

HWWAnalysisCode

Tutorial Sessions

Session 1/2

Basic classes and data management



Andreas Walz
andreas.walz@cern.ch

Albert-Ludwigs-Universität Freiburg

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Where to find...

- ▶ The code and ROOT files used for this tutorial can be found in the `HWWAnalysisCode` repository (including the solutions to the exercises):

`HWWAnalysisCode/trunk/tutorial`

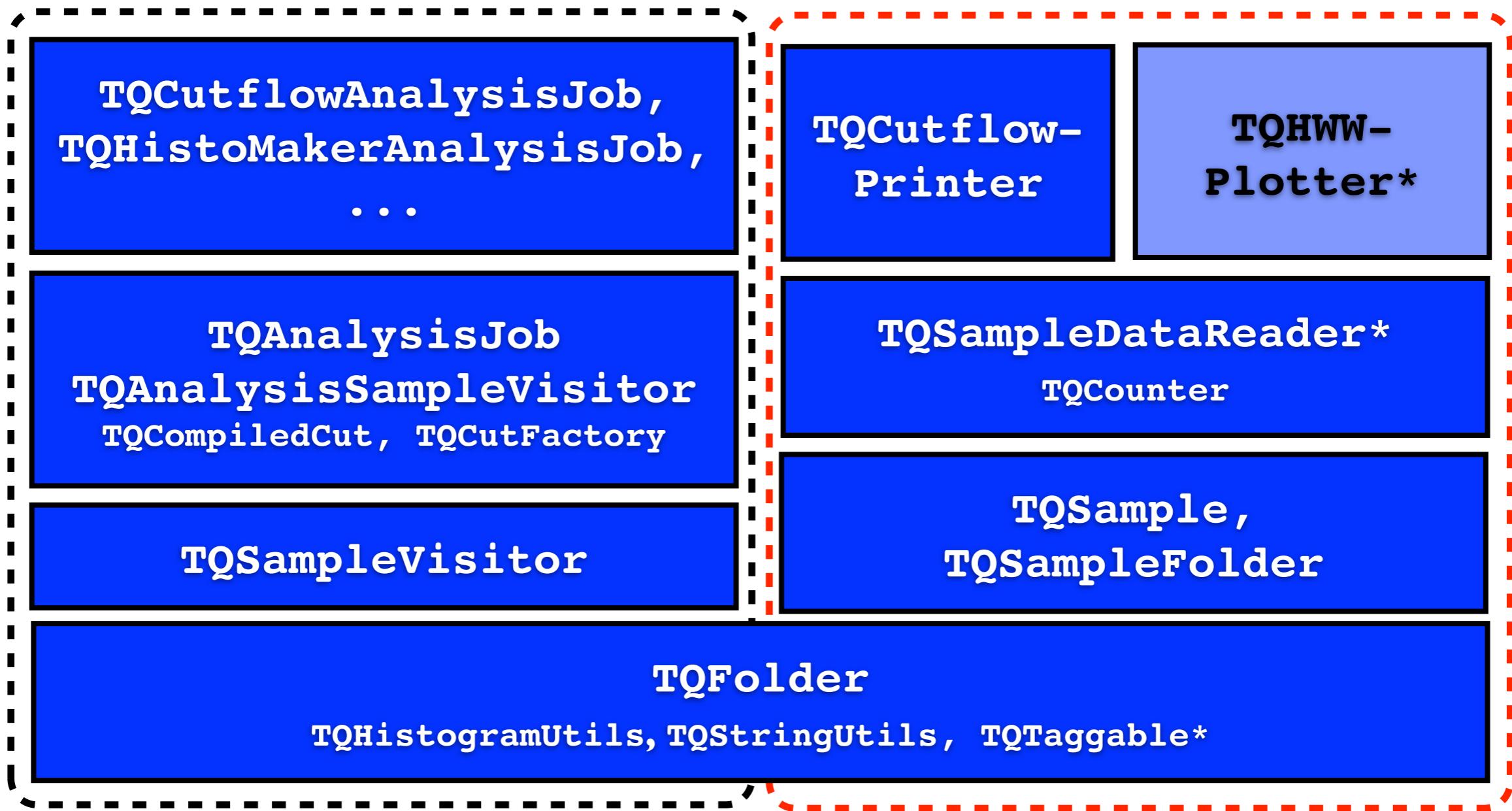
- ▶ A set of common ntuples (produced with `HWWNTupleCode -00-00-29`) to be used in some examples and exercises (in session 2) can be found at:

`/afs/cern.ch/work/a/awalz/public/ntuples`

Structure of the HWWAnalysisCode

Running analysis (producing analysis results, histograms, ...)

Management and presentation of analysis results (histograms, ...)



The TQFolder class

The TQFolder class is the basis for data management in the HWWAnalysisCode library. It is a **container** (similar to a directory in Linux) **for ROOT objects** (inheritors of **TObject**).

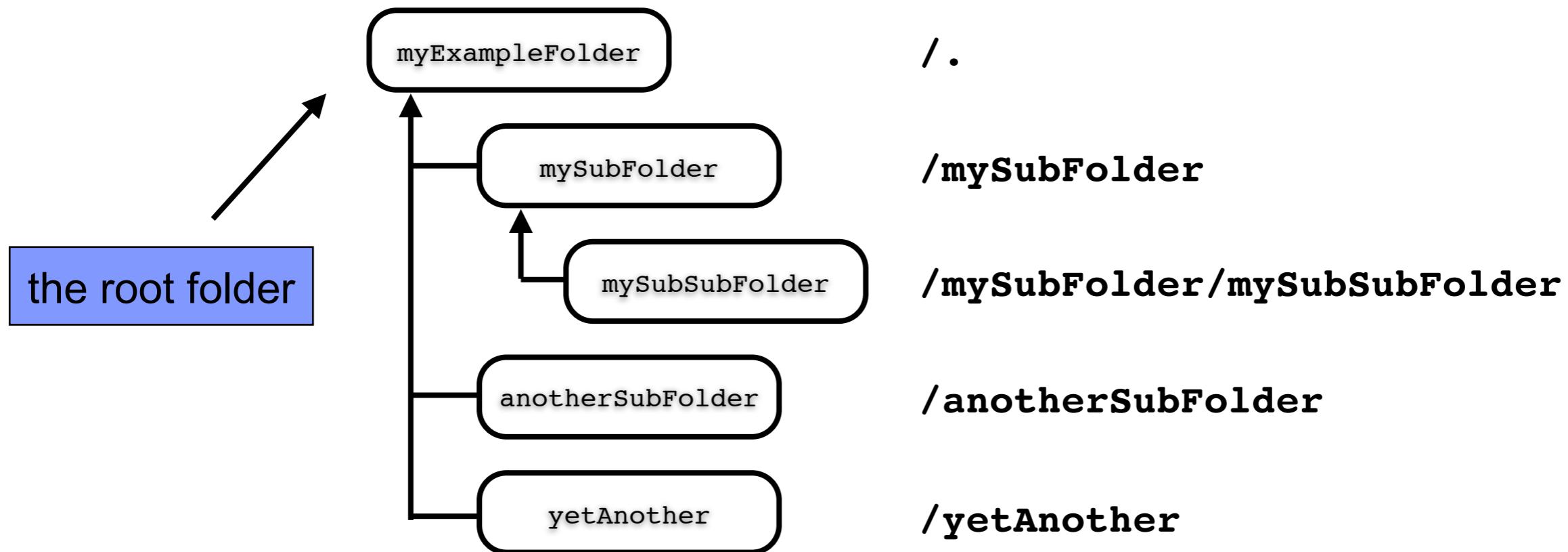
Some very useful/important functions:

- ▶ **folder->print();**
- ▶ print the contents of the folder (print-out can be controlled by parameters and flags)
- ▶ **folder->getFolder(„myFolder”);**
- ▶ return a pointer to a sub-folder

continued: The TQFolder class

The TQFolder class keeps a reference to its base folder (if existent) as well as to its sub-folders building a tree-like structure/hierarchy

- ▶ Writing the root folder to a ROOT file by using `folder->Write()` is sufficient to write the full hierarchy!



Example 1: TQFolder class

```
// always start in the ROOT command line loading the library  
root[0] .L ../lib/libQFramework.so
```



remember
this line

```
// load the example folder from the external ROOT file  
TQFolder * folder = TQFolder::loadFolder(  
    „example1.root:myExampleFolder”);
```

```
// print the contents of the folder  
folder->print();
```

```
// create a new sub-folder and print the contents again  
folder->getFolder(„myNewFolder+”);  
folder->print();
```



auto-create: creates the
folder if it doesn't exist

```
// create nested folders  
folder->getFolder(„newSub/newSubSub/foo+”);  
folder->print(„r”);
```

Example 2: TQFolder::print()

The print() method of the TQFolder class accepts several options to control the print-out:

// add a „details“ column to show additional information

```
folder->print(„d“);
```

use „rd“ in this example
to see the effect

// recursively print sub-folders (max. depth 2)

```
folder->print(„r2“);
```

can be any integer, no number
means unlimited depth

// use a filter (only show elements with matching name)

```
folder->print(„f[*LM*]“);
```

accepts wildcards „*“ and „?“,
usage like in Linux command

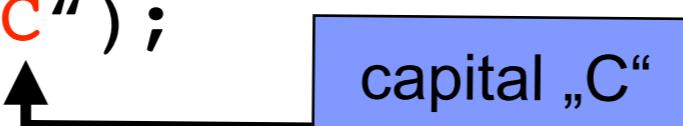
// combine options

```
folder->print(„df[*ALL*]r“);
```

continued: TQFolder::print()

```
// show the number of sub-elements  
folder->print( „c” );
```

```
// show the number of sub-elements (summed recursively)  
folder->print( „C” );
```



capital „C“

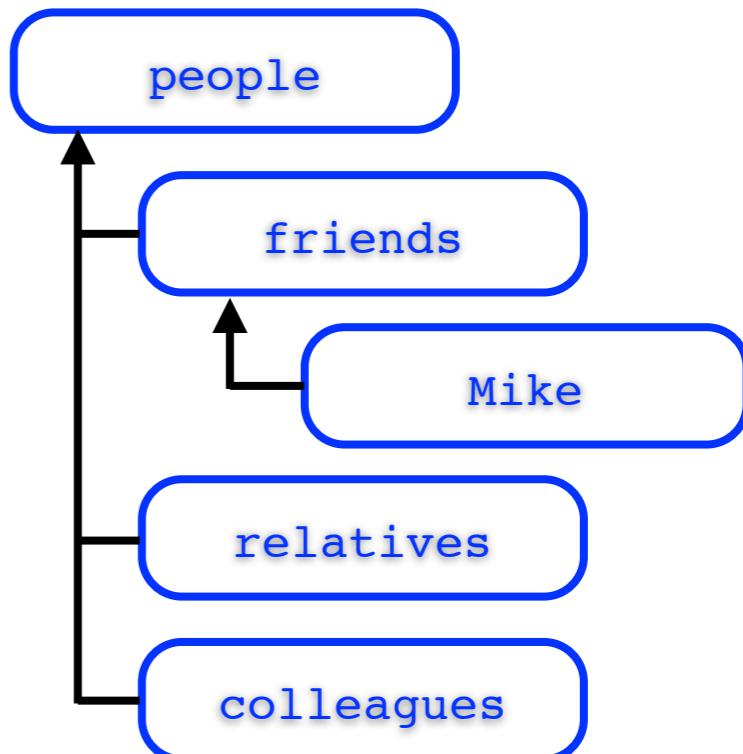
```
// print the contents of a sub-folder (here of „myNewFolder“)  
folder->print( „myNewFolder:” );
```

```
// ...combined with some options  
folder->print( „myNewFolder:dr2” );
```

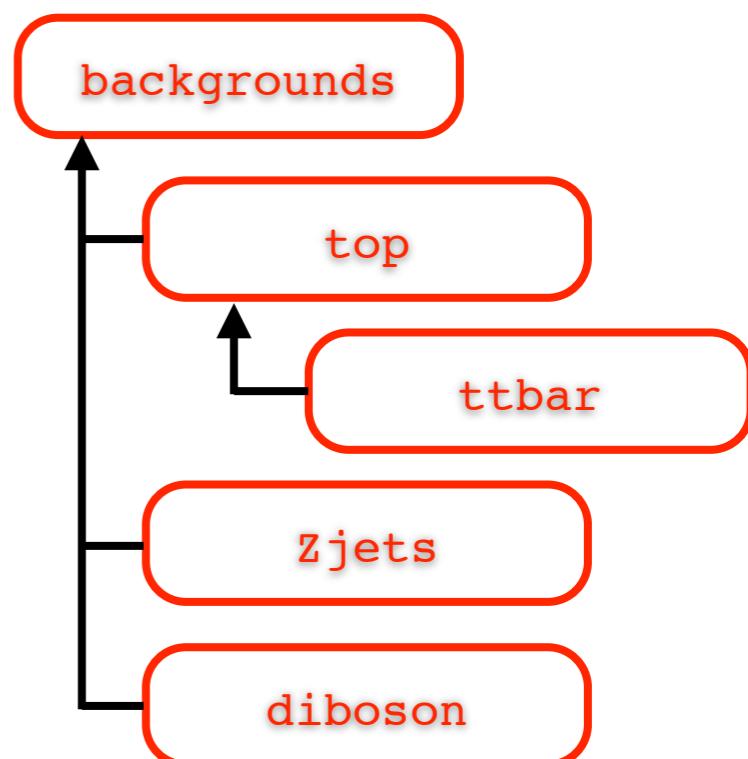
The `print()` method is the first tool to use if something does not work as it is expected !

The TQSampleFolder class

The **TQFolder** class is „only“ a container for objects derived from **TObject** and it **does not make any assumptions about the deeper meaning of its contents.**



The **TQSampleFolder** class is a specialization of the **TQFolder** class (inheriting all of its features) and it **is designed to represent a certain category of event samples.**



Example 3: TQSampleFolder class

There is a $H \rightarrow WW \rightarrow l\bar{l}l\bar{l}$ like example sample folder:

```
// load the example sample folder from the external ROOT file
TQSampleFolder * samples =
    TQSampleFolder::loadSampleFolder(
    „example3.root:samples“);

// print the contents of the sample folder
samples->print();

// explore the hierarchy, e.g.:
samples->print(„data:“);
samples->print(„data/ee:“);
samples->print("bkg/ee/diboson/ww/qqWW:d");

// print the full hierarchy
samples->print(„rH“);
```

hide system folders
starting with „..“

For ref: TQ(Sample)Folder classes

The `TQFolder` as well as the `TQSampleFolder` class provide several methods:

- ▶ **`folder->getListOfFolders(“name”)`**
 - ▶ return a `TList*` with a list of folders matching „name“
- ▶ **`folder->list(“name”)`**
 - ▶ print a list of folders matching „name“
- ▶ **`folder->getObject(“name”)`**
 - ▶ return a `TObject*` to the object matching „name“
- ▶ **`folder->deleteObject(“name”)`**
 - ▶ delete the object matching „name“
- ▶ **`sampleFolder->getListOfSampleFolders(“name”)`**
 - ▶ return a `TList*` with a list of sample folders matching „name“

Exercise 1: Locating a specific sample

Load the sample folder of the previous example and try to locate the sample with name „105200“ in the hierarchy. What is its path relative to the root folder?

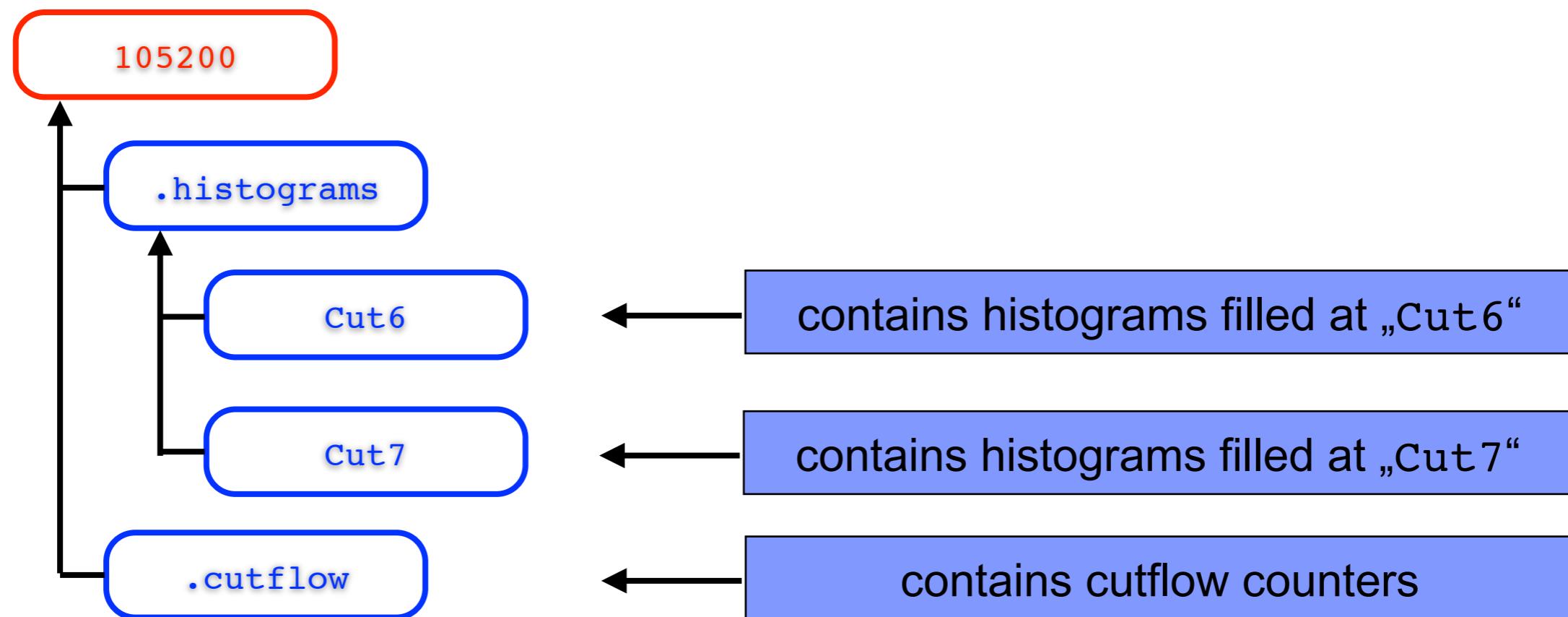
Some hints:

- ▶ use the `print()` method to locate the sample (there is more than one sample with this name)
- ▶ it is probably much faster doing this in the ROOT command line than writing a macro :-)

Analysis results in a TQSampleFolder

Reminder: the `TQSampleFolder` class is designed to represent a certain category of event samples.

Running an analysis means producing **histograms**, **cutflows**, **event lists**, ... („**analysis results**“) from event samples. These results are stored in the corresponding sample folder.



Example 4: Analysis results

```
// load the example sample folder from the external ROOT file
TQSsampleFolder * samples =
    TQSsampleFolder::loadSampleFolder(
„example3.root:samples”);

// print the contents of the sample folder representing the ttbar
sample (dataset ID 105200)
samples->print( „/bkg/ee/top/ttbar/105200:r1”);
samples->print( „/bkg/ee/top/ttbar/105200:r”);
```

Name	Class
.histograms/	TQFolder
Cut11/	TQFolder
Cut10/	TQFolder
Cut9/	TQFolder
Cut8/	TQFolder
Cut7/	TQFolder
Cut6/	TQFolder
.cutflow/	TQFolder
Cut11	TQCounter
Cut10	TQCounter
Cut9	TQCounter
Cut8	TQCounter
Cut7	TQCounter
Cut6	TQCounter

Cut6/	TQFolder
Mll	TH1F
Ptll	TH1F
DPhill	TH1F
m_jet_n	TH1F
MT	TH1F
METRel	TH1F
lepPt0	TH1F
lepPt1	TH1F
lepEta0	TH1F
lepEta1	TH1F

The TQSampleDataReader2 class

The TQSampleDataReader2 class provides powerful features to retrieve analysis results from a sample folder hierarchy:

- ▶ **reader->getHistogram(„path” , „histogram”);**
 - ▶ return a pointer to a histogram (TH1*) which is the sum of all histograms named „histogram” in „path”
- ▶ **reader->getCounter(„path” , „counter”);**
 - ▶ return a pointer to a cutflow counter (TQCounter*) which is the sum of all cutflow counters named „counter” in „path”

Example 5: TQSampleDataReader2

```
// load the example sample folder from the external ROOT file
TQSampleFolder * samples =
    TQSampleFolder::loadSampleFolder(
„example3.root:samples”);

// create an instance of the reader
TQSampleDataReader2 * rd =
    new TQSampleDataReader2(samples);

// make the reader printing error messages (not mandatory but
// sometimes helpful)
rd->setVerbose(1);

// retrieve and draw a histogram
TH1 * h = rd->getHistogram(„bkg”, „Cut7/Mll”);
h->Draw(„hist”);
```

continued: TQSampleDataReader2

```
// get histogram as sum of „bkg“ and „sig“
```

```
rd->getHistogram( „bkg + sig“, „Cut7/Mll“ );
```



```
// get histogram as sum of „Cut7“ and „Cut8“
```

```
rd->getHistogram( „bkg“, „Cut7/Mll + Cut8/Mll“ );
```

```
// use wildcards (returns the sum of matching folders)
```

```
rd->getHistogram( „bkg/?/top“, „Cut7/Mll“ );
```



„?“ matches any sample folder! In this example:

bkg/ee/top

bkg/em/top

bkg/mm/top

```
// return a normalized histogram (integral = 1.)
```

```
rd->getHistogram( „bkg“, „Cut7/Mll“, „norm=true“ );
```

Exercise 2: nJets of Z+jets samples

Load the sample folder of exercise 1 and plot the normalized nJets („m_jet_n“) distributions of different parton multiplicity ($Z \rightarrow \mu\mu$)+jets samples at „Cut6“ in one plot.

Some hints:

- ▶ use the `print()` method to locate the Z+jets samples.
Recall that option „d“ displays additional information
- ▶ concentrate on $\mu\mu$ generated and reconstructed final state,
e.g.

/bkg/mm/Zjets/Nom/z/mm/107660



reconstructed final state



generated final state

The TQCounter class

The TQCounter class is a representation of an event counter including **final event yields**, **uncertainties** and **raw event counts** (without any normalization or event weights applied):

- ▶ **counter->getCounter();**
- ▶ return the final event yield
- ▶ **counter->getError();**
- ▶ return the (statistical) uncertainty
- ▶ **counter->getRawCounter();**
- ▶ return the raw number of events (integer)

Example 6: Reading cutflows

```
// load the example sample folder from the external ROOT file
TQSampleFolder * samples =
    TQSampleFolder::loadSampleFolder(
        „example3.root:samples”);

// create an instance of the reader
TQSampleDataReader2 * rd =
    new TQSampleDataReader2(samples);

// retrieve the counter for total background at Cut7
TQCounter * c = rd->getCounter(„bkg”, „Cut7”);

// do whatever you like with the numbers
cout << „Total Background = ” << c->getCounter()
    << „ +/- ” << c->getError() << endl;
```

Exercise 3: Cut efficiencies

Load the sample folder of exercise 1 and calculate and print the cut efficiencies of „Cut11“ with respect to „Cut10“ for the total background and for the signal of a 125 GeV Higgs for ee, e μ and $\mu\mu$ final state, respectively.

Some hints:

- ▶ In the default $H \rightarrow WW \rightarrow l\nu l\nu$ structure, the path for different lepton flavour final states is
- ▶ total background: „bkg/ee“, „bkg/em“, „bkg/mm“
- ▶ 125 GeV Higgs signal: „sig/ee/mh125“, „sig/em/mh125“, „sig/mm/mh125“

Tags

You can associate tags (key-value-pairs) to a `TQFolder`, `TQSampleFolder`, ... object to store arbitrary additional information or control the behavior of classes accessing the folder hierarchy

► **`folder->setTagInteger(“mass”, 300);`**

or print the list of tags associated to the folder

► **`folder->printTags();`**

Key	Type	Value
datasetid	integer	105200
xsection	double	166.779999
kfactor	double	1.000000
filtereff	double	0.543000
priority	integer	1
generator	string	"MC@NLO"
processinfo	string	"ttbar(w/oFullHad)"
.init.filepath.alias	string	"MC11c"
.init.nevents	double	11584563.696994
.init.neventsbin	integer	1
~style.default.title	string	"t#bar{t}"
~style.default.histLineColor	integer	219
~usemcweights	bool	true

continued: TQSampleDataReader2

When reading histograms, the `TQSampleDataReader2` class reads specific tags associated to the sample folder (or its base sample folders) and applies corresponding style settings to the histogram to be returned:

- ▶ `style.default.histLineColor`
- ▶ `style.default.histFillColor`

- ▶ ...

„`style.`“ = common
style prefix

„`default`“ = style
scheme

style parameter

For reference: style tags

The style tags currently supported by the `TQSampleDataReader2` class:

- ▶ `(Integer) style.default.color`
- ▶ `(Integer) style.default.histLineColor`
- ▶ `(Integer) style.default.histLineStyle`
- ▶ `(Integer) style.default.histLineWidth`
- ▶ `(Integer) style.default.histFillColor`
- ▶ `(Integer) style.default.histFillStyle`
- ▶ `(Integer) style.default.histMarkerStyle`
- ▶ `(Double) style.default.histMarkerSize`
- ▶ `(Integer) style.default.histMarkerColor`
- ▶ `(String) style.default.title`

Exercise 4: Applying styles

Repeat exercise 2 but now using different line colors for distributions from different samples.

Some hints:

- ▶ a third argument of the `setTagInteger()` method can be used to specify the sub folder(s) to apply the tag to

```
folder->setTagInteger(  
    „style.default.histLineColor“, kRed,  
    „/bkg/mm/zjets/Nom/z/mm/107660“);
```

Generalizing histograms

By default histograms are filled and stored for every input sample individually potentially resulting in a very large data structure (you may remember the code crashing when trying to write such a large folder to a file).

In most of the cases it is sufficient to have histograms filled for a set of major sample categories.

The `TQSampleFolder` class provides a method which automatically „generalizes“ histograms of its sub sample folders to the level of the sample folder you call this method at:

- ▶ **sampleFolder->generalizeHistograms();**

Example 5: Generalizing histograms

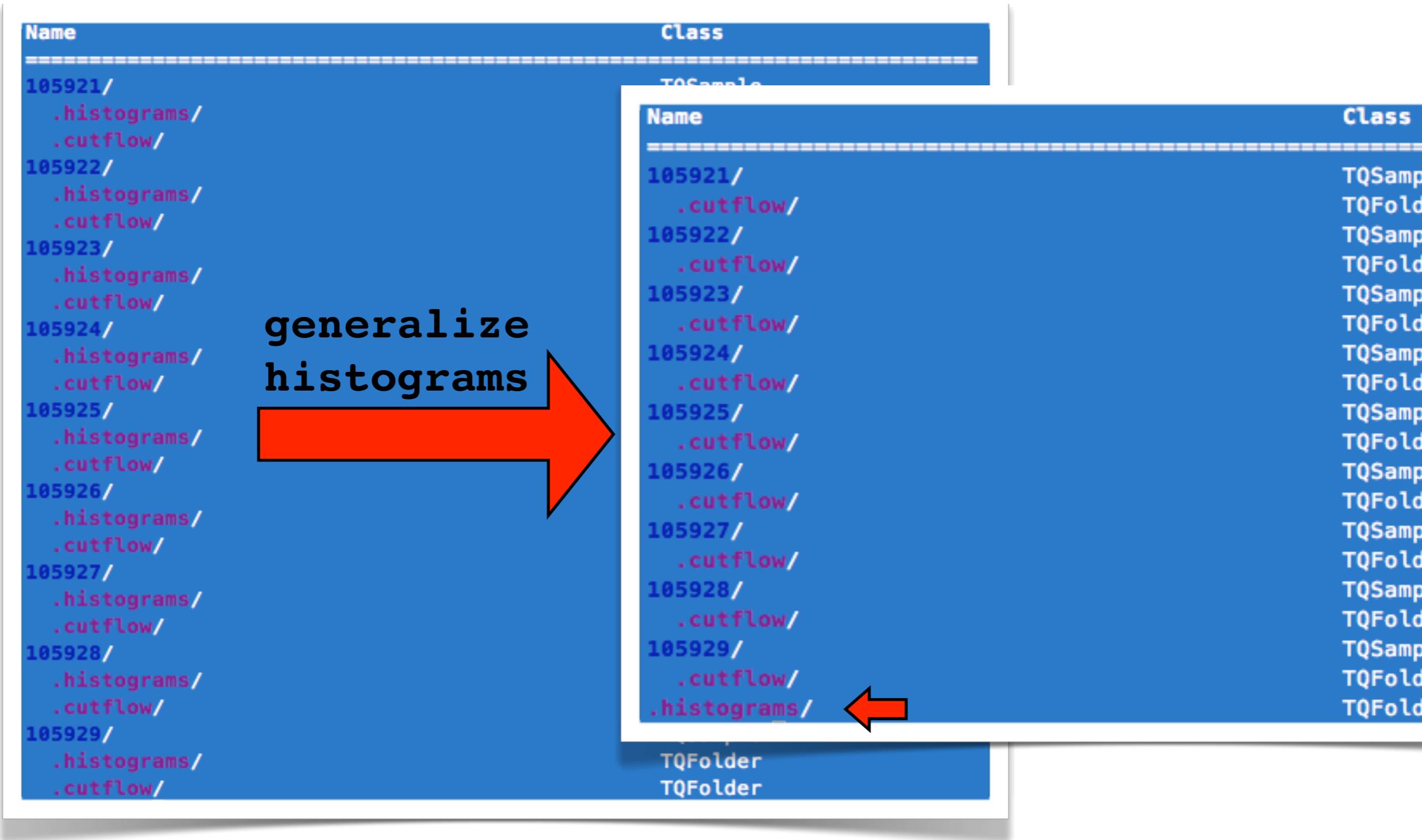
```
// load the example sample folder from the external ROOT file
TQSsampleFolder * samples =
    TQSsampleFolder::loadSampleFolder(
„example3.root:samples”);

// show individual histogram folders of qq->WW->evev samples
samples->print(„bkg/ee/diboson/WW/qqWW:rh”);

// generalize the histograms of qq->WW->evev samples
samples->getSampleFolder(
„bkg/ee/diboson/WW/qqWW”)
->generalizeHistograms( );

// see what happened to the individual histogram folders
samples->print(„bkg/ee/diboson/WW/qqWW:rh”);
```

Example 5: Generalizing histograms



Exercise 5: Generalizing histograms

Load the sample folder of a default $H \rightarrow WW \rightarrow l\bar{l}l\bar{l}$ analysis file and write it to a new file with reduced size by generalizing histograms to the level of the main background categories (W+jets, Z+jets, WW, WZ/ZZ/Z γ , ttbar, single top).

Some hints:

- ▶ use `hww_dataMC.root` as analysis input file
- ▶ each background category is present three times (for ee, e μ and $\mu\mu$ final state respectively)

For ref: $H \rightarrow WW \rightarrow l\nu l\nu$ sample structure

The default $H \rightarrow WW \rightarrow l\nu l\nu$ sample structure is given by

- ▶ bkg/
 - ▶ ee/ (em/, mm/)
 - ▶ diboson/
 - ▶ WW/
 - ▶ ...
 - ▶ NonWW/
 - ▶ ...
 - ▶ Zjets/
 - ▶ ...
 - ▶ Wjets/
 - ▶ ...
 - ▶ top/
 - ▶ ttbar/
 - ▶ singletop/
 - ▶ sig/
 - ▶ ee/ (em/, mm/)
 - ▶ mh125/ (mh110/, ..., mh600/)
 - ▶ ggf/
 - ▶ vbf/
 - ▶ WH/
 - ▶ ZH/
 - ▶ data/
 - ▶ ee/ (em/, mm/)

The TQHWWPlotter2 class

Producing plots in the $H \rightarrow WW \rightarrow l\nu l\nu$ style from a corresponding sample folder hierarchy can be as easy as two lines of code (the TQHWWPlotter2 class is specific to $H \rightarrow WW \rightarrow l\nu l\nu$)

```
▶ TQHWWPlotter2 * pl =  
    new TQHWWPlotter2(samples);
```

```
pl->plot(„Cut10_ALL/METRel”);
```

only $\mu\mu$ final state

```
pl->plot(„mm:Cut10_ALL/METrel”);
```

Set additional parameter to control the plotter (e.g. show a ratio plot of data/MC):

```
▶ pl->plot(„Cut10_ALL/METRel”,  
    „style.showRatio=true”);
```

Example 6: Plots in $H \rightarrow WW \rightarrow l\nu l\nu$ style

```
// load the sample folder of exercise 5
TQSsampleFolder * samples =
    TQSsampleFolder::loadSampleFolder(
        „hww_dataMC_genHisto.root:samples”);

// create an instance of the HWW plotter:
TQHWWPlotter2 * pl = new TQHWWPlotter2(samples);

// create and save a plot with data/MC ratio
TCanvas * c = pl->plot(„Cut10_ALL/METRel”,
    „style.showRatio = true”);
c->SaveAs(„MyHWWPlot.eps”);
```

For reference: TQHWWPlotter2

The parameter currently supported by the `TQHWWPlotter2` class:

- ▶ `input.mh`
- ▶ `style.logScale`
- ▶ `style.showRatio`
- ▶ `style.ratioMin`
- ▶ `style.ratioMax`
- ▶ `style.forceRatioLimits`
- ▶ `style.showUnderflow`
- ▶ `style.showOverflow`
- ▶ `labels.info`
- ▶ `labels.lumi`
- ▶ `labels.process`

The TQCutflowPrinter2 class

The TQCutflowPrinter2 class is an analysis independent class producing cutflow tables from corresponding sample folder hierarchies.

A cutflow table is matrix-like structure with different processes and cut stages in its rows and columns.

Construct a cutflow table using:

- ▶ **printer->addCut(„cut”, „title”);**
- ▶ **printer->addProcess(„path”, „title”);**
- ▶ **printer->createTable();**
- ▶ **printer->writeTable(...);**

Example 7: Cutflow tables

```
// load the sample folder of exercise 5
TQSampleFolder * samples =
    TQSampleFolder::loadSampleFolder(
        „hww_dataMC_genHisto.root:samples”);

// create an instance of the cutflow printer:
TQCutflowPrinter2 * pr = new
    TQCutflowPrinter2(samples);

// define the table
pr->addCut(„Cut10_ALL”, „Z veto”);
pr->addCut(„Cut11_ALL”, „METRel cut”);
pr->addProcess(„sig/ee/mh125”, „Signal”);
pr->addProcess(„bkg/ee/zjets”, „Z+jets”);
pr->addProcess(„bkg/ee”, „Total Background”);
pr->addProcess(„data/ee”, „Observed”);
```

continued: Example 7: Cutflow tables

...

// create the table

```
TList * table = pr->createTable(  
    „style.cellWidth = 30“);
```

// print the table

```
pr->writeTable(table);
```

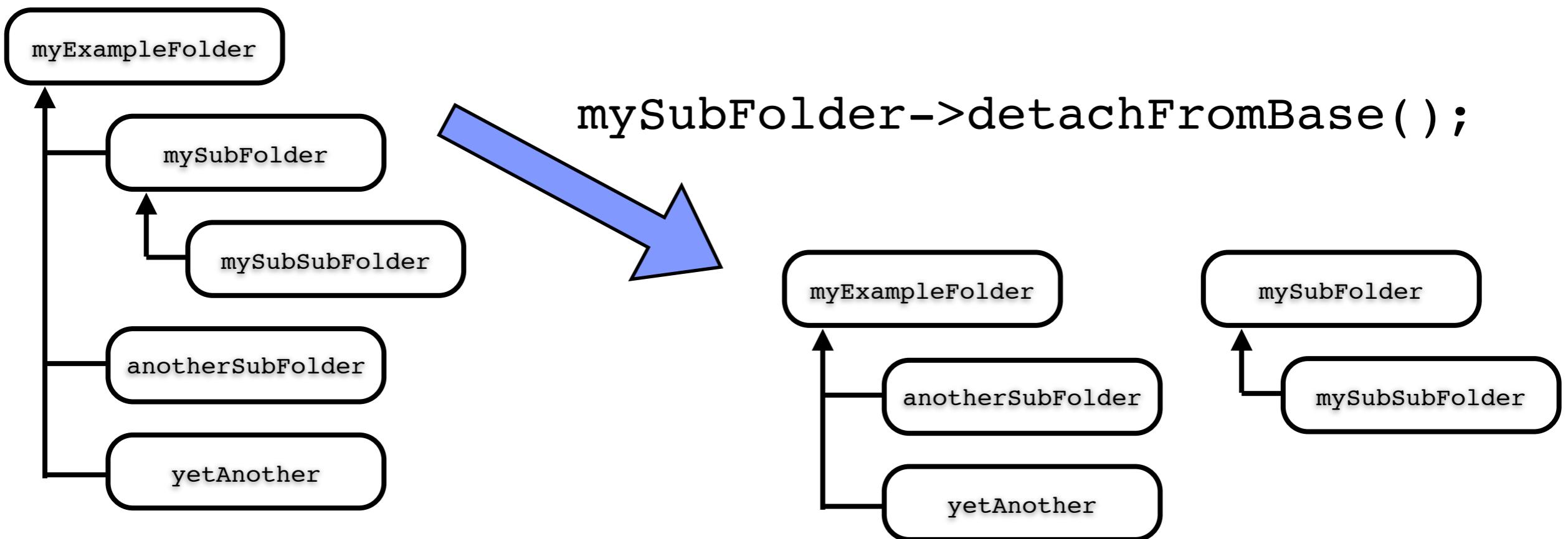
// write table to a file

```
pr->writeTable(table, „file.txt“);
```

Moving folders

Folders in a hierarchy may be deleted, moved, copied

- ▶ **detachFromBase()**
- ▶ **addFolder(...), addSampleFolder(...)**
- ▶ **copy()**



Example 8: Deleting folders

```
// load the example sample folder from the external ROOT file
TQSsampleFolder * samples =
    TQSsampleFolder::loadSampleFolder(
„example3.root:samples”);

// print the contents of the sample folder
samples->print();

// delete the signal sample folder
delete samples->getSampleFolder(„sig”)
->detachFromBase();

// print the contents of the sample folder again
samples->print();
```

Example 9: Moving folders

```
// load the example sample folder from the external ROOT file
TQSampleFolder * samples =
    TQSampleFolder::loadSampleFolder(
„example3.root:samples”);

// create a new folder to put the Wjets folder into
TQSampleFolder * dest = samples
    ->getSampleFolder(„newWjetsSampleFolder+”);

// detach the Wjets folder from its base...
TQSampleFolder * Wjets = (TQSampleFolder*)samples
    ->getSampleFolder(„bkg/ee/Wjets”)
    ->detachFromBase();

// ...and put it into its new destination folder
dest->addSampleFolder(Wjets);
```

Exercise 6: Moving folders

Load the sample folder written in exercise 5 and the sample folder containing the data-driven W+jets estimate and replace the Monte Carlo W+jets expectation by the data-driven one (for ee final state only).

Some hints:

- ▶ use `hww_ddWjets_genHist.root` as data-driven W+jets input file (the root sample folder in this file is named „Wjets“)
- ▶ the data-driven W+jets estimate for the ee final state is the sum of „data/ee“ and „bkg/ee“ in the data-driven W+jets sample folder, so you need to copy/move two folders
- ▶ please note: these two folders have the same name („ee“)

Additional exercises

Load the `hww_dataMC_genHist.root` file and produce a plot (using the `TQHWWPlotter2` class) of the transverse mass („MT“ or „MT_wide“) distribution in the 0 jet signal region („Cutt2_0jet_lowmass_SF_ALL“), but blinding the data.

Some hints:

- ▶ `samples->getListOfFolders(„data/.histograms/Cutt2_0jet_lowmass_SF_ALL“)` returns a `TList*` of every folder containing data histograms after cut Cutt2_0jet_lowmass_SF_ALL