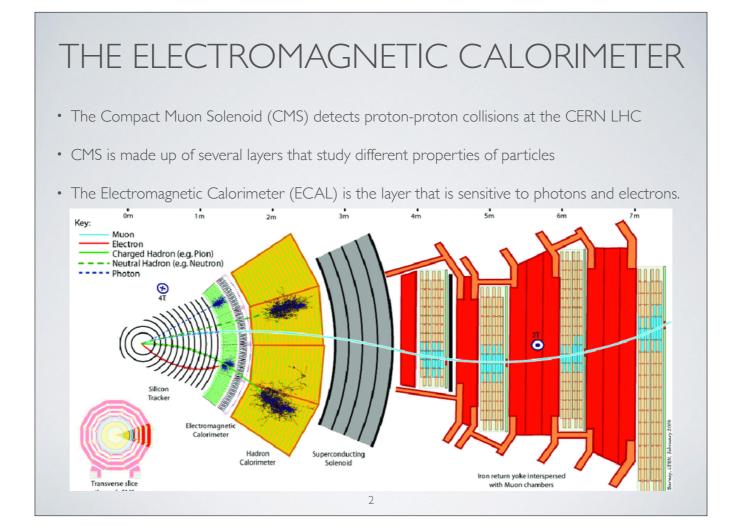


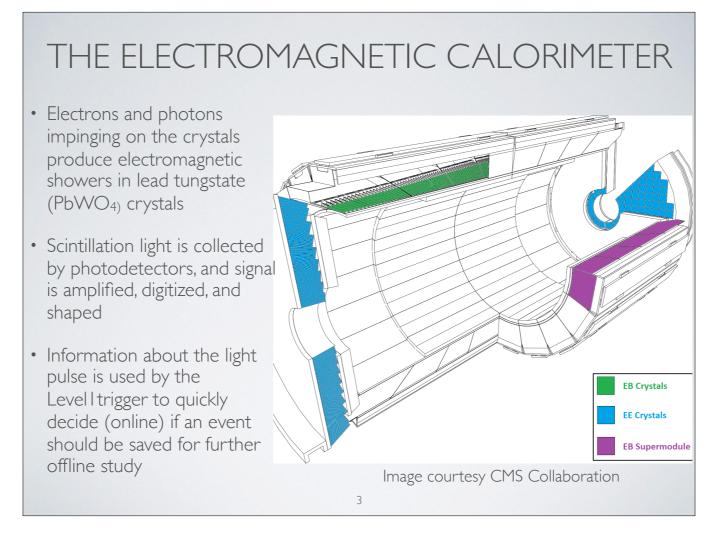
IMPROVING THE CMS ECALTRIGGER FOR RUN 3 OF THE CERN LHC

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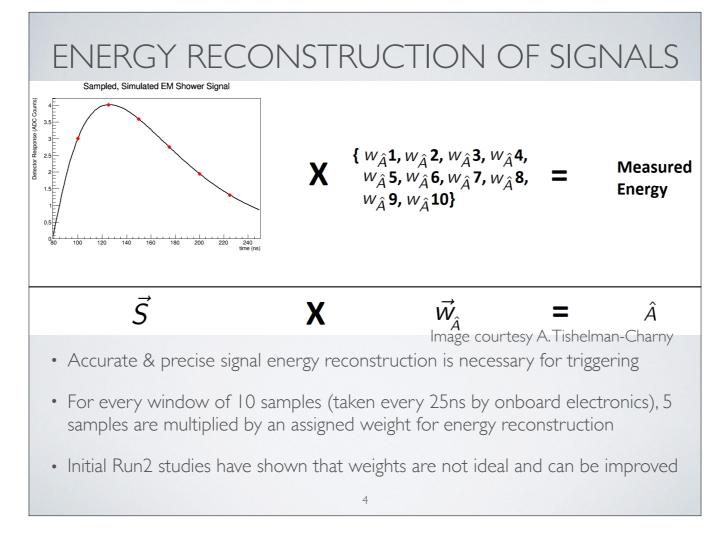


layers: -silicon tracker, -e cal, -hcal, -solenoid (not a detector, produces B-field) , - muon chambers



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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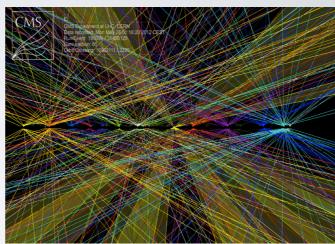
Reminder: VFE -very front end

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

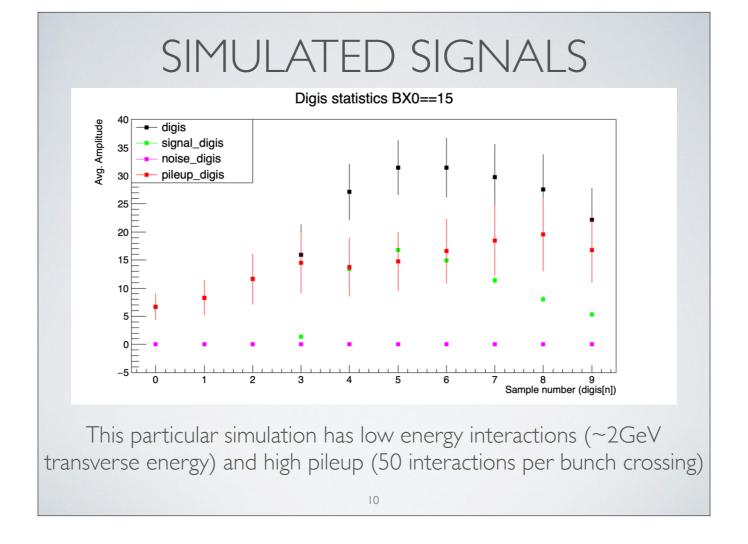
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ROOT is C++ based particle physics analysis framework pyROOT — root libraries can be imported into python do two layers of code: first use a layer

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)

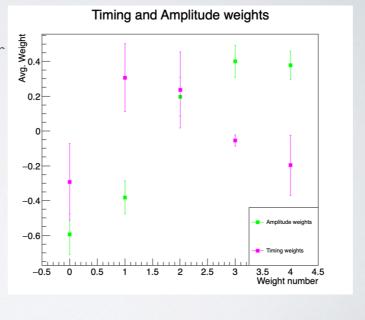
This plot is at eta ring 28, bunch crossing 15

Statistics are generated from histograms of each digis[n]



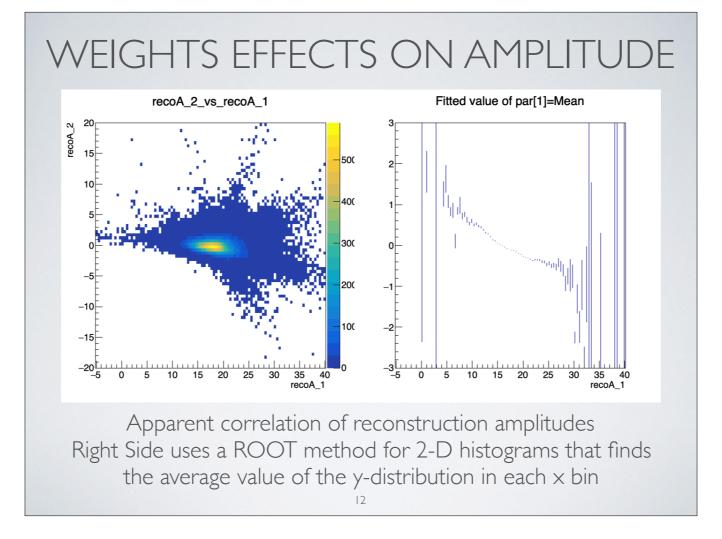
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- The model uses the 'true' amplitude to choose the weights that accurately reconstruct the amplitude for each event
- The model similarly chooses the timing weights that identify if the peak arrives sooner or later than the expected peak
- This plot represents the average weights selected over 320,000 events



Check with Abe on how accurate description of timing weights is peak is at w3

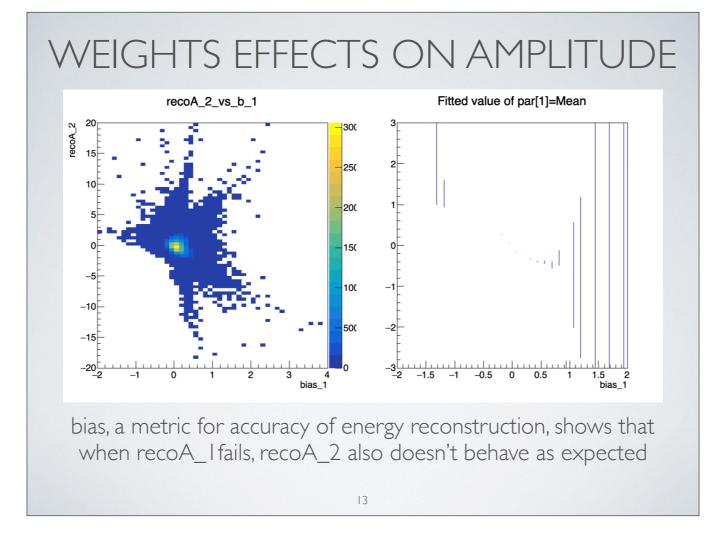
1st two amplitude weights are negative because they are attempting to subtract background (no signal here) two timing weights before peak are high because they are trying to return difference between actual peak time and expected peak time



Examining behavior of recoA_2 vs recoA_1:

we expect the average of recoA_2 to be zero when the waveform is not shifted, and for recoA_1 and recoA_2 to be uncorrelated recall Z is entries

on right: y is the mean from a gaussian fit of each slice



Examining behavior of recoA_2 vs bias_1:

recoA_1 and bias_1 are correlated because the range of true energies is very small, so a high or low reconstructed energy must therefore be inaccurate, and this shows that the correlation between recoA_2 and bias_1 is conserved, and suggests that out-of time peaks are correlated with bad energy reconstruction

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

