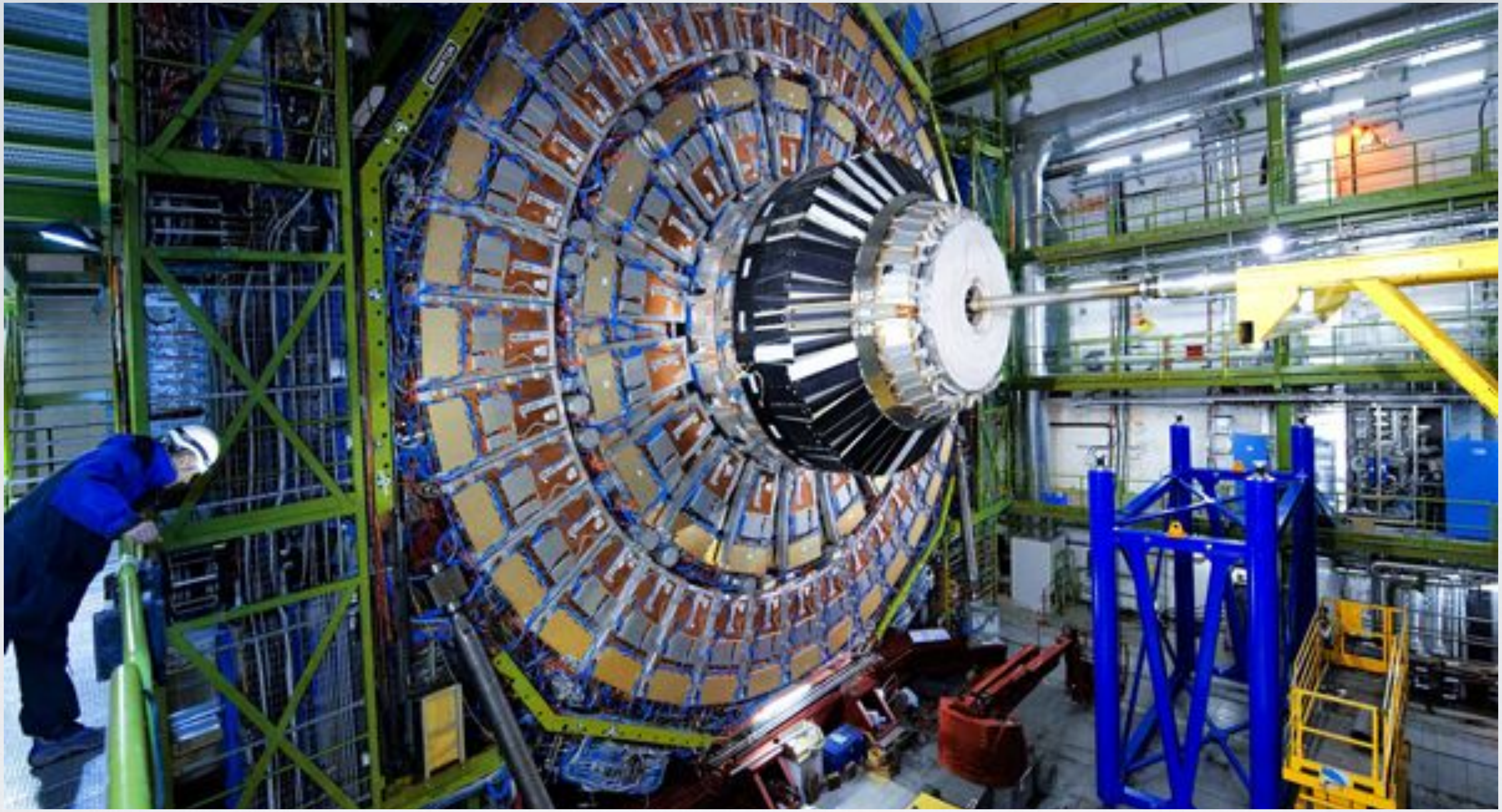

IMPROVING THE CMS ECAL TRIGGER FOR RUN 3 OF THE CERN LHC

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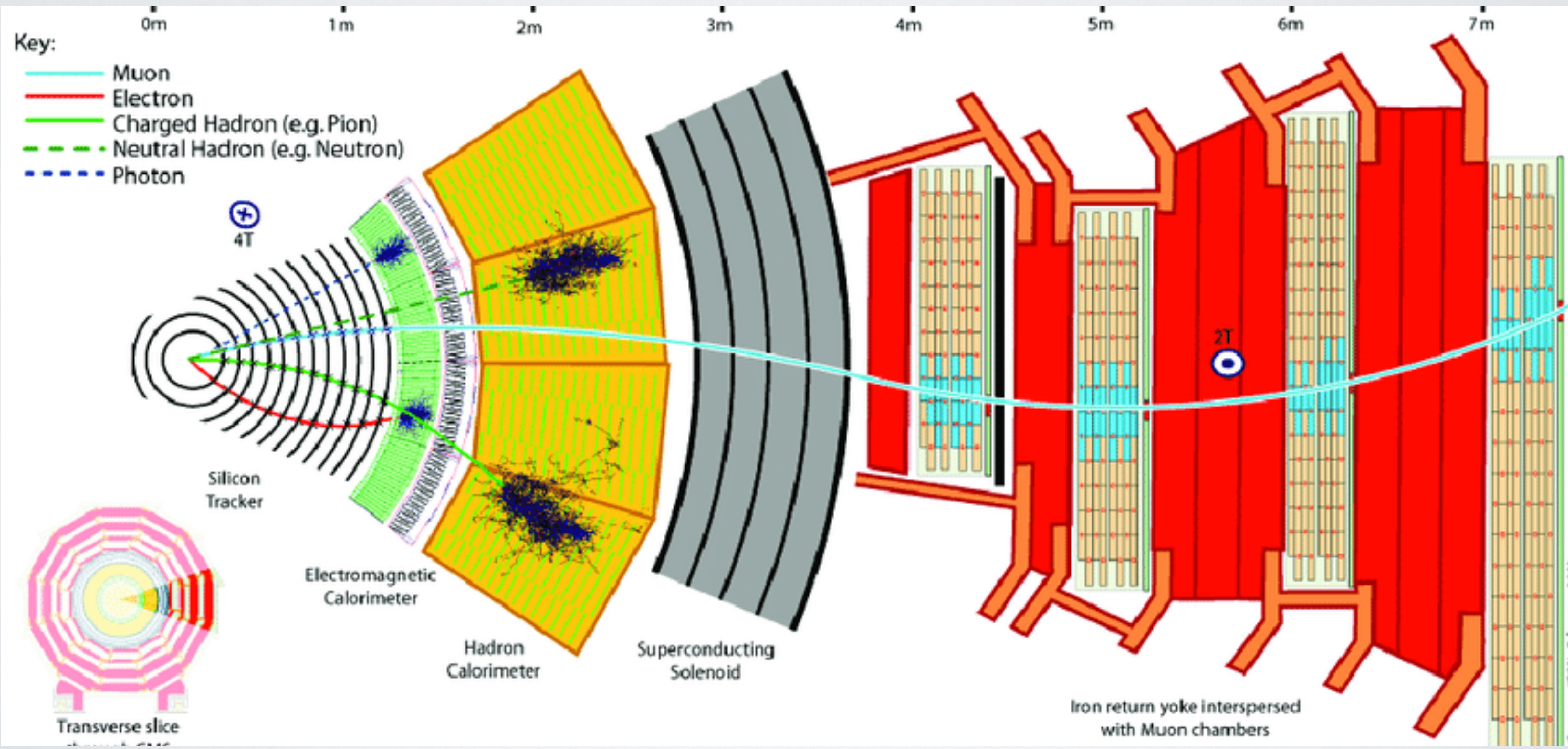
THE ELECTROMAGNETIC CALORIMETER

- 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 ($PbWO_4$) crystals
- Scintillation light is collected by photodetectors, and signal is amplified, digitized, and shaped
- Information about the light pulse is used by the Level1trigger to quickly decide (online) if an event should be saved for further offline study

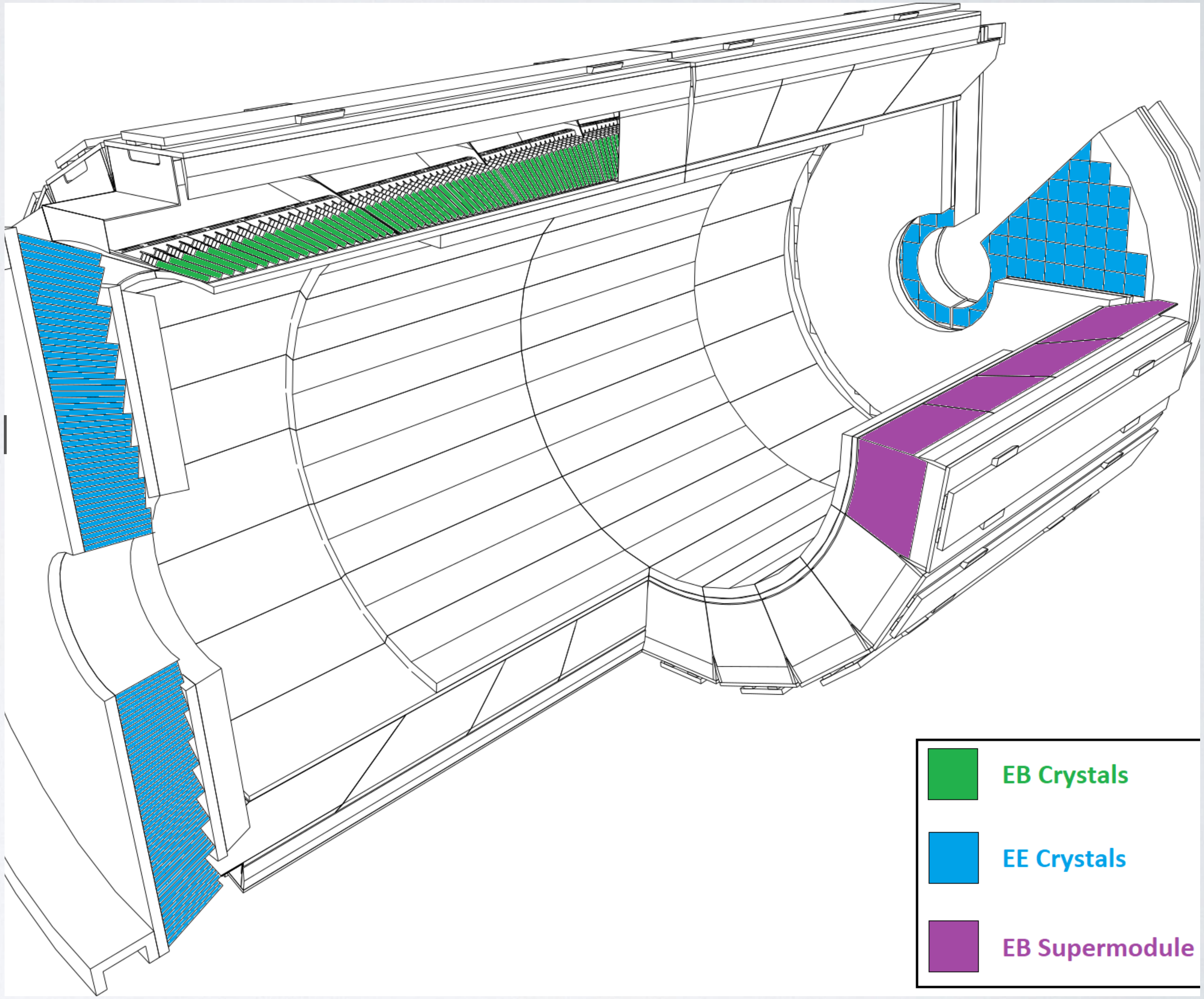
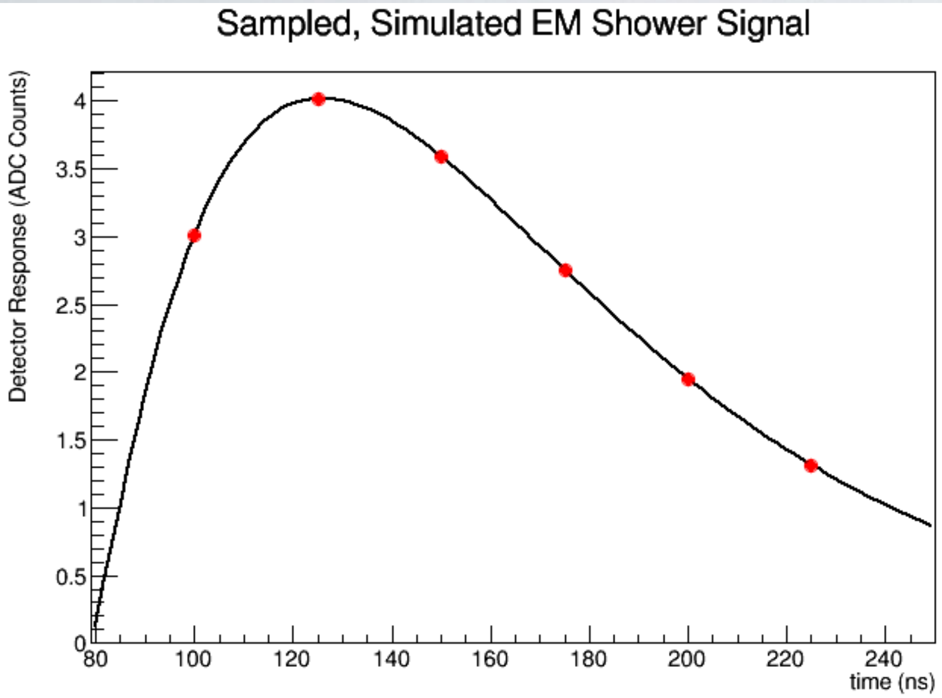


Image courtesy CMS Collaboration

ENERGY RECONSTRUCTION OF SIGNALS



$$\times \quad \{ w_{\hat{A}}1, w_{\hat{A}}2, w_{\hat{A}}3, w_{\hat{A}}4, w_{\hat{A}}5, w_{\hat{A}}6, w_{\hat{A}}7, w_{\hat{A}}8, w_{\hat{A}}9, w_{\hat{A}}10 \} \quad = \quad \textbf{Measured Energy}$$

$$\vec{S} \quad \times \quad \vec{w}_{\hat{A}} \quad = \quad \hat{A}$$

Image courtesy A. Tishelman-Charny

- Accurate & precise signal energy reconstruction is necessary for triggering
- For every window of 10 samples (taken every 25ns by onboard electronics), 5 samples are 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 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

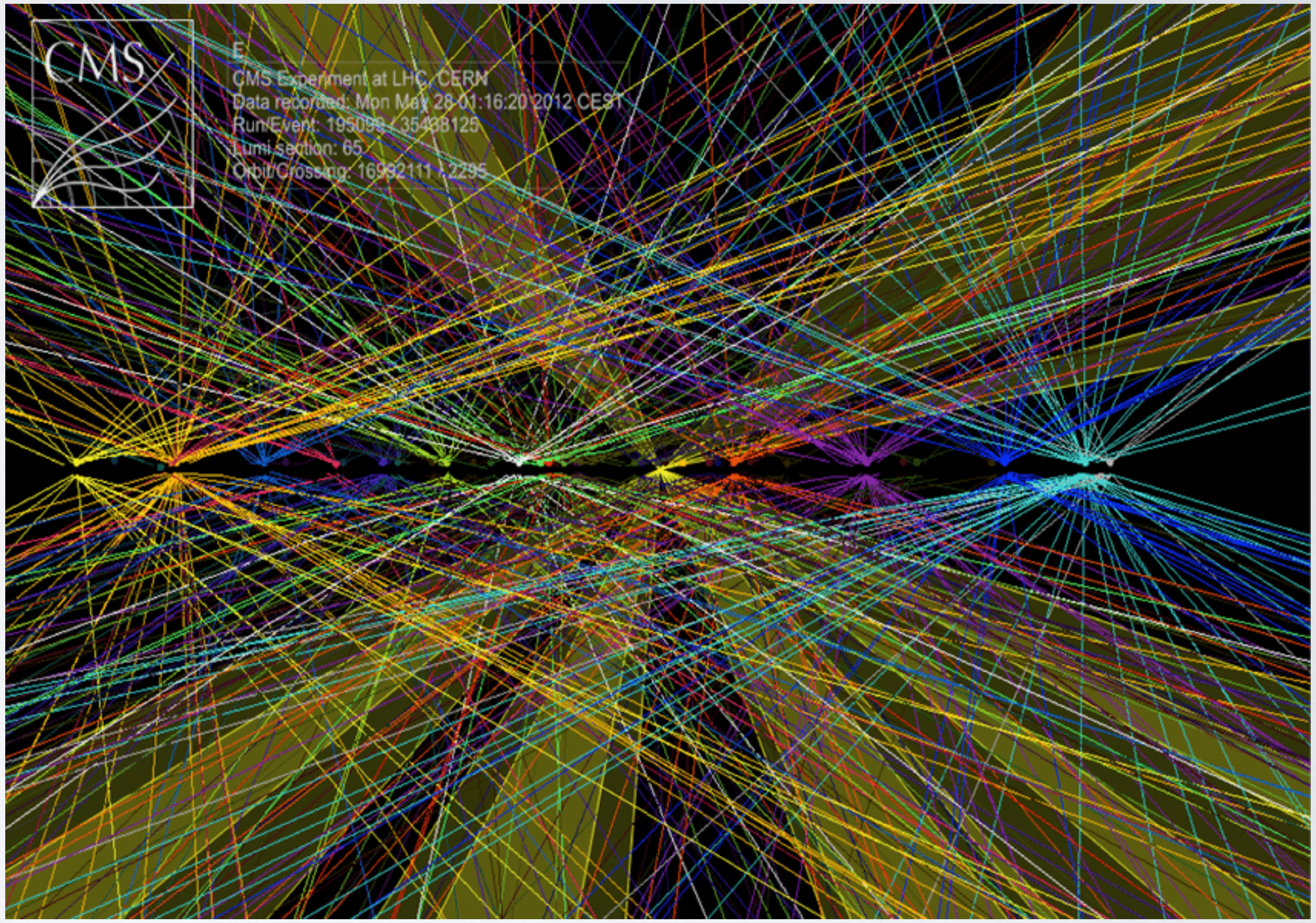
PILEUP

- 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

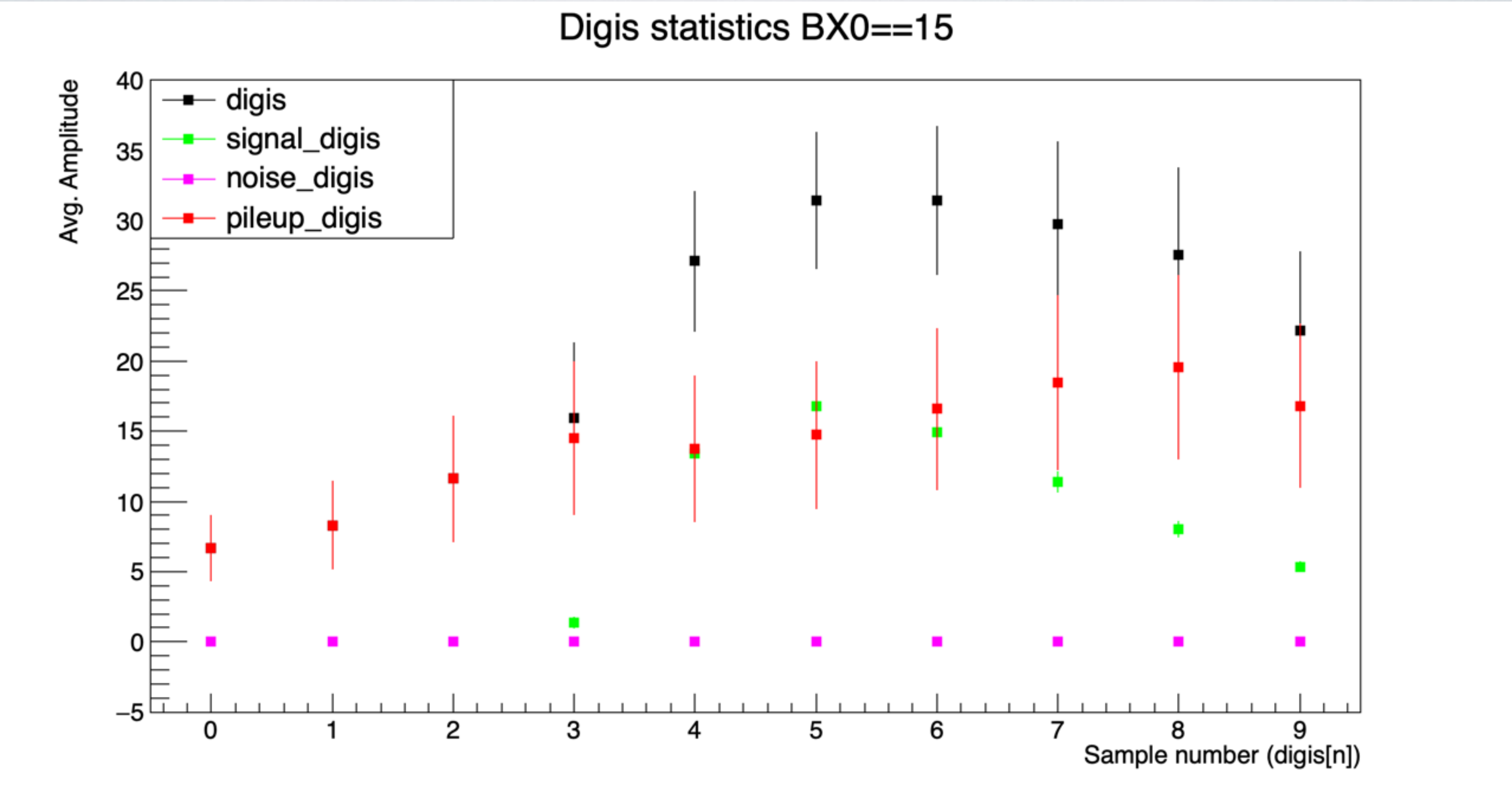
```python
def create_1Dhisto(bias_tree,histo_name,binparams,parameter,cuts):
    h = TH1F(histo_name,parameter,binparams[0],binparams[1],binparams[2])
    h.GetXaxis().SetTitle(parameter)
    h.GetYaxis().SetTitle('Entries')
    drawstatement = parameter + ' >> ' + histo_name
    bias_tree.Draw(drawstatement,cuts,'hist')
    h.SetDirectory(0)
    return h

def create_2Dhisto(bias_tree,histo_name,binparams,parameters,cuts):
    h = 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])
    h.GetYaxis().SetTitle(parameters[1])
    drawstatement= parameters[1] + ':' + parameters[0] + ' >> ' + histo_name
    bias_tree.Draw(drawstatement,cuts,'COLZ1')
    h.SetDirectory(0)
    return h

def slicefity(histo,func,slicebins,options):
    fitparams = TObjArray()
    histo.FitSlicesY(func,slicebins[0],slicebins[1],slicebins[2],options,fitparams)
    return fitparams

def iterate_curves(tree,curvelist,type):
    list = [0 for x in range(len(curvelist))]
    for i,p in enumerate(curvelist):
        if type == 'TH1F':
            list[i] = create_1Dhisto(tree,p[0],p[2],p[3],p[4])
        if type == 'TH2F':
            list[i] = create_2Dhisto(tree,p[0],p[2],p[3],p[4])
    return list
```

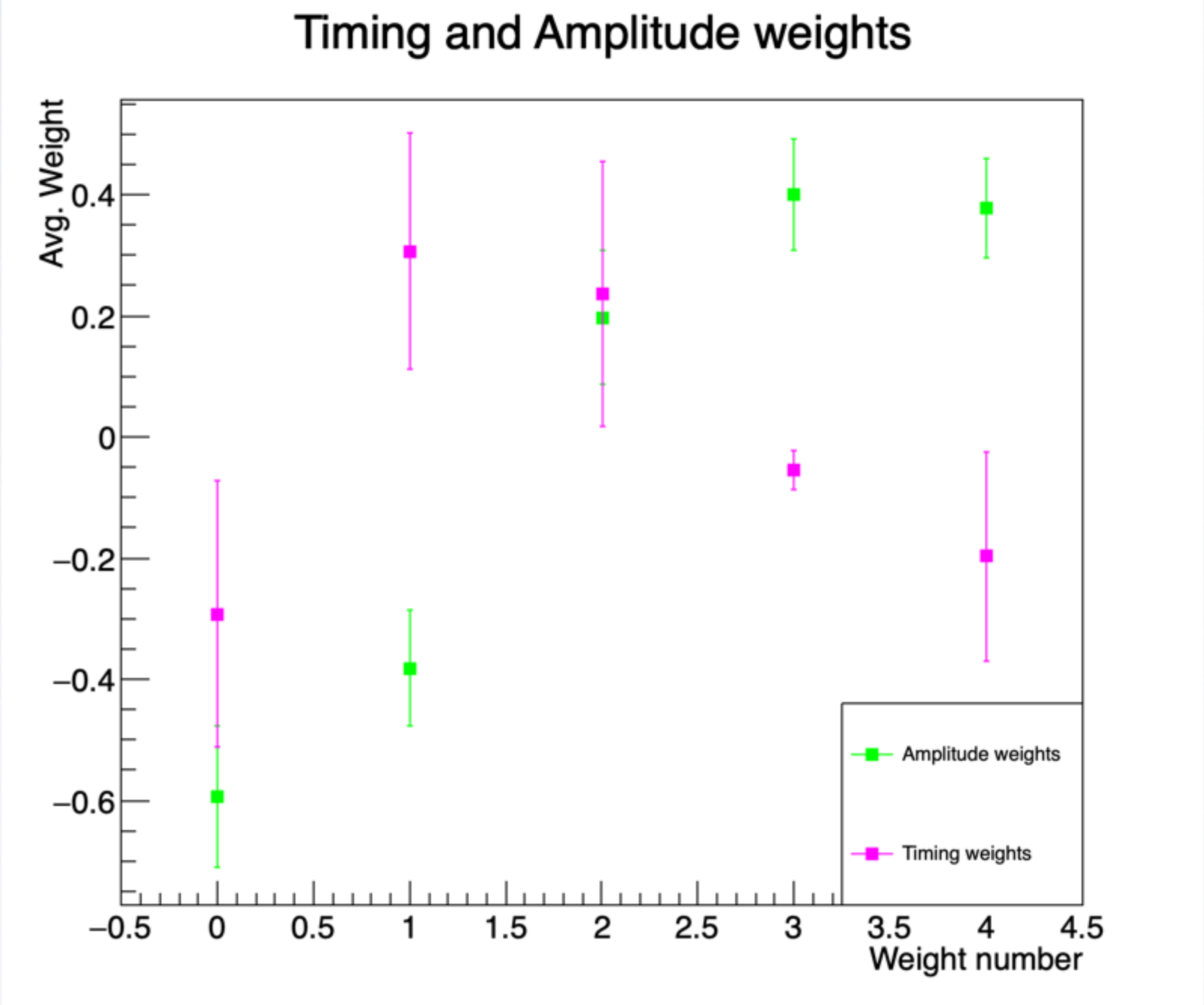
SIMULATED SIGNALS



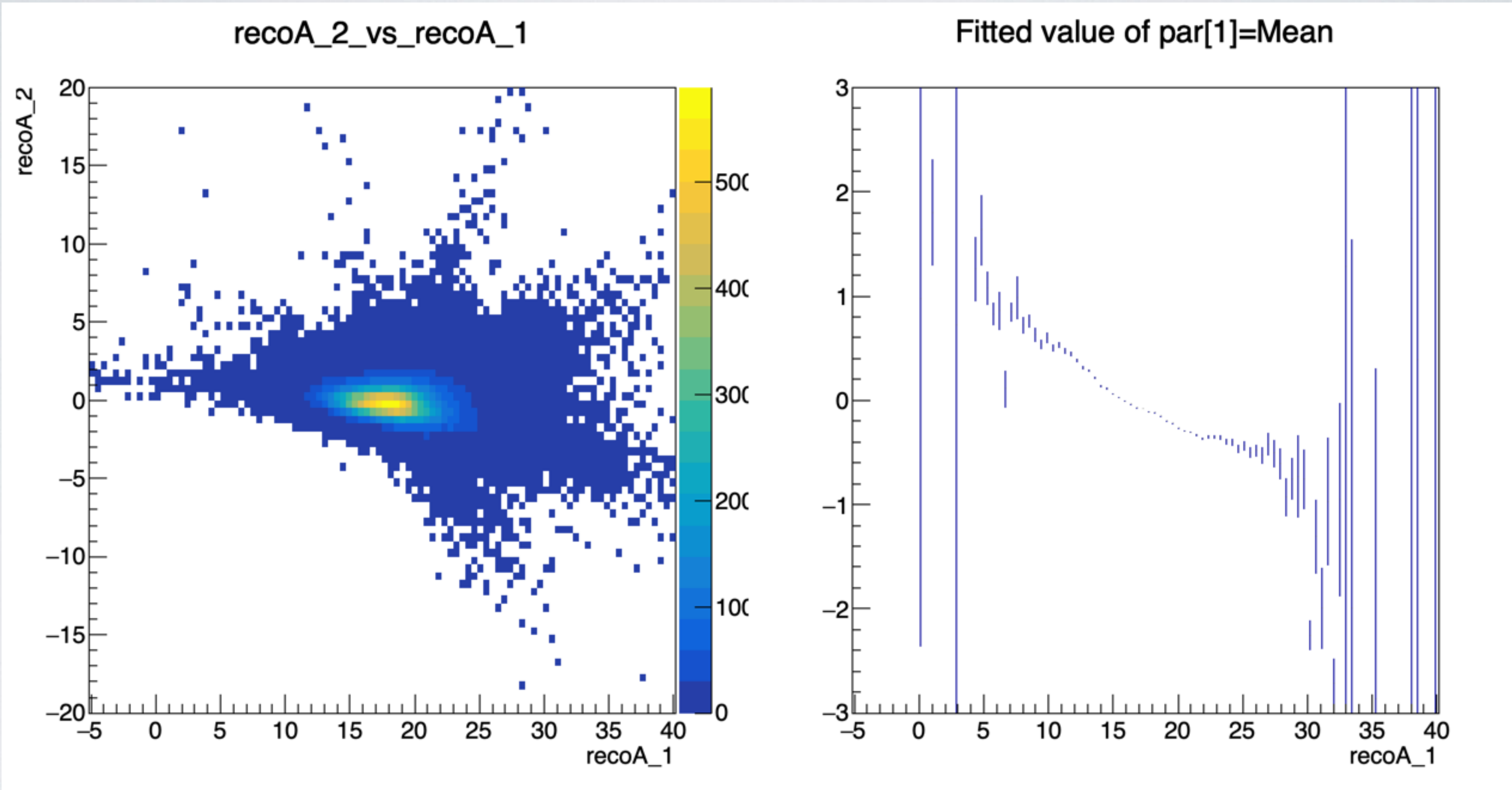
This particular simulation has low energy interactions (~2GeV transverse energy) and high pileup (50 interactions per bunch crossing)

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
- This plot represents the average weights selected over 320,000 events



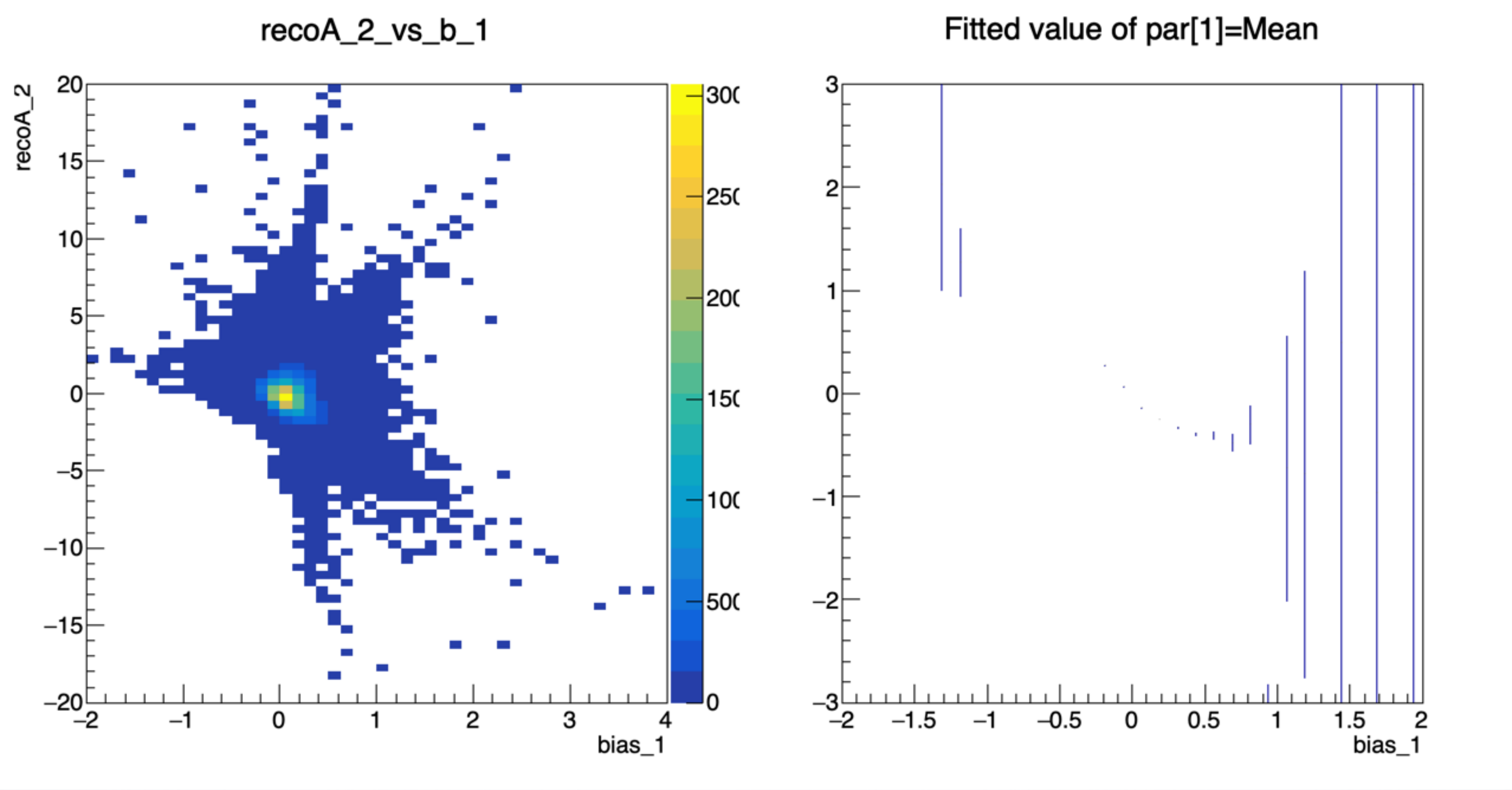
WEIGHTS EFFECTS ON AMPLITUDE



Apparent correlation of reconstruction amplitudes

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

WEIGHTS EFFECTS ON AMPLITUDE



bias, a metric for accuracy of energy reconstruction, shows that when recoA_1fails, 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