mu+ mu-
ma5>plot PT[mu] 501 0 500
ma5>plot N[mu] 6 0 5
ma5>plot MET 101 0 100
ma5>plot TET 501 0 500
ma5>
```

The command display allows to show the analysis to be performed.

```
▶ Displaying the analysis to be performed.
   ma5>display selection
      ****** Selection ******
      1. Plot : PT [ mu ]
      2. Plot : N [ mu ]
      3. Plot: MET
      4. Plot: TET
      **********
   ma5>
▶ Removing a selection.
   ma5>remove selection[4]
   ma5>display selection
      ******* Selection *******
      1. Plot : PT [ mu ]
      2. Plot: N [ mu ]
      3. Plot: MET
      ***********
   ma5>
```

Implementing an analysis.

 The command set allows to modify the properties of the histograms to be plotted.

Performing an analysis. 0000000

Performing the analysis.

- The command submit allows to perform the selected analysis.
 - * Reading of the lhe event files.
 - * Updating the information associated to each dataset.
 - * Creating a ROOT file with the analysis itself.
 - * Syntax: submit <dirname> (where the produced files are stored).

► Submitting the analysis.

```
ma5>submit testdir
  Creating folder 'testdir' ...
  Copying 'LHEanalyzer' source files...
  Inserting your selection into 'LHEanalyzer' ...
  Writing the list of datasets...
  Creating a 'Makefile' ...
  Compiling 'LHEanalyzer' ...
  Linking 'LHEanalyzer' ...
```

- The command submit allows to perform the selected analysis.
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 - * Reading of the lhe event files.
 - * Updating the information associated to each dataset.
 - * Creating a ROOT file with the analysis itself.
 - * Syntax: submit <dirname> (where the produced files are stored).

```
Checking LHEanalyzer output...
 Extracting data from the output files...
ma5>
```

- The command display_datasets.
 - * The information is now updated.

```
ma5>display ww
  *************
  Name of the dataset = ww (signal).
  User-imposed cross section = 0.0.
  User-imposed weight of the set = 1.0.
  Line color in histograms = black.
  Background color in histograms = invisible.
  List of LHE files included in this dataset:
   - .../demo/ww.lhe
  *************
  Cross section = 72.6 pb.
  Total number of events = 10000.
  *************
ma5>
```

- A ROOT-file is already there.
- Three additional output format can be generated.
 - * HTML: generate_html <dirname>.
 - * LATEX: generate_latex <dirname> or generate_pdflatex <dirname>.

Let's go for a live demo

Outline.

- Current to-do list.

Short term plans.

- * update command ('replot' in GNUPLOT).
- * preview/open command.
- * Hardcoded analyses (multiplicities, p_T , invariant masses,
- * The all keyword.
- * Sanity check routine of the LHE files.
- * p_T ordering of the particles (leading muon, ...).
- * Particle history (mother, grand-mother, ...).
- * Restoring Madanalysis configuration from the ROOT files.

Longer term plans.

- * Cuts and calculation of the efficiencies.
- * Automated treatment for the binning of the histograms.
- * Basic detector simulation.
- * Manual.