# STUDYING THE CMS ECAL TRIGGER AT THE CERN LHC

Kirsten Randle Prof. Toyoko Orimoto

#### THE ELECTROMAGNETIC CALORIMETER

- The Compact Muon Solenoid (CMS) detects proton-proton collisions at the CERN LHC
- CMS is made up of several layers that study different properties of particles
- The Electromagnetic Calorimeter (ECAL) is the layer that is sensitive to photons and electrons.

#### THE ELECTROMAGNETIC CALORIMETER

 Electrons and photons impinging on the crystals produce electromagnetic showers in lead tungstate (PbWO<sub>4)</sub> crystals

 Scintillation light is collected by photodetectors, and signal is amplified, digitized, and shaped

 Information about the light pulse is used by the Level I trigger to quickly decide (online) if an event should be saved for further offline study

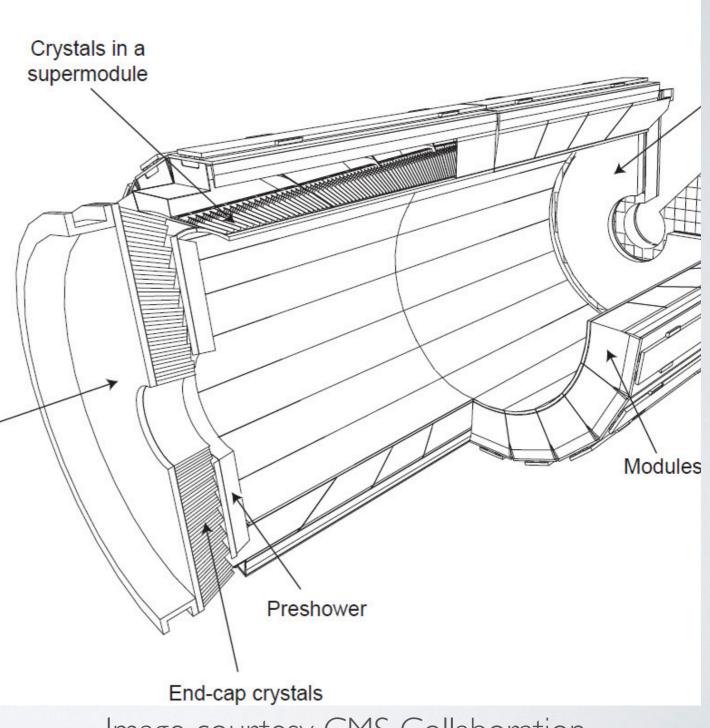
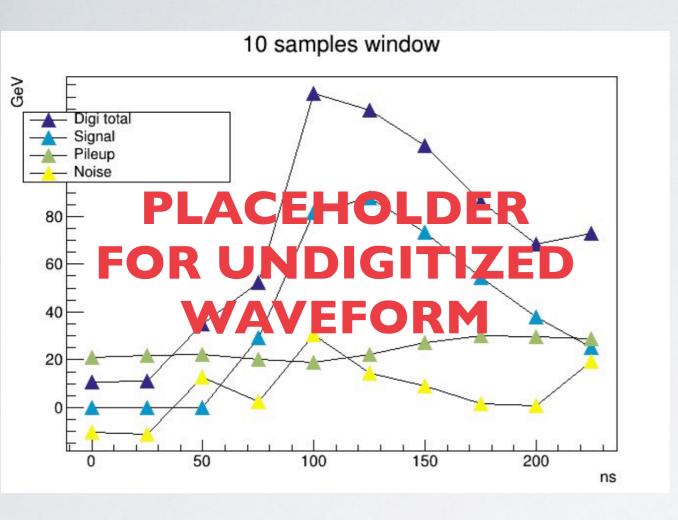
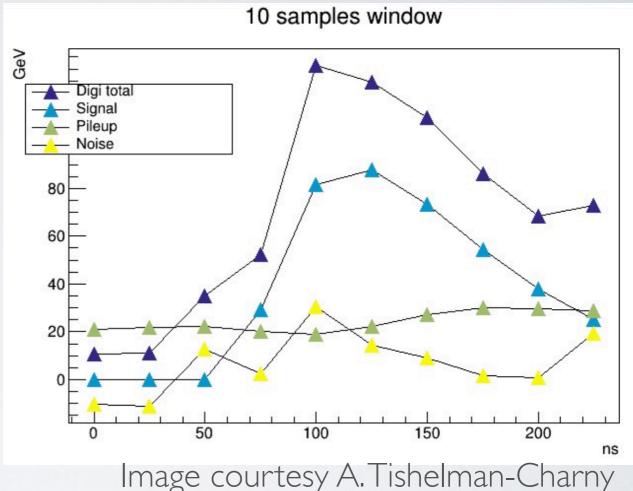


Image courtesy CMS Collaboration

#### ENERGY RECONSTRUCTION OF SIGNALS





- image courtesy A. Hisrielman-Charn
- Accurate & precise signal energy reconstruction is necessary for triggering
- For every window of 10 samples (taken every 25ns by onboard electronics), each sample is multiplied by an assigned weight for energy reconstruction
- · Initial Run2 studies have shown that weights are not ideal and can be improved

### WEIGHTS

- Amplitude weights
  - · active, uniform for whole detector
  - Current study: update weights and vary weights for different parts of the detector
- 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

#### 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

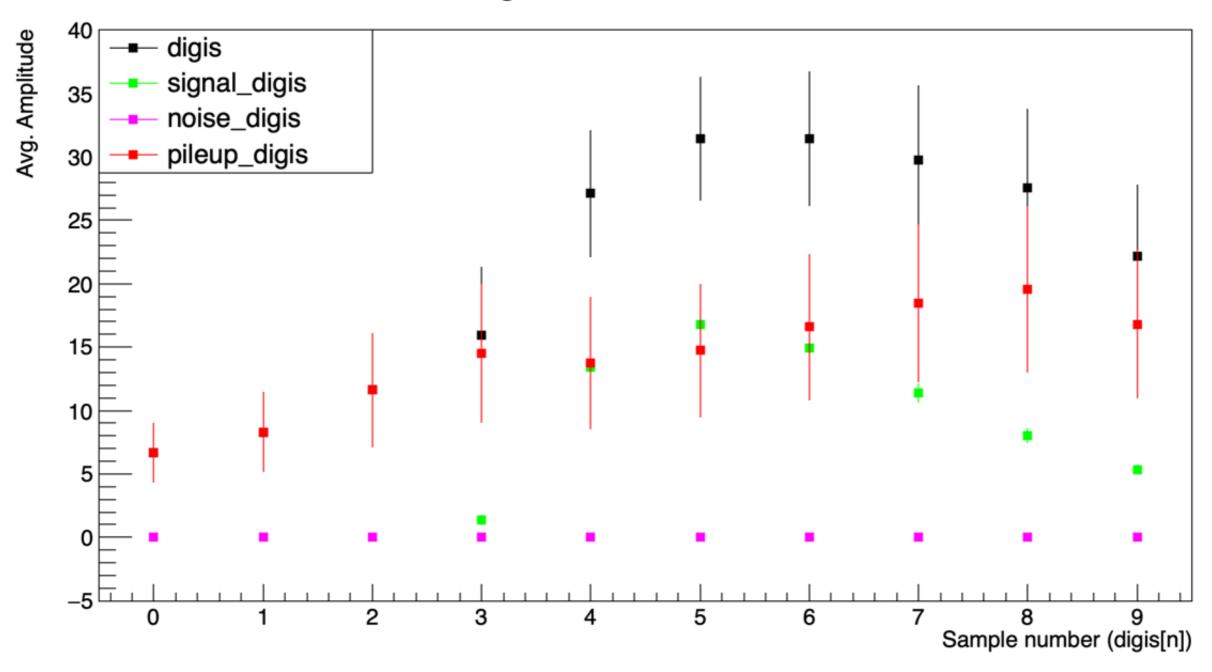
#### PLOTTER

- 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

```
def create_1Dhisto(bias_tree, histo_name, binparams, parameter, cuts):
65
       h = TH1F(histo_name, parameter, binparams[0], binparams[1], binparams[2])
66
       h.GetXaxis().SetTitle(parameter)
67
       h.GetYaxis().SetTitle('Entries')
68
       drawstatement = parameter + ' >> ' + histo_name
69
       bias_tree.Draw(drawstatement,cuts,'hist')
70
       h.SetDirectory(♥)
71
       return h
72
73
   def create_2Dhisto(bias_tree, histo_name, binparams, parameters, cuts):
74
       h =
75
           TH2F(histo_name, histo_name, binparams[0][0], binparams[0][1], binparams[0][2], binparams[1][0], binparams
           [1][1],binparams[1][2])
       h.GetXaxis().SetTitle(parameters[0])
76
       h.GetYaxis().SetTitle(parameters[1])
77
       drawstatement= parameters[1] + ':' + parameters[0] + ' >> ' + histo_name
78
       bias_tree.Draw(drawstatement,cuts,'COLZ1')
79
       h.SetDirectory(♥)
80
       return h
81
82
   def slicefity(histo, func, slicebins, options):
83
       fitparams = TObjArray()
84
       histo.FitSlicesY(func, slicebins[0], slicebins[1], slicebins[2], options, fitparams)
85
       return fitparams
86
87
   def iterate_curves(tree,curvelist,type):
88
       list = [0 for x in range(len(curvelist))]
89
       for i,p in enumerate(curvelist):
90
           if type == 'TH1F':
91
                list[i] = create_1Dhisto(tree,p[0],p[2],p[3],p[4])
92
           if type == 'TH2F':
93
                list[i] = create_2Dhisto(tree, p[0], p[2], p[3], p[4])
94
       return list
95
```

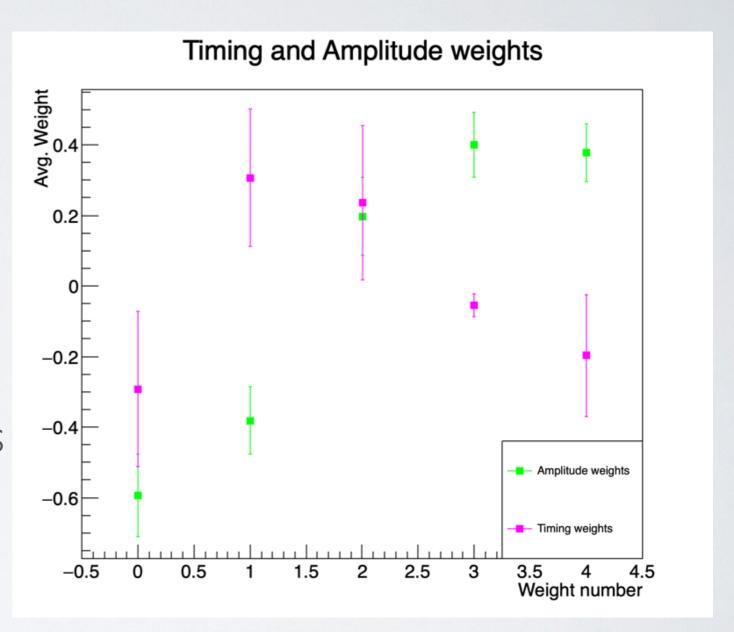
## SIMULATED SIGNALS

#### Digis statistics BX0==15

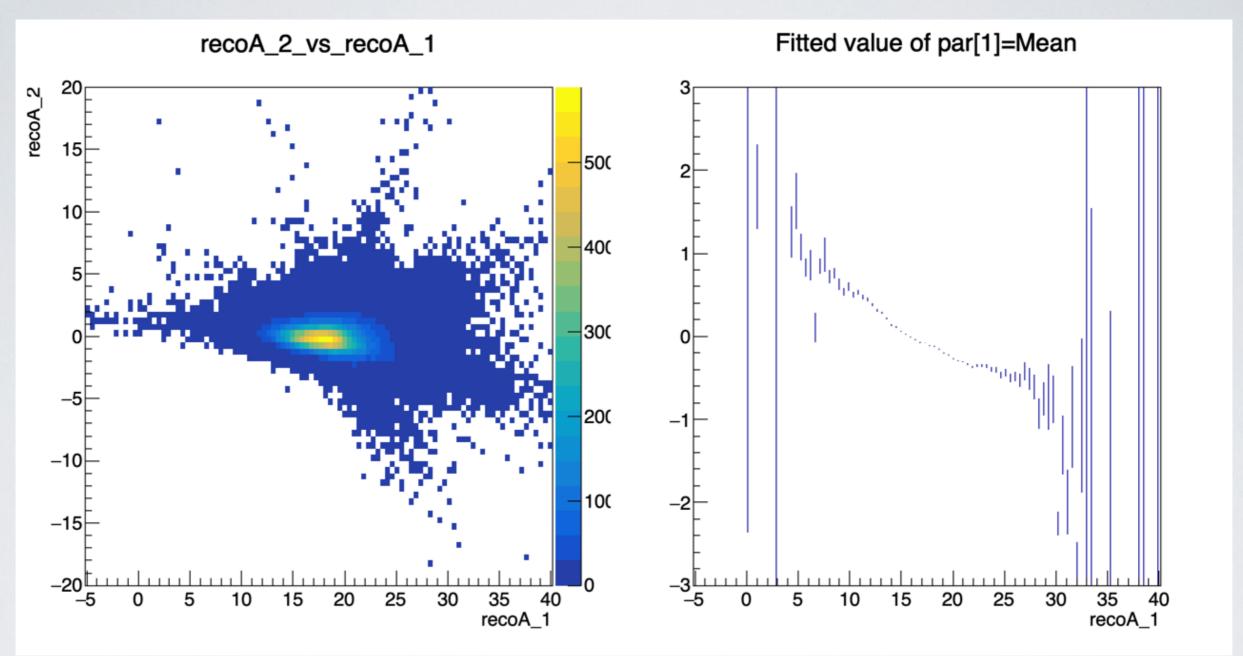


### WEIGHTS

- 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



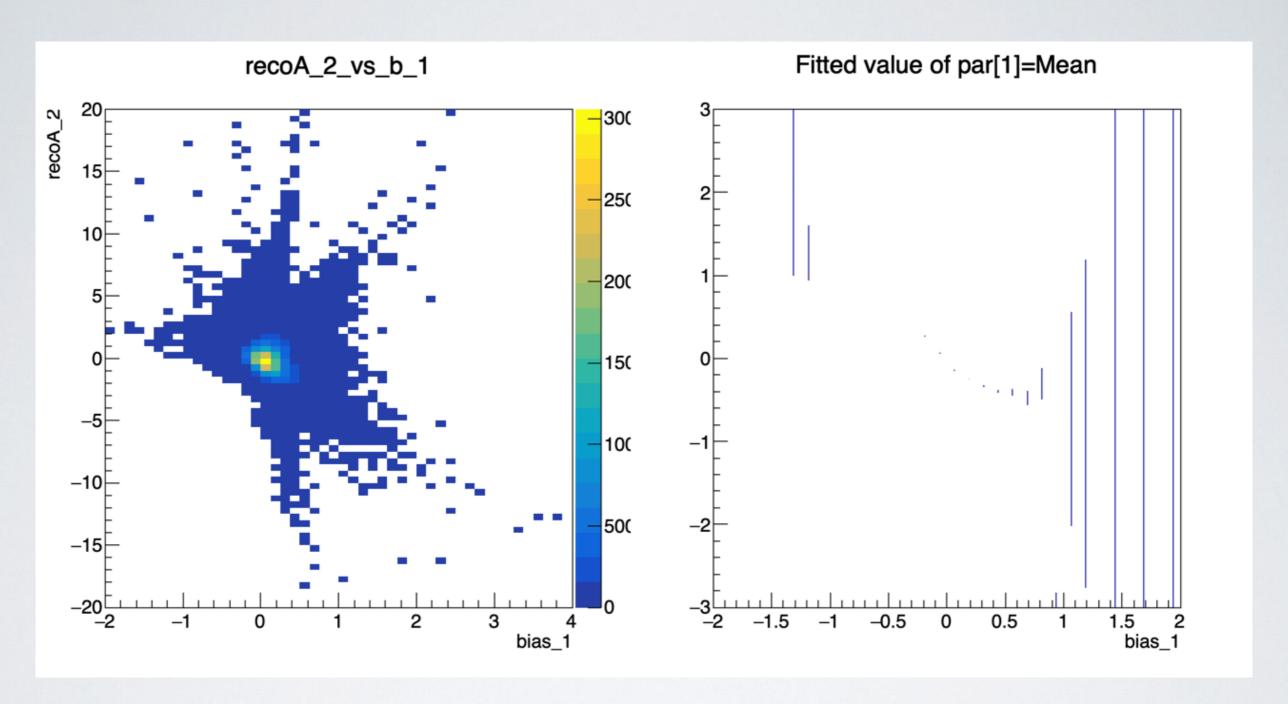
#### WEIGHTS EFFECTS ON AMPLITUDE



Apparent correlation of reconstruction amplitudes

Right Side uses a ROOT method for 2-D histograms that finds the average value of the  $\chi_2$ -distribution in each x bin

#### WEIGHTS EFFECTS ON AMPLITUDE



bias, a metric for accuracy of energy reconstruction, shows that when recoA\_I fails, recoA\_2 also doesn't behave as expected

# 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