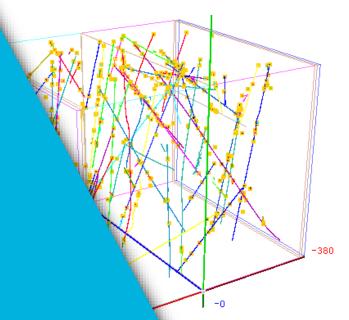
Running the reconstruction

(Exercise)

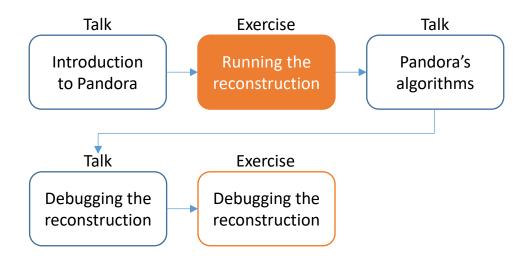
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10th UK LArTPC Software and Analysis Workshop



Reconstruction session



Credit: These slides are based on previous LArSoft workshop slides by Lorena Escudero and Andrew Smith-Jones

Key references:

Pandora ProtoDUNE paper
Pandora MicroBooNE paper

Goals

- This session is scheduled for 50 minutes.
- Main goal 1 Find and get to grips with the SBND reconstruction FHiCL files
 - Find the standard_reco1_sbnd.fcl and standard_reco2_sbnd.fcl configuration files
 - Look at the different reconstruction steps that we run
 - Understand what each of them do
- Main goal 2 Run the reconstruction
 - Run the reconstruction on the files we simulated yesterday
 - This includes running Pandora
 - Dump out the new output products to confirm we produced what we wanted

Before we get started...

- Later today we'll be running the event display
- Follow the steps from yesterday to get everything set up. To check, make sure MRB_TOP points to where you expect. Let us know if you have any issues here

\$ echo \$MRB_TOP # This should print the path to your development area

Main Goal 1

Understanding the SBND reconstruction FHiCL files

SBND reconstruction FHiCL files

- The full reconstruction is split into two fcls:
 - standard_reco1_sbnd.fcl
 - 2) standard_reco2_sbnd.fcl

which are run in succession

- reco1 generally corresponds to the 'lower-level' reconstruction e.g. signal processing, hit formation, etc...
- reco2 generally corresponds to the 'higher-level' reconstruction e.g. particle creation
- Let's take a closer look at the fcls to see what these stages in more detail

reco1

Find the location of standard_reco1_sbnd.fcl using SBND's handy fhicl finder script

```
$ find_fhicl.sh standard_reco1_sbnd.fcl
```

Didn't work? Make sure you've setup your working area!

Open the file with your favourite code editor

\$ vim /path/from/command/above

Wow, vim is your favourite? Great choice!

To exit vim do Esc :qa Return

To exit emacs do Ctrl-x Ctrl-C

- Find the trigger_paths: [...] block
- You'll learn more in the analysis tutorial, but this block tells us the names of the 'producer' modules that will add products to our .root file
- You'll find the trigger_paths block filled by the name 'reco1'
- You'll find 'reco1' defined as @local::sbnd_reco1_producer_sequence, which itself is defined in workflow_reco1.fcl
- Repeat the above steps to search and view the reco1 producer sequence in workflow_reco1.fcl

Following a seemingly never-ending series of fhicl files to find the thing you want is pretty standard - keep going, you'll get there eventually!

```
sbnd_reco1_producer_sequence: [
    rns
    , opdecopmt
    , opdecoxarapuca
    , ophitpmt
    , ophitxarapuca
    , opflashtpc0
    , opflashtpc1
    , opflashtpc1xarapuca
    , opflashtpc1xarapuca
    , conflashttc1, conflashtt2, conflashttc1, conflashtt1, conflashtt2, conflash
```

reco1 – what are these modules?

```
sbnd_reco1_producer_sequence: [
                                            random number saver
  rns
, opdecopmt
, opdecoxarapuca
, ophitpmt
, ophitxarapuca
                                            photon detection system reco
, opflashtpc0
, opflashtpc1
, opflashtpc0xarapuca
, opflashtpc1xarapuca
                                            TPC hit formation
, gaushit
, gaushitTruthMatch
                                            identify to which MC particle each hit belongs
, crtstrips
                                            cosmic ray tagger (CRT) reco
                                            machine learning (ML) reco inputs
 , cluster3d
```

reco2

Follow the same procedure to examine standard_reco2_sbnd.fcl

```
$ find_fhicl.sh standard_reco2_sbnd.fcl
$ vim /path/from/command/above
```

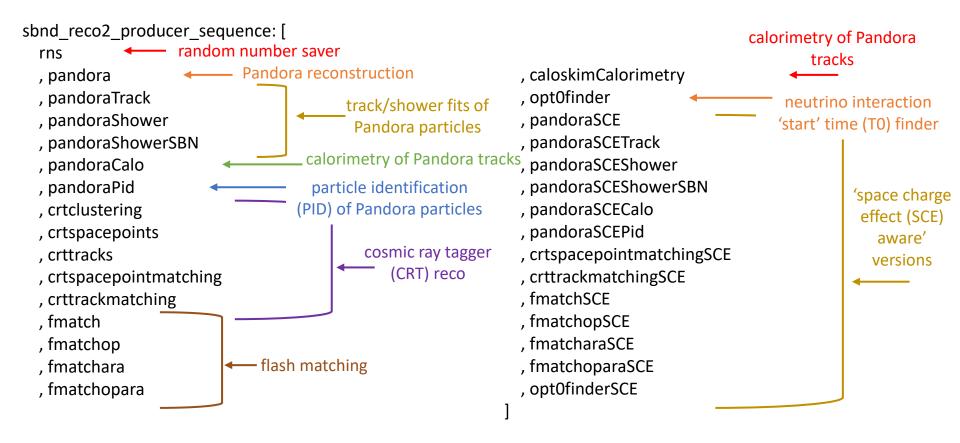
Hunting through fcls is something you'll be doing time and time again, and one day they will suddenly make sense

- Find the trigger paths: [...] block
- You'll find the trigger_paths block filled by the name 'reco2'
- You'll find 'reco2' defined as @local::sbnd_reco2_producer_sequence, which itself is defined in workflow_reco2.fcl
- Repeat the above steps to search and view the reco2 producer sequence in workflow_reco2.fcl

In the last session, we were introduced to pandora, but there are many other steps in the reconstruction chain too!

```
sbnd reco2 producer sequence: [
  rns
                                                     , caloskimCalorimetry
  , pandora
  , pandoraTrack
                                                     , opt0finder
  . pandoraShower
                                                     . pandoraSCE
  , pandoraShowerSBN
                                                     , pandoraSCETrack
  . pandoraCalo
                                                     , pandoraSCEShower
  , pandoraPid
                                                     , pandoraSCEShowerSBN
  , crtclustering
                                                     , pandoraSCECalo
  , crtspacepoints
                                                     . pandoraSCEPid
  . crttracks
                                                     , crtspacepointmatchingSCE
                                                     , crttrackmatchingSCE
  , crtspacepointmatching
                                                     , fmatchSCE
  , crttrackmatching
                                                     , fmatchopSCE
  . fmatch
  . fmatchop
                                                     . fmatcharaSCE
  . fmatchara
                                                     , fmatchoparaSCE
                                                     , opt0finderSCE
  , fmatchopara
```

reco2 – what are these modules?

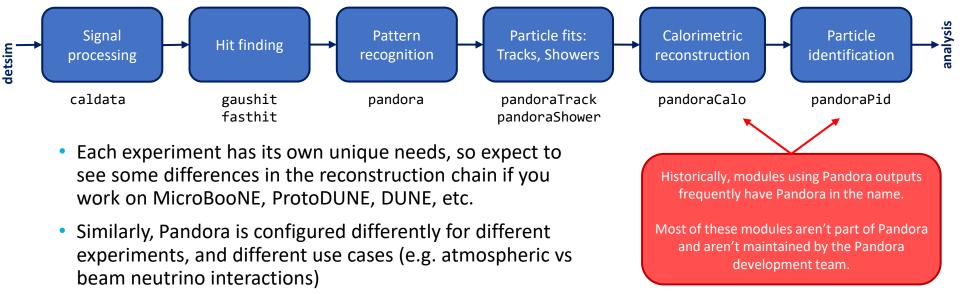


A cumulative reconstruction output

There's a lot happening in the reconstruction stage

Next, let's see how Pandora is configured for SBND

• To demonstrate how these modules accumulate a reconstruction output, let's focus on the Pandora 'workflow' i.e. ignoring the optical and CRT reconstruction

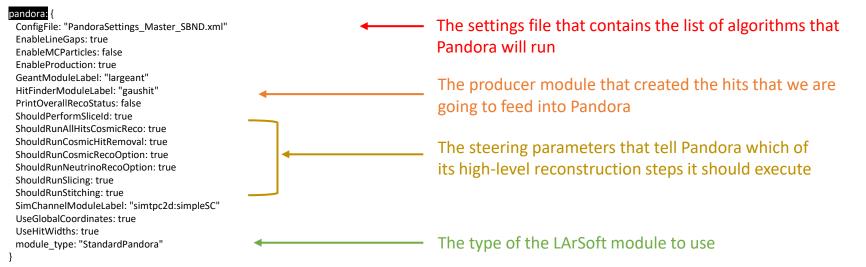


Pandora's configuration

- Practically all LArSoft modules have a configuration defined by parameters e.g. say that we had a producer module that added a number to our root file it would likely have a parameter which defines the number that is added, and that parameter is set in our fcl files
- fcl files often #include many other fcl files, and it can be unclear where our parameters are set
- fhicl-dump answers this question, and follows all #includes to get the bottom-line configuration
- We can pipe (|) its output to less and search for a producer to learn more:

\$ fhicl-dump standard_reco2_sbnd.fcl | less -p "pandora:"

Less's –p option allows us to jump straight to the part of the file that we are interested in. In less, use the ↑/↓ keys to move up and down, and 'q' to exist.



Main Goal 2

Running the reconstruction

Running the reconstruction

• We are now poised to run the reconstruction! Make a directory to work in, and run it:

```
$ mkdir -p $MRB_TOP/reco/work
$ cd $MRB_TOP/reco/work
$ lar -c standard_reco1_sbnd.fcl -n -1 -s /path/to/my/detsim/file.root -o reco1_tutorial.root
$ lar -c standard_reco2_sbnd.fcl -n -1 -s reco1_tutorial.root -o reco2_tutorial.root
```

Can also run on pre-made gen+g4+detsim files in:
/scratch/LAR25/TPCSimulation/detsim tutorial.root

reco1 can take some time to run

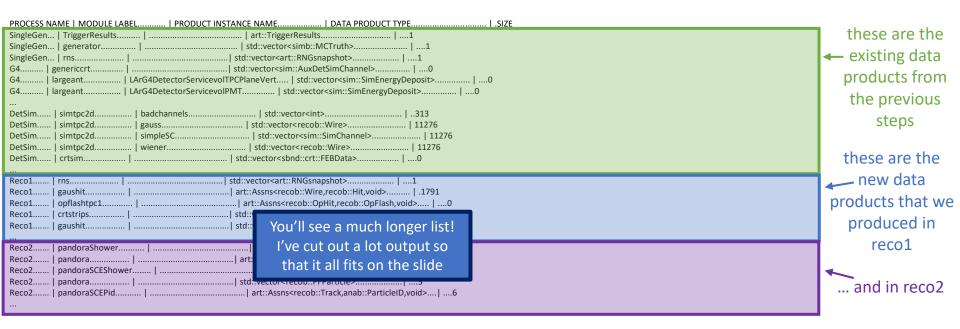
```
TimeTracker printout (sec)
                               0.515653
source:RootInput(read)
                                    0.000233144 0.00842777 0.0686421 0.000264929 0.0204246
reco2:rns:RandomNumberSaver
                                         2.0329e-05 4.86311e-05 0.000269584 2.2662e-05 7.37634e-05
reco2:pandora:StandardPandora
reco2:pandoraTrack:LArPandoraTrackCreation
                                             0.00274014 0.00409558 0.00646157 0.00403605
reco2:pandoraShower:LArPandoraModularShowerCreation
reco2:pandoraShowerSBN:LArPandoraModularShowerCreation
                                                                           0.00280114
reco2:pandoraCalo:GnocchiCalorimetry
                                           0.00202068 0.00246629 0.00407539 0.00228423 0.000550001
reco2:pandoraPid:Chi2ParticleID
                                       0.00011043 0.000201841 0.000876067 0.000128414 0.000225004
reco2:pandoraSCEPid:Chi2ParticleID
                                        end path:out1:RootOutput
                                       2.194e-06 3.3834e-06 1.1412e-05 2.4845e-06
end_path:out1:RootOutput(write
```

We can check to see that everything we expected has been executed, and see how long each took

So... what's new?

Run eventdump.fcl on the .root file to see the new collections we've just made

\$ lar -c eventdump.fcl -s reco2_tutorial.root -n 1



Additional information

Configuring Pandora steps

(For reference)

- Pandora's full reconstruction workflow is designed to handle neutrino interactions in dense cosmic ray environments
- You'll see in the upcoming lecture how we've designed algorithms, and entire algorithm chains to target specific topologies and have:
 - a cosmic-ray optimised algorithm chain
 - a beam neutrino interaction optimised algorithm chain
 - an atmospheric neutrino interaction optimised algorithm chain
 - a test-beam optimised algorithm chain (for use at ProtoDUNE)
 - a low energy neutrino interaction optimised reconstruction chain (for e.g. supernova neutrinos at the DUNE FD)
- SBND is a surface detector in a neutrino beam, so sees both neutrino and cosmic ray interactions
- We want to run the cosmic algorithm chain over cosmic interactions, and the neutrino algorithm chain over neutrino
 interactions ⇒ we'll want to run the so called 'consolidated reconstruction' chain
- We can configure Pandora to run one, many, or all steps by modifying its steering parameters
- Let's have a go at changing how Pandora runs!

Pandora Config

(For reference)

Firstly, what steering parameters do we have and what do they mean?

should we initially run a fast and simplified version of the reconstruction? ShouldRunAllHitsCosmicReco and do we need to 'stitch' this reconstruction output between different detector chambers? ShouldRunStitching should we try to remove 'obvious' cosmic rays from this output e.g. tracks that both enter and leave the detector? ShouldRunCosmicHitRemoval do we expect multiple cosmic rays? or multiple neutrinos? or both? ShouldRunSlicing And therefore, do we need to group (or slice) the input hits such that each individual interaction exists within it's own slice? ShouldRunCosmicRecoOption do we expect cosmic-rays? and so should we run the cosmic algorithm ShouldRunNeutrinoRecoOption chain over these slices? ShouldPerformSliceId do we expect neutrinos? and so should we run the neutrino algorithm chain over these slices?

do we need to determine whether a slice contains a cosmic ray or neutrino interaction?

Let's do it! (For reference)

- Make a new directory to work in during this session
- Then create a new FHiCL file with the lines below
- Then save and close the file

```
$ mkdir -p $MRB_TOP/reco/config
$ cd $MRB_TOP/reco/config # Put your new .fcl file here
$ vim my_reco_sbnd_basic.fcl
```

Please use your favourite text editor, here we use vim. If you accidentally opened vim and want to close it type Esc, :qa, Return &

include the standard configuration

#include "standard_reco2_sbnd.fcl"

physics.producers.pandora.ShouldRunAllHitsCosmicReco: false physics.producers.pandora.ShouldRunStitching: false physics.producers.pandora.ShouldRunCosmicHitRemoval: false physics.producers.pandora.ShouldRunSlicing: false physics.producers.pandora.ShouldRunCosmicRecoOption: false physics.producers.pandora.ShouldRunNeutrinoRecoOption: true physics.producers.pandora.ShouldPerformSliceId: false

With these values of the steering parameters:

- we'll treat the input hits as if they belong to a single interaction
- and will reconstruct them under the neutrino algorithm chain
- performing no stitching across the central gap in SBND – JINKIES!

Pointing to a new configuration

(For reference)

• We'll need to make sure that LArSoft will know where to look for our new FHiCL file, to do this we add it to the FHICL FILE PATH environment variable. Start by printing it to the terminal:

```
$ echo $FHICL FILE PATH
```

- You will see many many directories, all separated by a ':'.
- To add our reco/config folder to this list run the following command:

```
$ export FHICL FILE PATH=$MRB TOP/reco/config:$FHICL FILE PATH
```

- echo the FHICL FILE PATH again to check that everything worked (it should be the first in the list)
- Now run fhicl-dump again to make sure our new configuration file is set up as we want

```
$ fhicl-dump my reco sbnd basic.fcl | less -p "pandora:"
```

• Now try to run reco2 (on your reco1 output) with your new fcl file!

```
$ lar -c my reco sbnd basic.fcl -n -1 -s /path/to/my/reco1/file.root -o alternative reco2 events.root
```