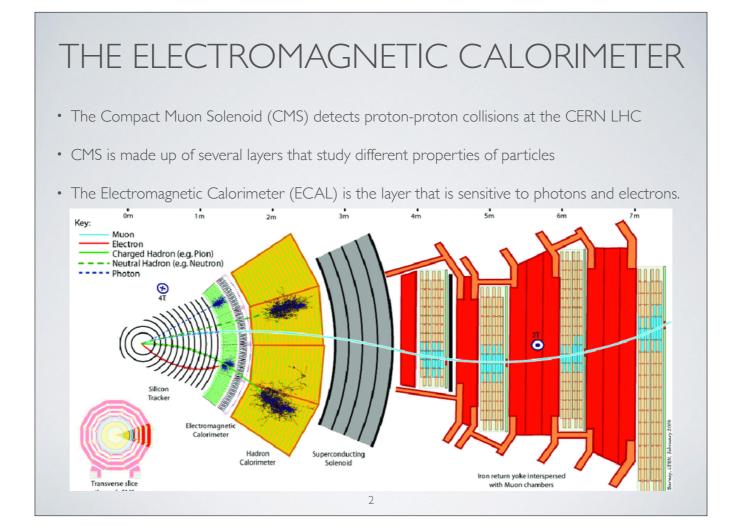
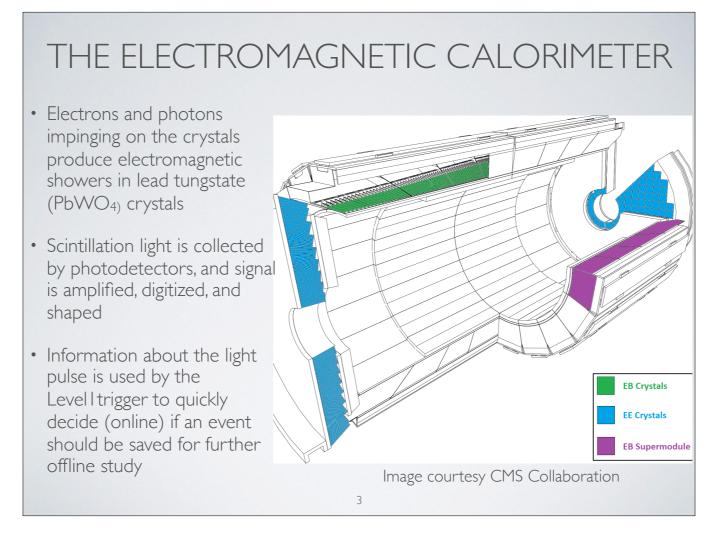


IMPROVING THE CMS ECALTRIGGER FOR RUN 3 OF THE CERN LHC

Kirsten Randle Prof. Toyoko Orimoto

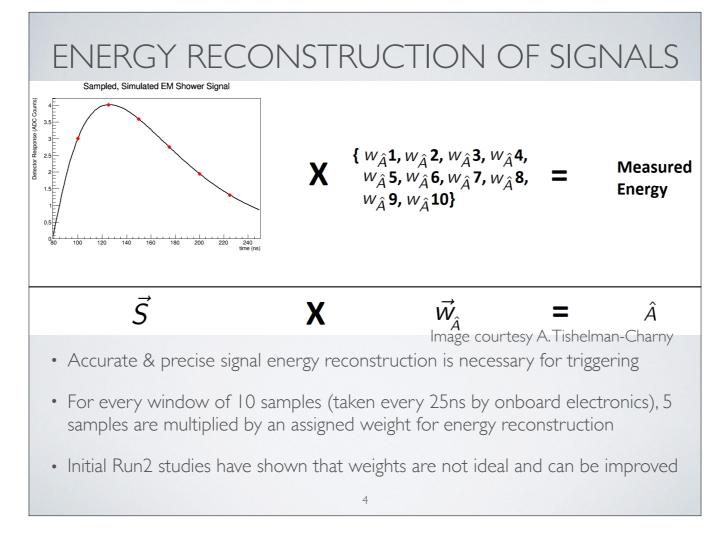


layers: -silicon tracker, -e cal, -hcal, -solenoid (not a detector, produces B-field) , - muon chambers



CMS is split into 2 parts -barrel -endcaps

Crystals are first in groups of 5 crystals for one board (VFE -very front-end) and then in larger groups that depend on the angle from the beam-axis ($eta = -ln(tan(\theta)/2)$)



Weights are designed to preferentially amplify the peak amplitude over the background to get more accurate energy reconstruction of the event

WEIGHTS

- · Amplitude weights
 - active, uniform for whole detector in run 2
 - Current study: update weights and increase granularity by choosing different weights for different parts of the detector for run 3
- Timing weights
 - very front end readout electronics have unused capacity for second set of weights
 - Current study: optimize timing weights to identify out-of-time pileup in a signal for run 3

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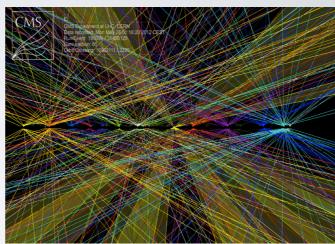
Reminder: VFE -very front end

Crystals in high-eta regions, like the endcaps are especially susceptible to radiation damage and may need unique weights, while weights are currently uniform for the whole detector

Run 2 — 2015-2018 Run 3 — 2021-2024

PILEUP

- Bunches of protons meet inside the detector every 25 ns, this is called a Bunch Crossing (BX)
- Scintillation in the detector takes much longer than this (10 samples, 250 ns)
- Pieces of signals from other bunch crossings can add to overall amplitude
- Out-of-time pileup not only changes the amplitude but also the pulse shape



SCOPE OF PROJECT

- Get ROOT and pyROOT running
- Write a flexible plotter that uses ROOT
- Produce interesting of plots of various parameters used to study the amplitude and timing weights from simulated data

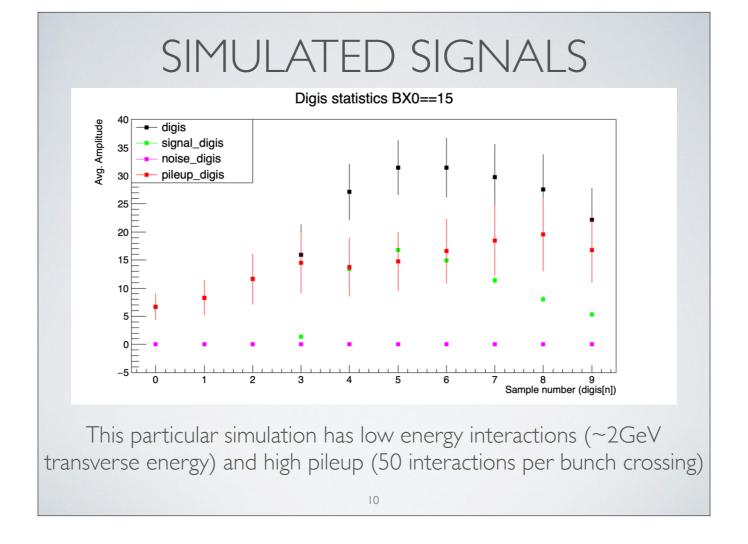
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ROOT is C++ based particle physics analysis framework pyROOT — root libraries can be imported into python do two layers of code: first use a layer

PLOTTER

- Plotter is split into 2 layers to separate time-heavy processes for ease of use
- ROOT libraries are used for power to deal with large datasets
- First layer takes data and produces histogram objects and saves them to a file that can be accessed for the second layer
- Second layer takes histogram objects and plots them, since second layer runs quickly it can be tweaked easily
- Flexibility makes changes to cuts, studying different parameters, or repeating the same studies on different data sets easy

```
65 def create_1Dhisto(bias_tree, histo_name, binparams, parameter, cuts):
                 h = TH1F(histo_name,parameter,binparams[0],binparams[1],binparams[2])
                 h.GetXaxis().SetTitle(parameter)
67
68
                 h.GetYaxis().SetTitle('Entries')
                  drawstatement = parameter + ' >> ' + histo_name
69
70
                  bias_tree.Draw(drawstatement,cuts,'hist')
71
                  h.SetDirectory(0)
72
                  return h
73
74 def create_2Dhisto(bias_tree,histo_name,binparams,parameters,cuts):
75
                             TH2F (histo\_name, histo\_name, binparams[0][0], binparams[0][1], binparams[0][2], binparams[1][0], binparam
                             [1][1],binparams[1][2])
76
                  h.GetXaxis().SetTitle(parameters[0])
77
                  h.GetYaxis().SetTitle(parameters[1])
                  drawstatement= parameters[1] + ':' + parameters[0] + ' >> ' + histo_name
78
                  bias_tree.Draw(drawstatement,cuts,'COLZ1')
79
80
                  h.SetDirectory(0)
81
                  return h
82
83 def slicefity(histo,func,slicebins,options):
                   fitparams = TObjArray()
                  histo.FitSlicesY(func,slicebins[0],slicebins[1],slicebins[2],options,fitparams)
                  return fitparams
86
87
88 def iterate_curves(tree,curvelist,type):
                  list = [0 for x in range(len(curvelist))]
89
90
                  for i,p in enumerate(curvelist):
91
                           if type == 'TH1F':
                                     list[i] = create_1Dhisto(tree,p[0],p[2],p[3],p[4])
92
                            if type == 'TH2F':
93
94
                                     list[i] = create_2Dhisto(tree,p[0],p[2],p[3],p[4])
95
                 return list
                                                                                                                                       9
```



The average signal (green)

Average electronic noise (magenta) (can be positive or negative, small amplitude, average is zero)

Pileup (red) is very variable— only 1 BX is studied, which means that different bunch crossings may have different pileup behavior (BXs with more or fewer events preceding/following it will have different distributions)

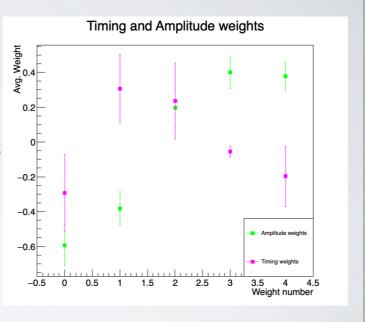
This plot is at eta ring 28, bunch crossing 15

Statistics are generated from histograms of each digis[n]

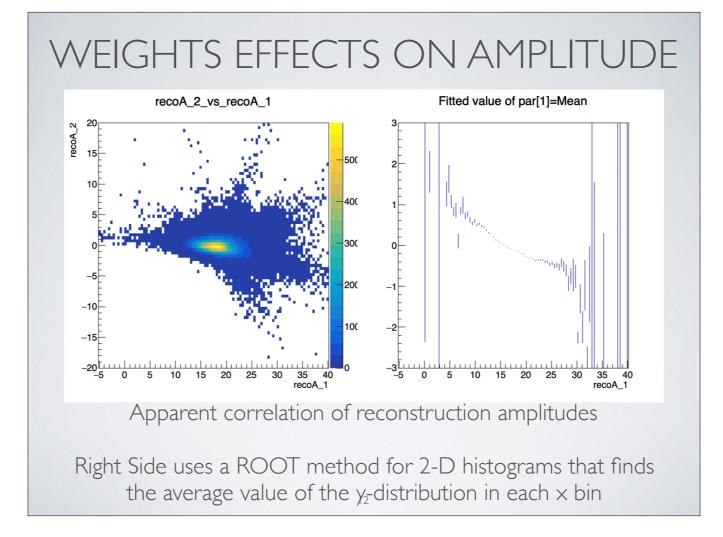
WEIGHTS

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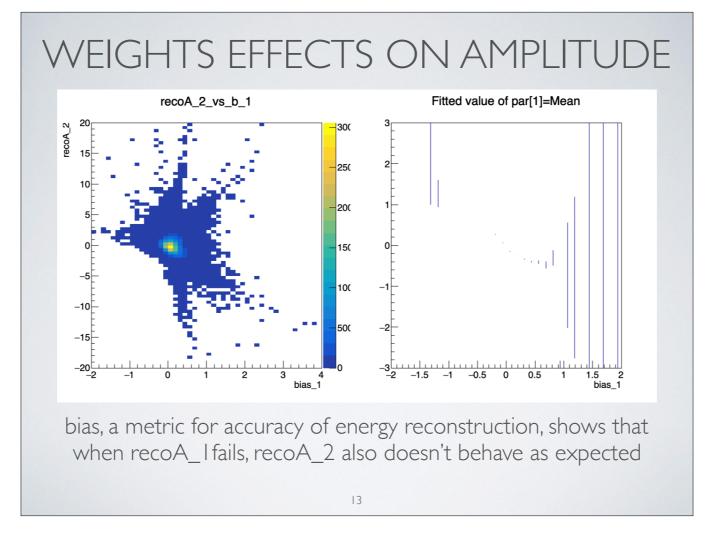
- The model uses the 'true' amplitude to choose the weights that accurately reconstruct the amplitude for each event
- The model similarly chooses the timing weights that reconstruct the amplitude to zero for each event
- This plot represents the average weights selected over 320,000 events



Check with Abe on how accurate description of timing weights is



Examining behavior of recoA_2 vs recoA_1: we expect the average of recoA_2 to be constant at zero, and for recoA_1 and recoA_2 to be uncorrelated



Examining behavior of recoA_2 vs recoA_1: we expect the average of recoA_2 to be constant at zero, and for recoA_1 and recoA_2 to be uncorrelated

FUTURE DIRECTIONS

- Determine what types of events are failing to produce appropriate timing weights (recoA are far from expected values)
- Quantify in which sample high pileup affects amplitude reconstruction the most
- Study failure modes for amplitude weight failure, specifically the pileup dependence
- Examine Timing weight's ability to identify out-of-time pileup

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Determine which events are failing — this may require examining single events ${\sf Quantify}$