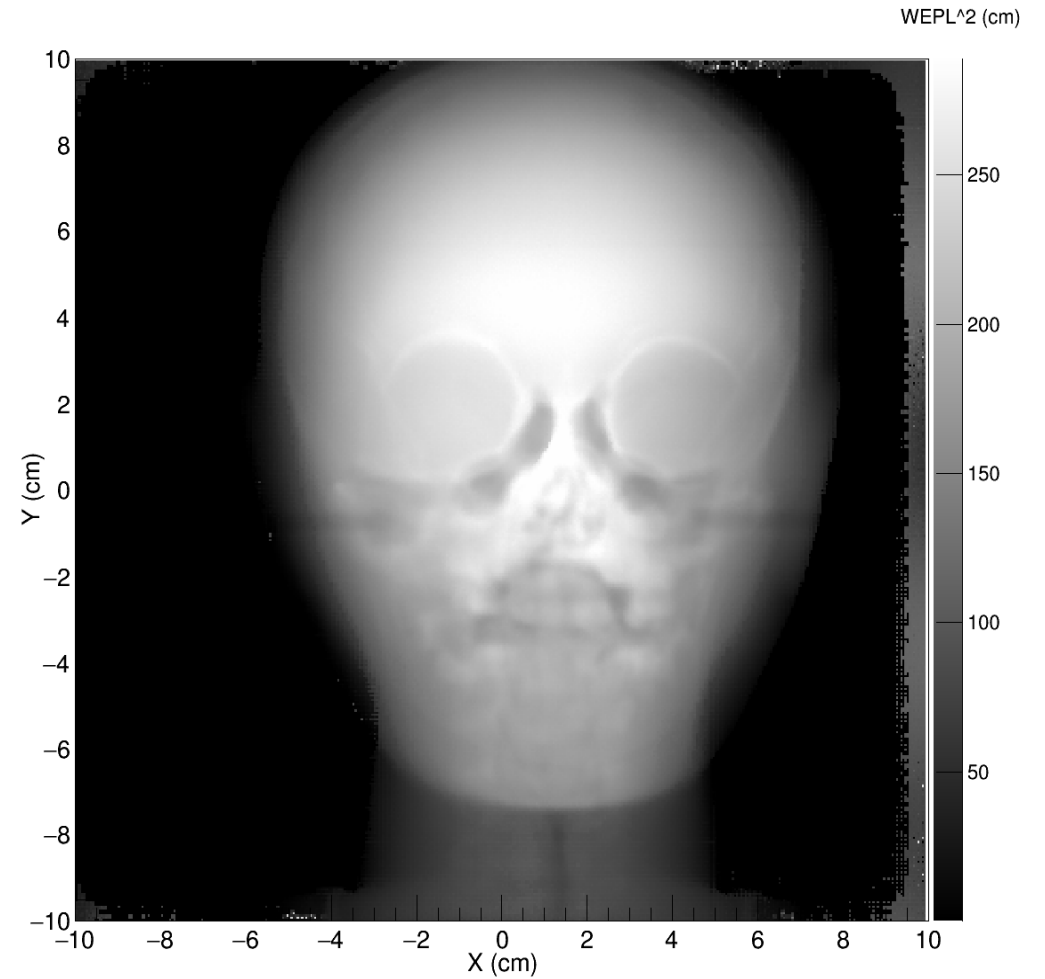
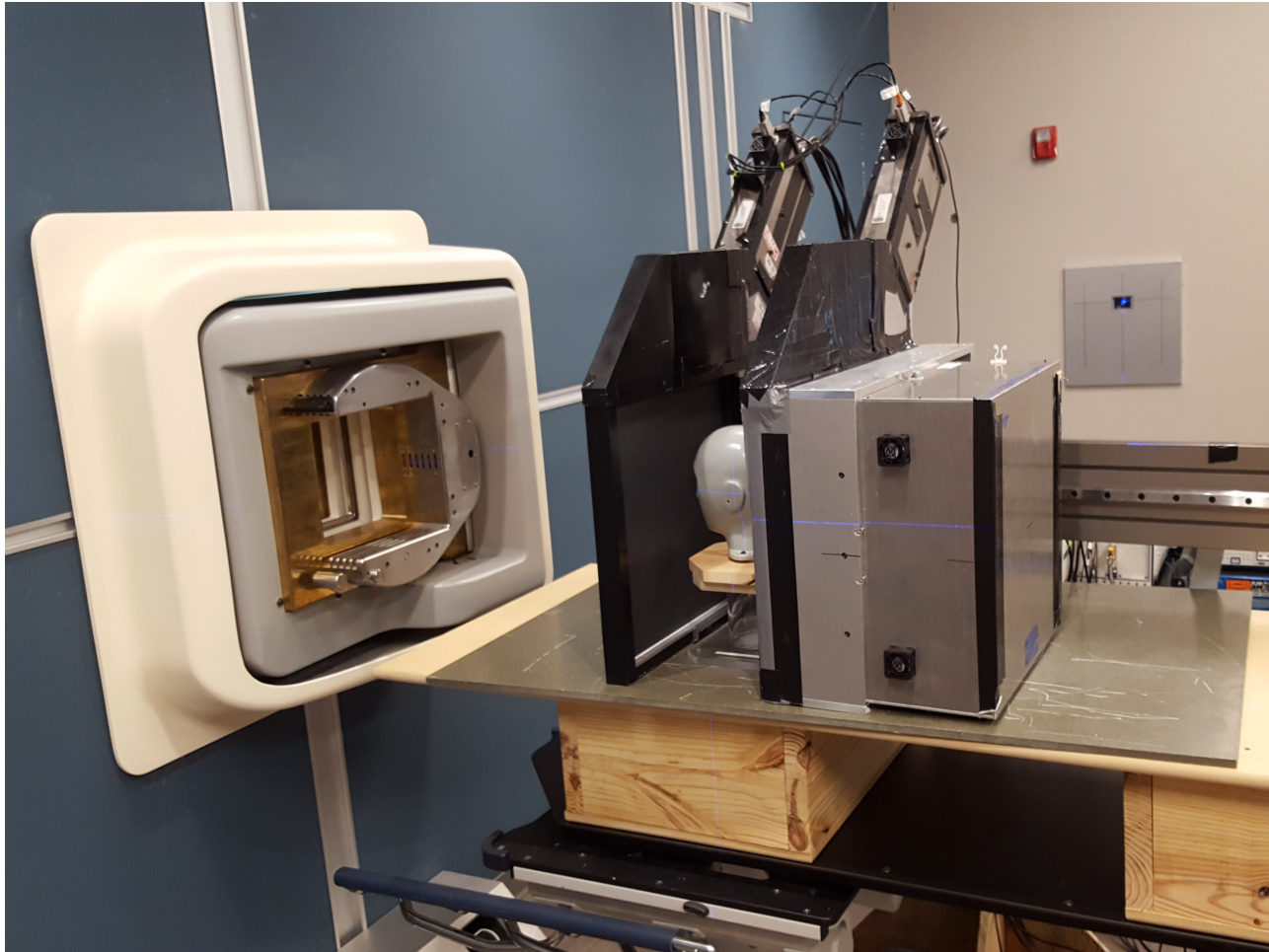
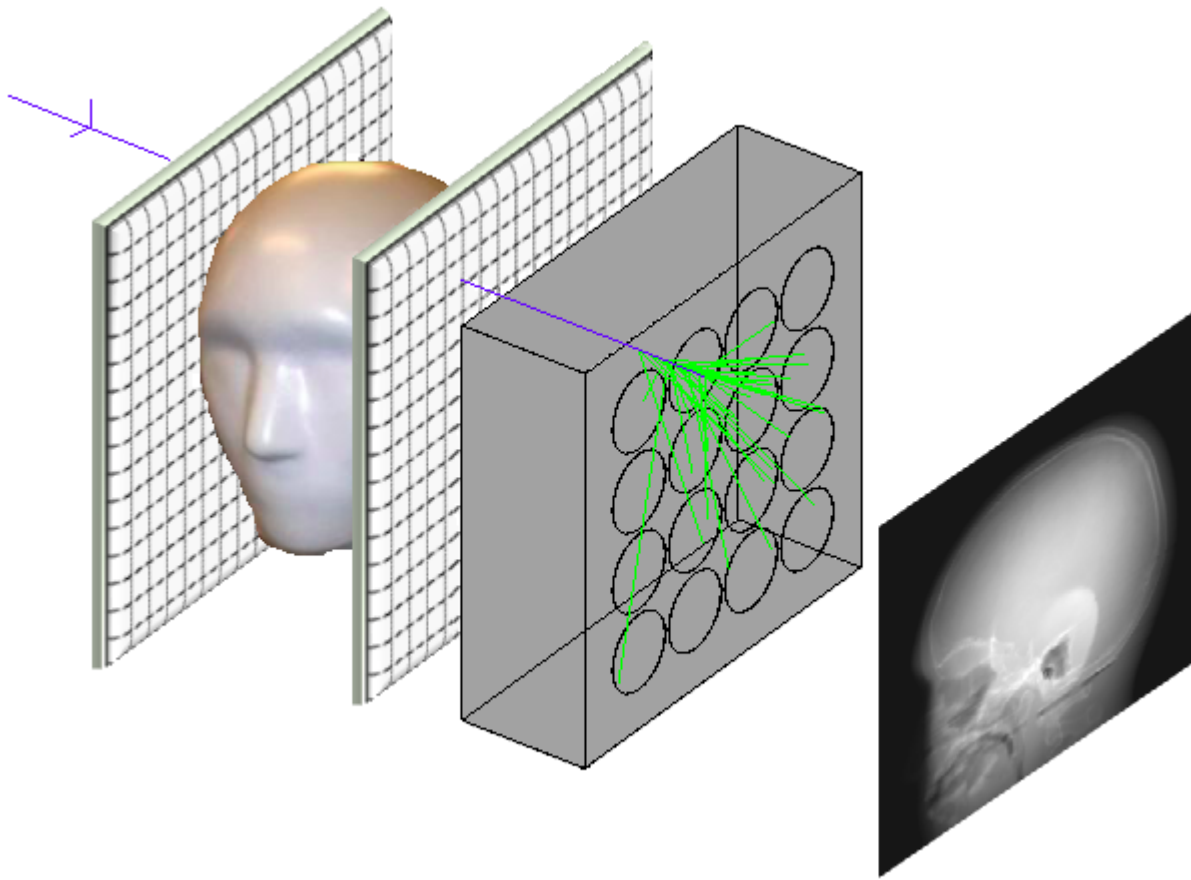
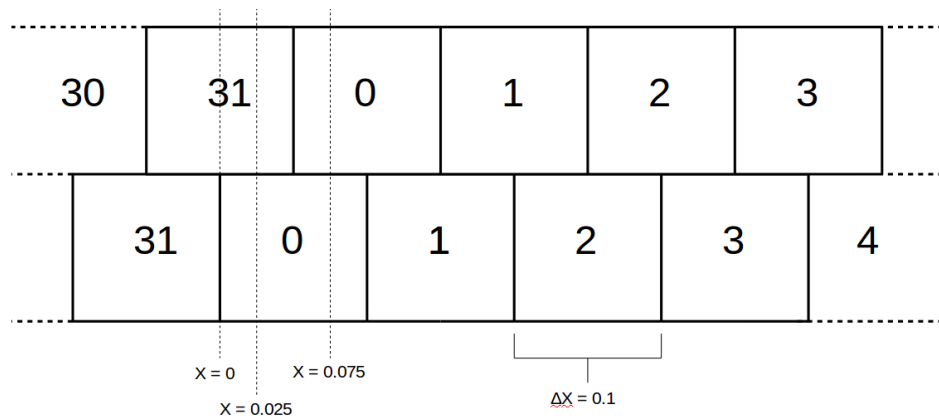


ProtonVDA Radiography Results

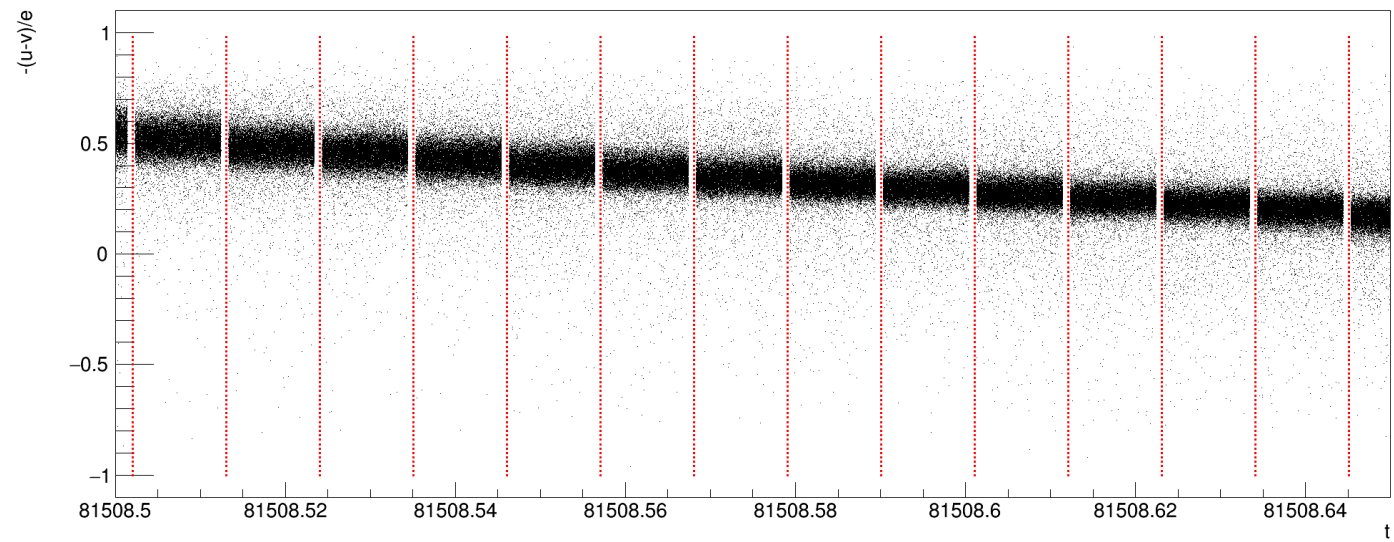
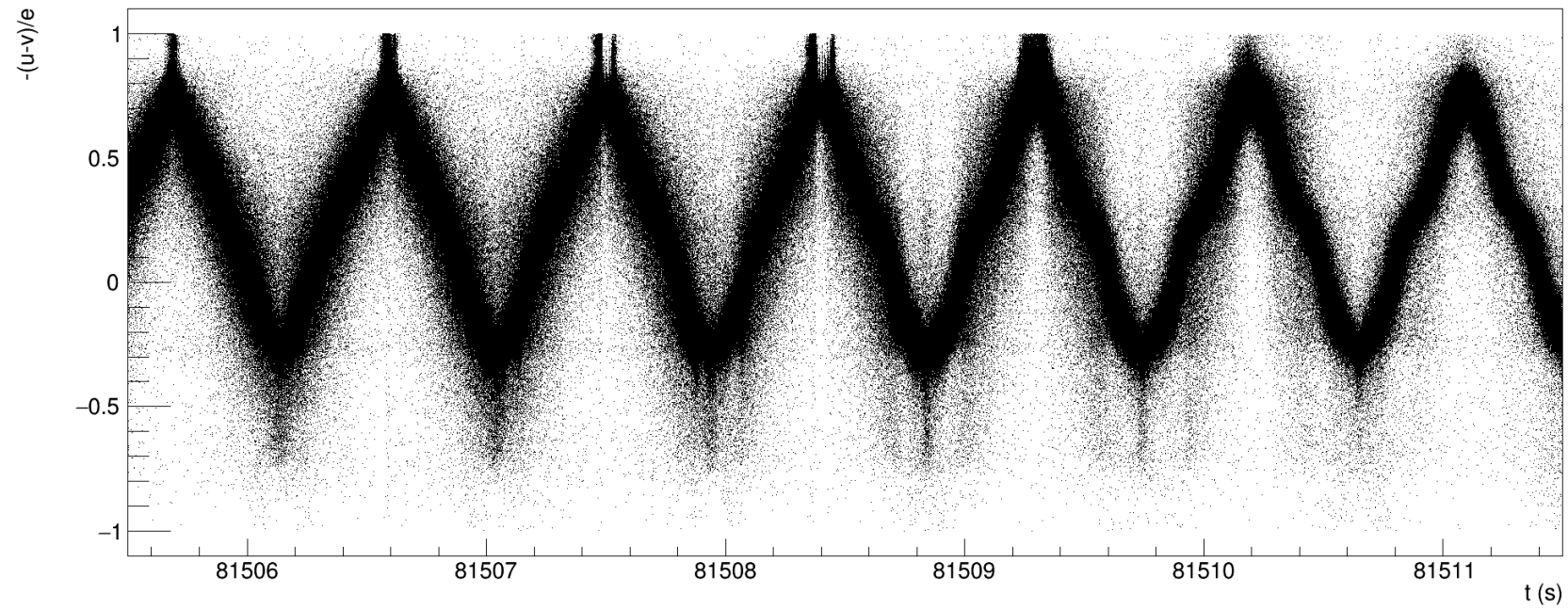




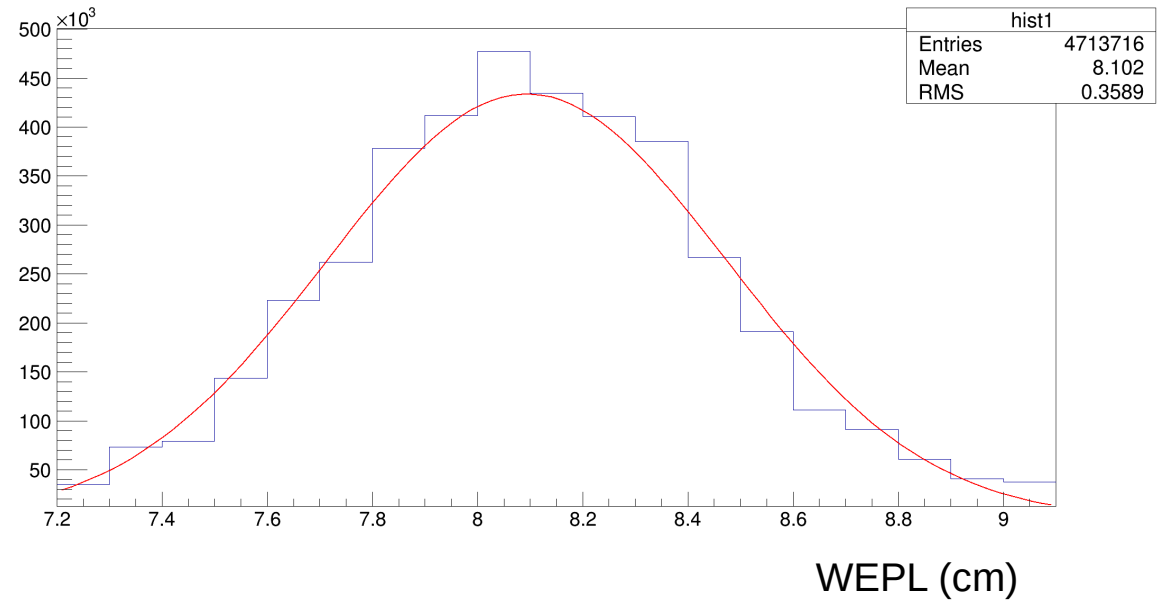
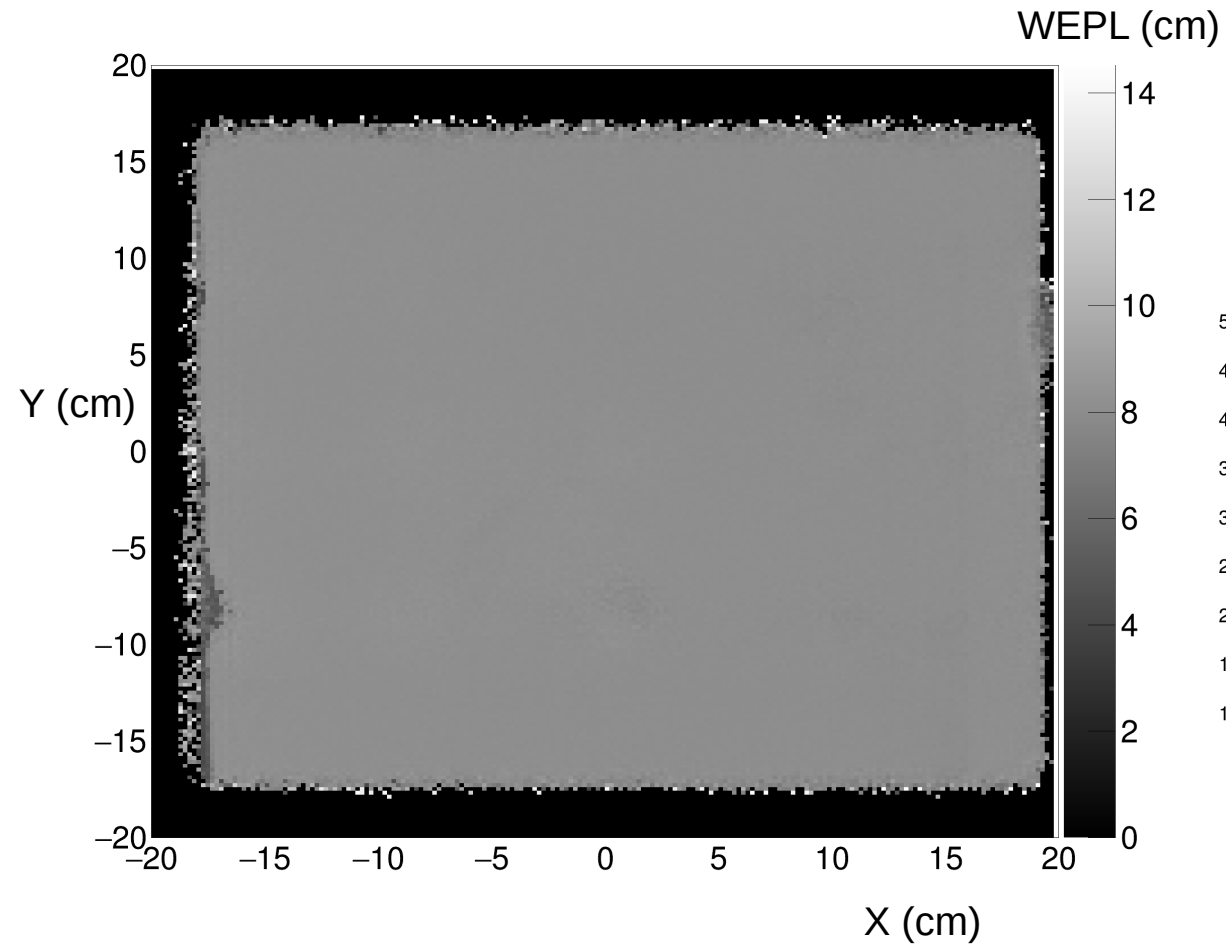
- X-Y tracking planes upstream and downstream – two layers of 1 mm scintillating fibers, offset for 0.5 mm granularity
- Multiplexed fiber readout
 - 32 channels per plane
 - fibers divided into 12 strips
- 40 x 40 x 13 cm block of scintillator for range detector
 - 4 x 4 array of PMTs
 - Output digitized into four channels: E, U, V,C
- Individual protons tracked at up to 10 MHz
- > 99% tracking efficiency
- WEPL resolution ~ 3 mm per proton
- 40 x 40 cm image field size
- Fast (<1 min) image reconstruction for radiograph

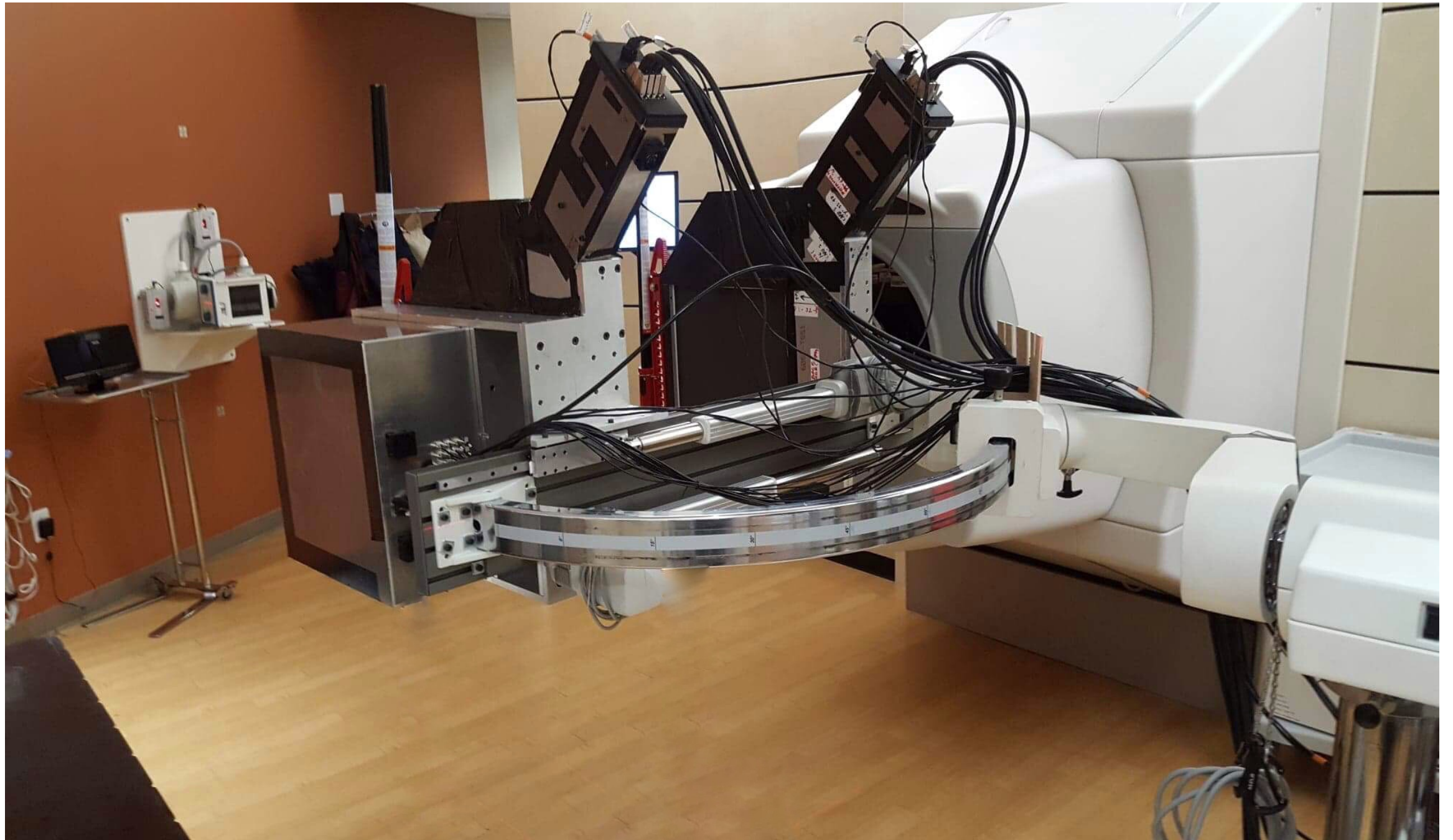


$-(u-v)/e:t\{-(u-v)/e>-1 \ \&\& \ -(u-v)/e<1 \ \&\& \ t>81505 \ \&\& \ t<81512\}$

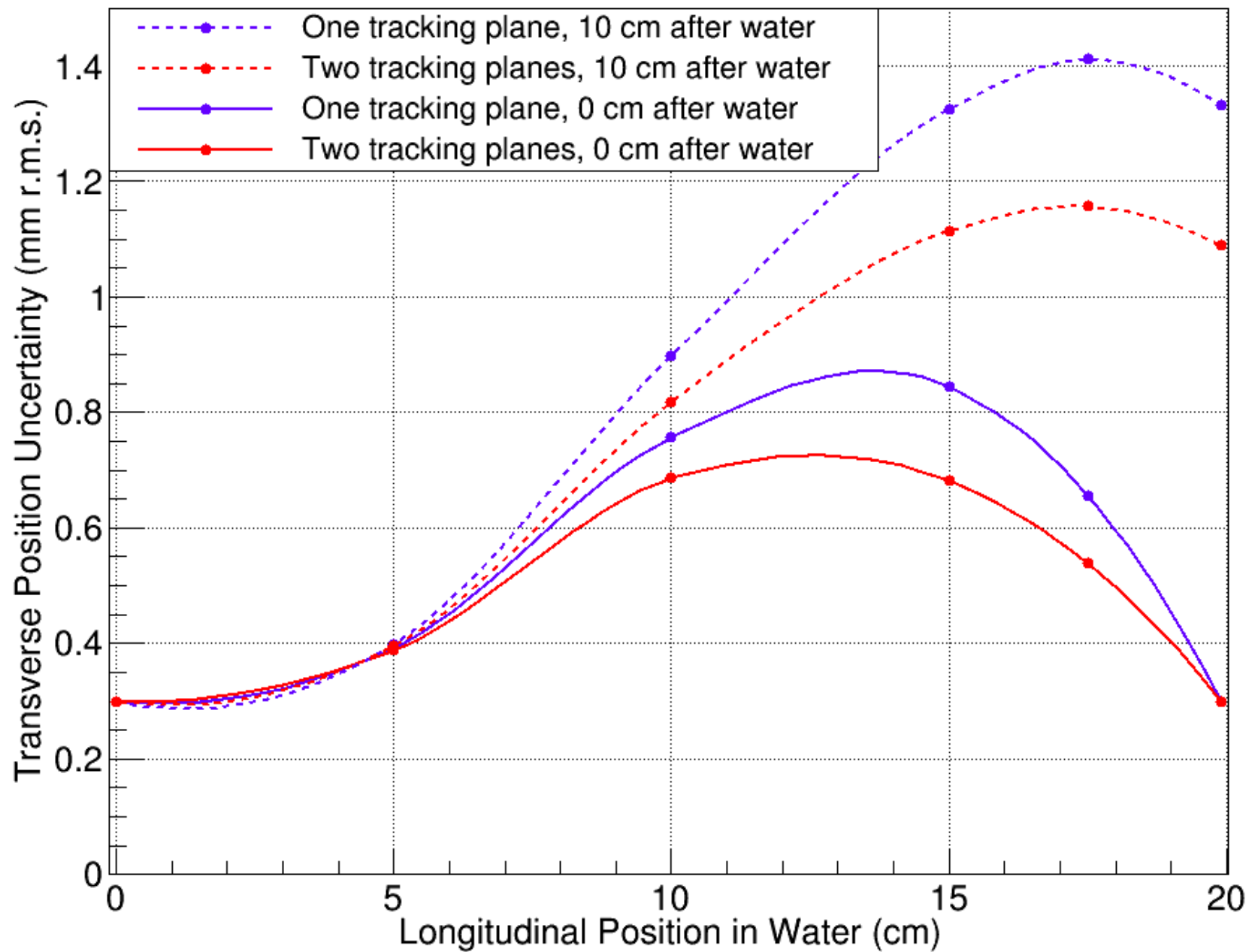


151 MeV scan through 8.1 cm block of water

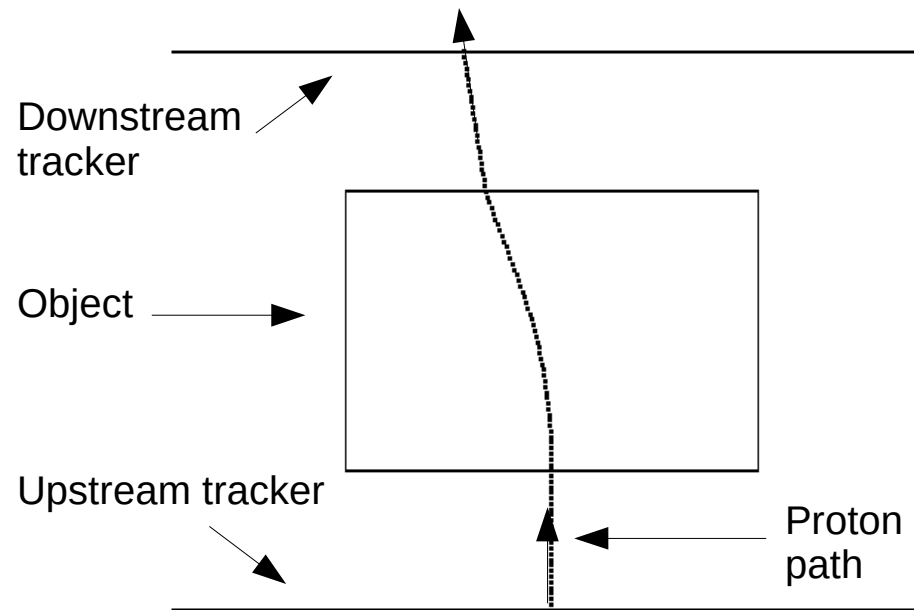




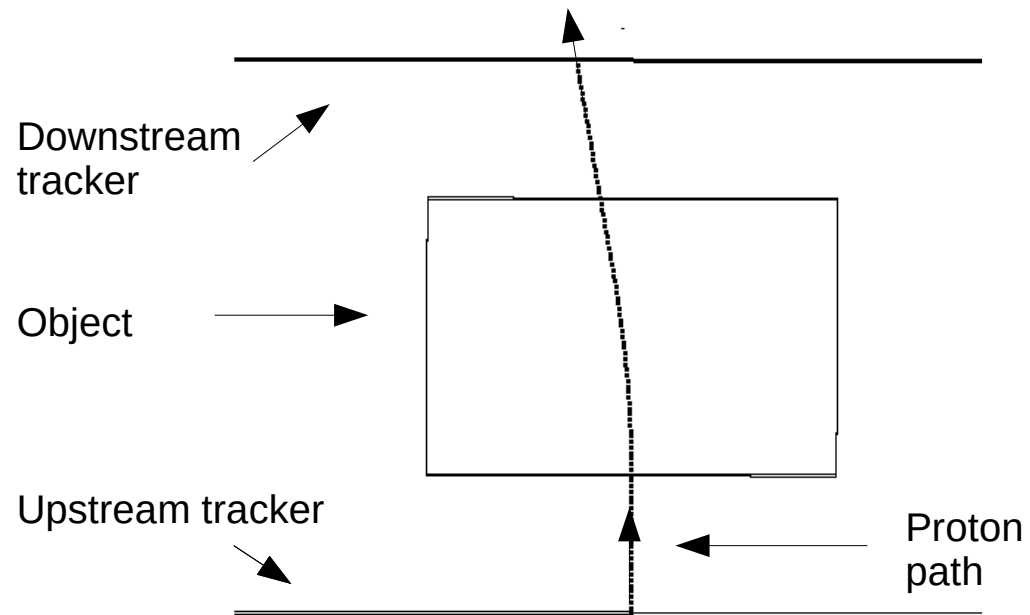
Precision of Most Likely Path Reconstruction in 20 cm Water



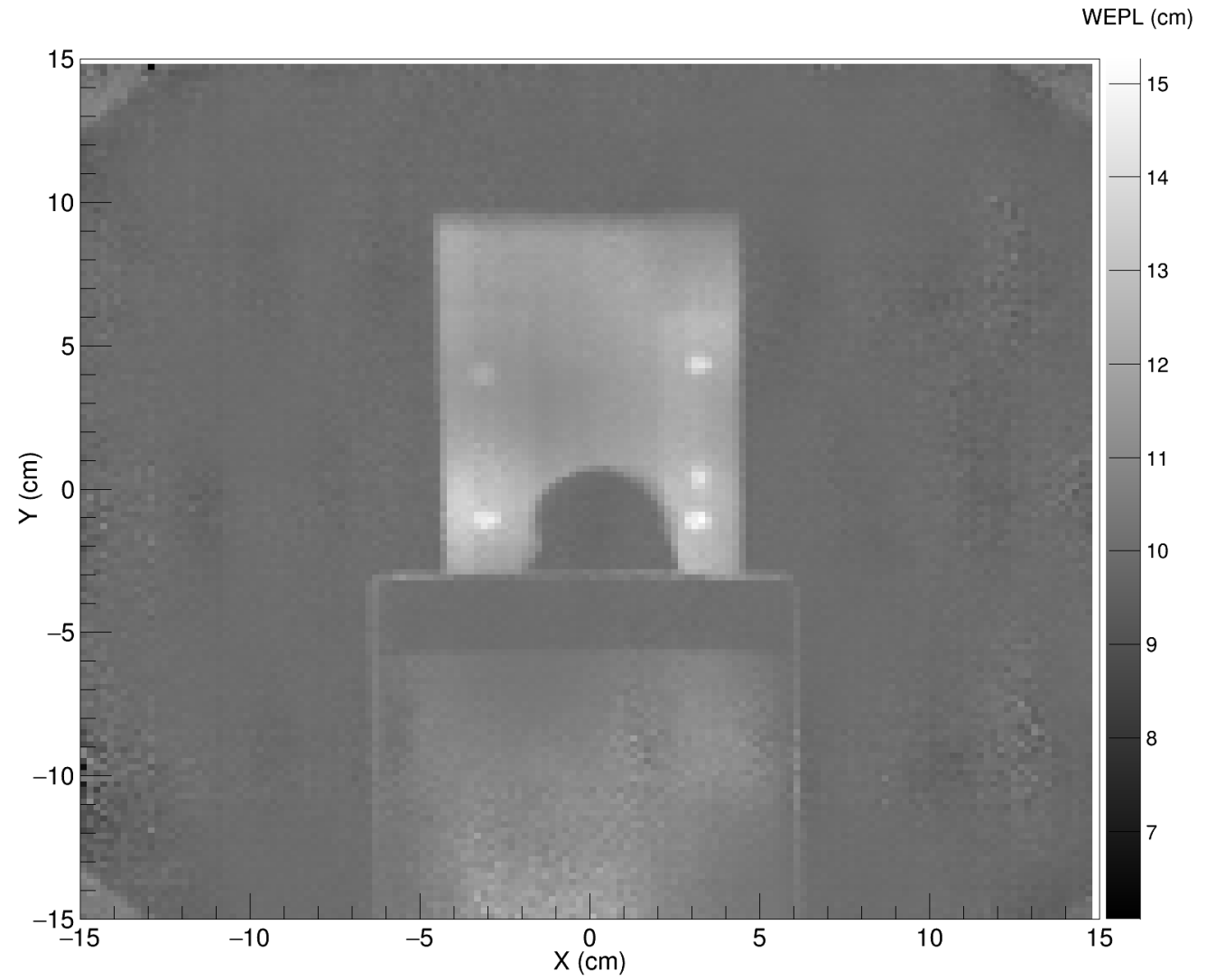
Hull exit point from straight line approximation

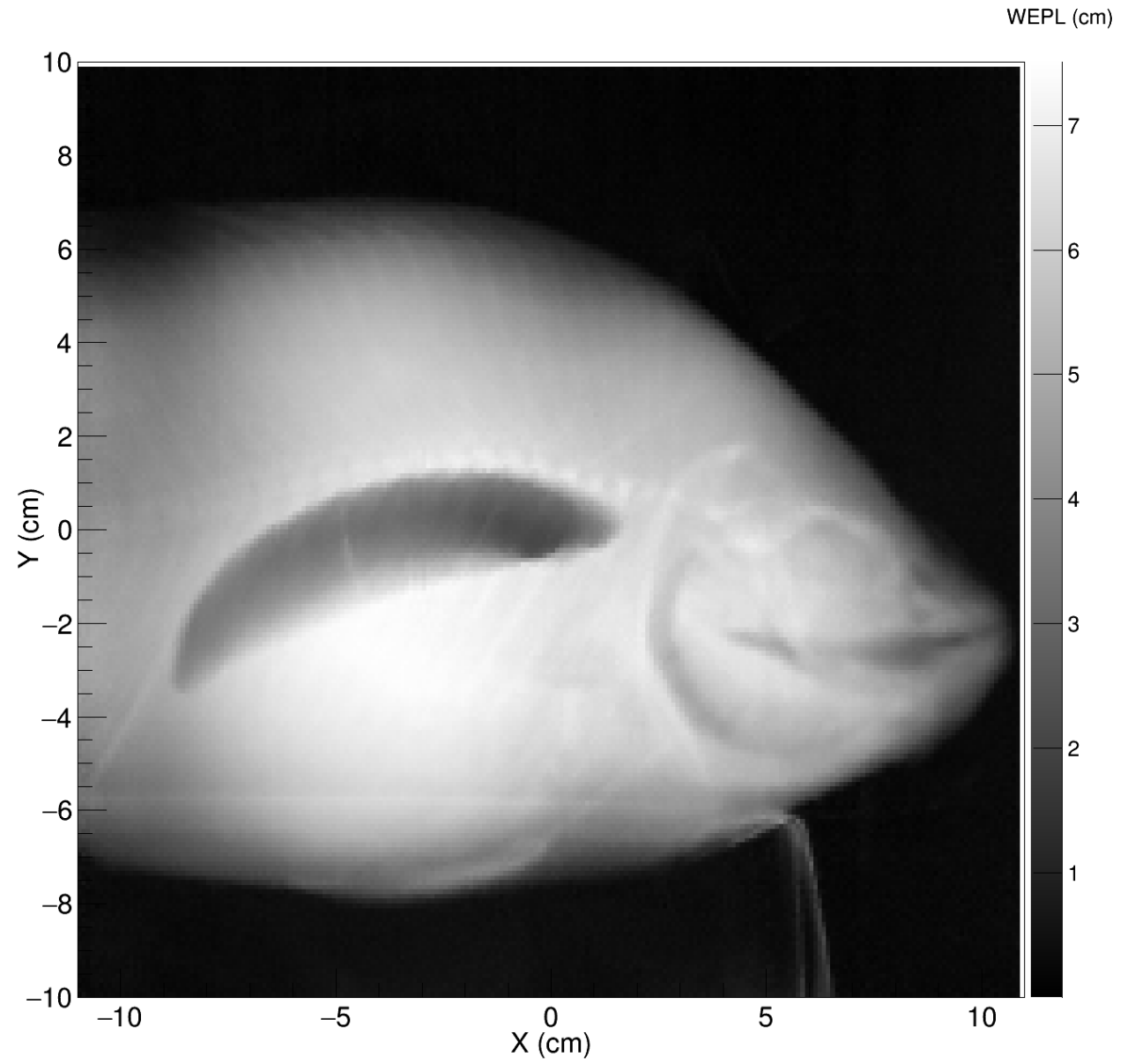


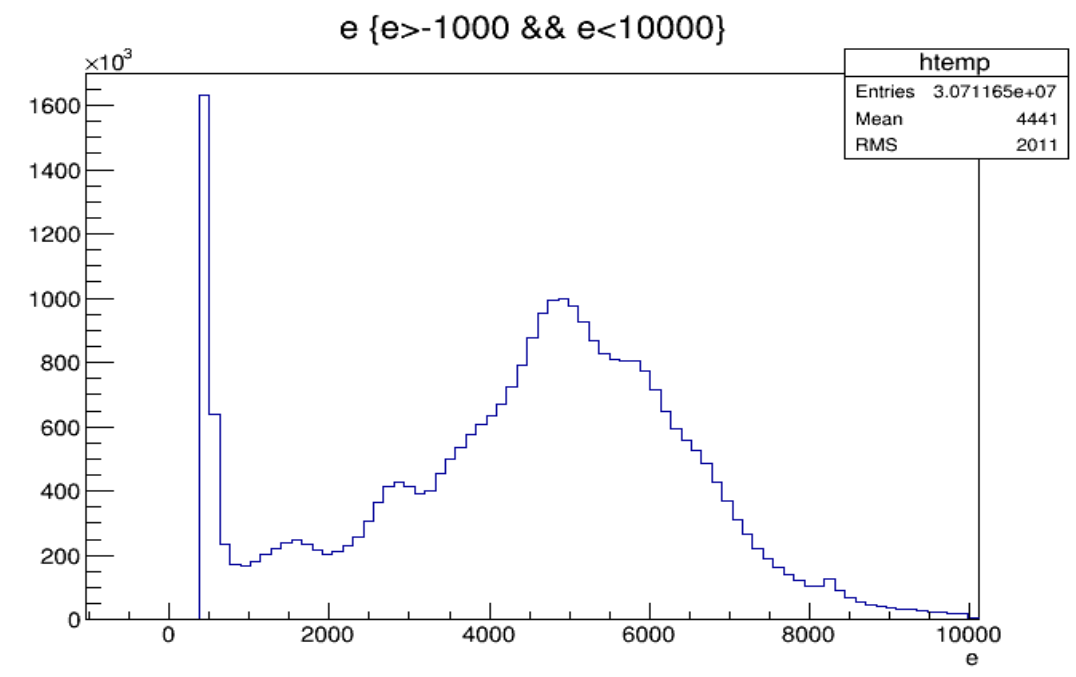
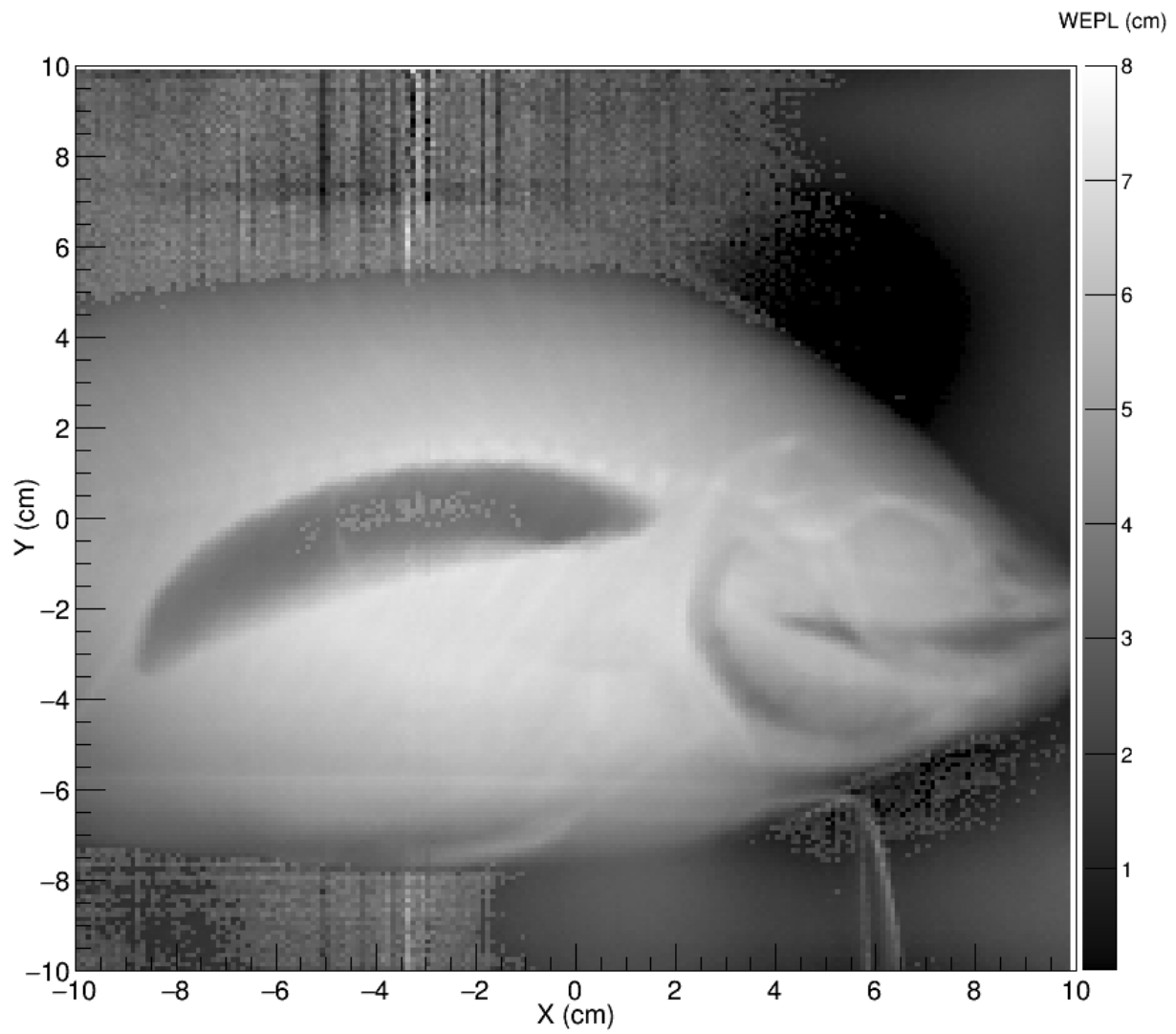
Hull exit point from Most Likely Intersection method



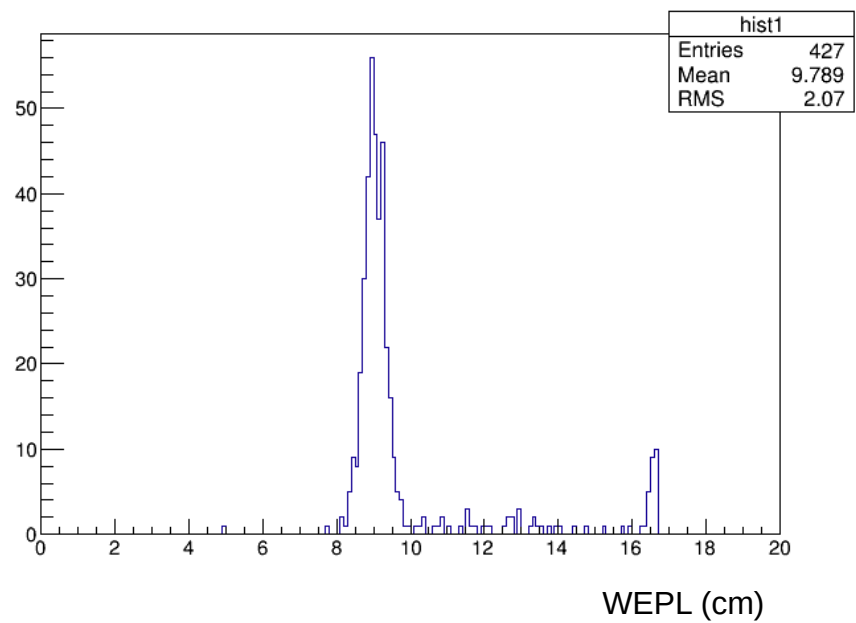
Our first real image – block of wood with screws



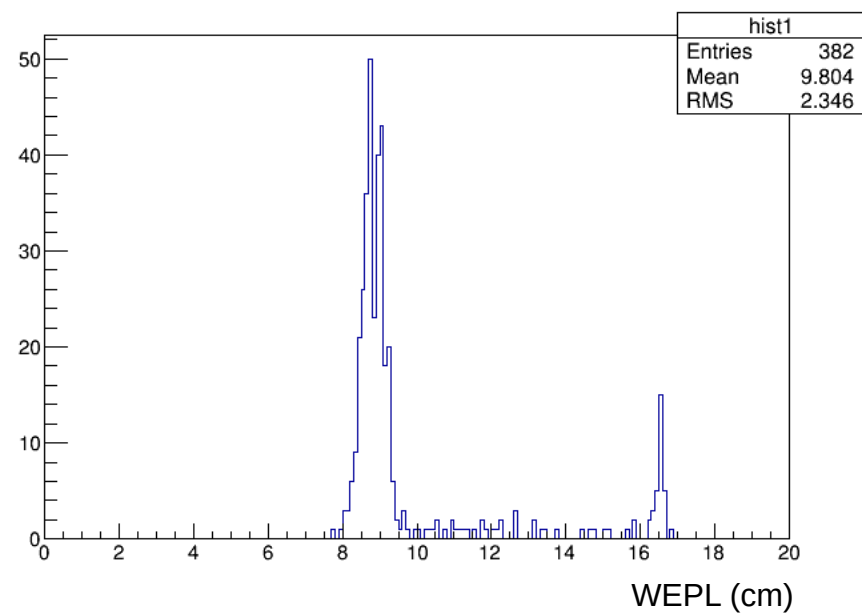




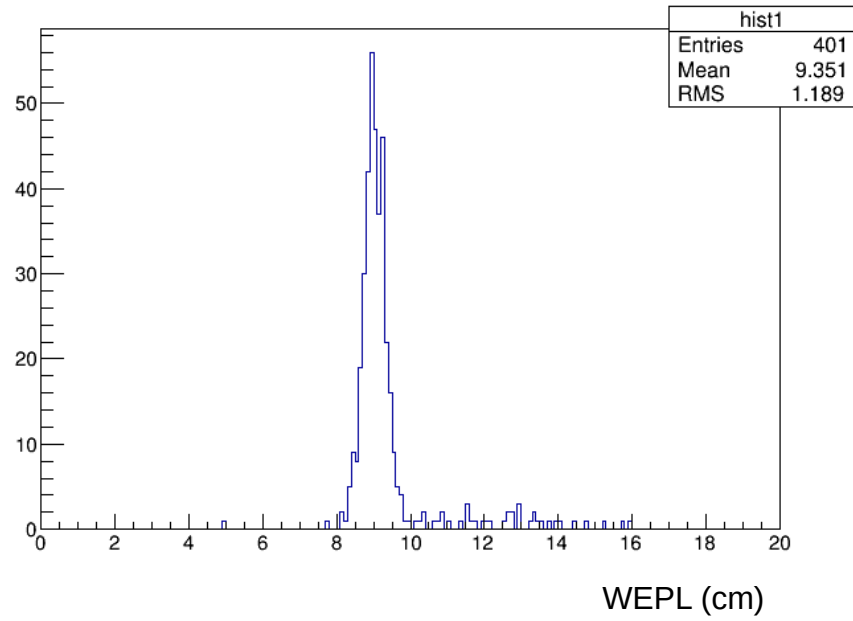
Pixel 1



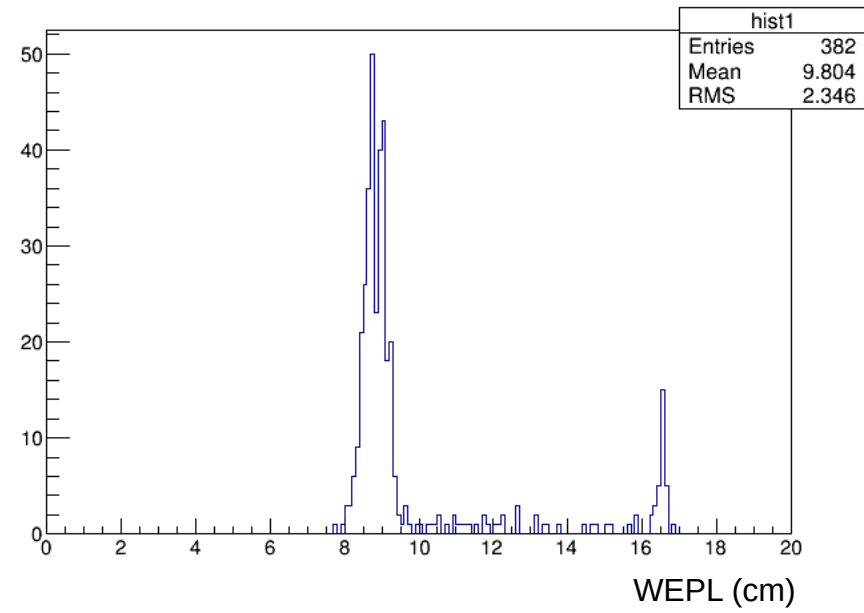
Pixel 2



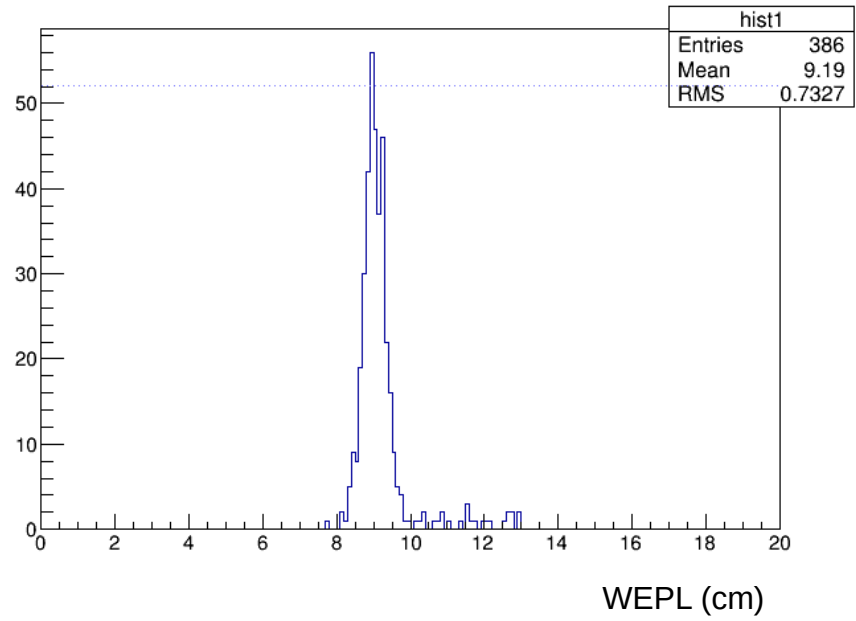
Pixel 1



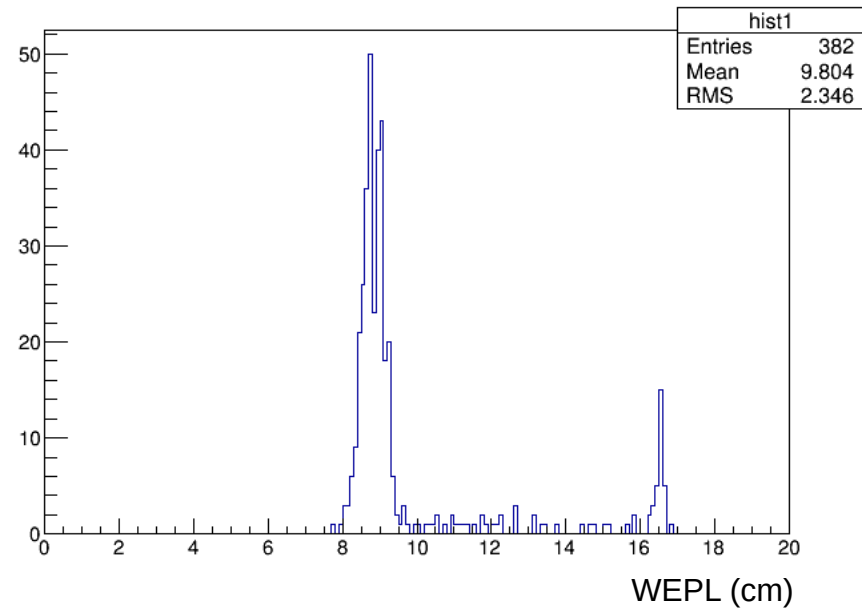
Pixel 2



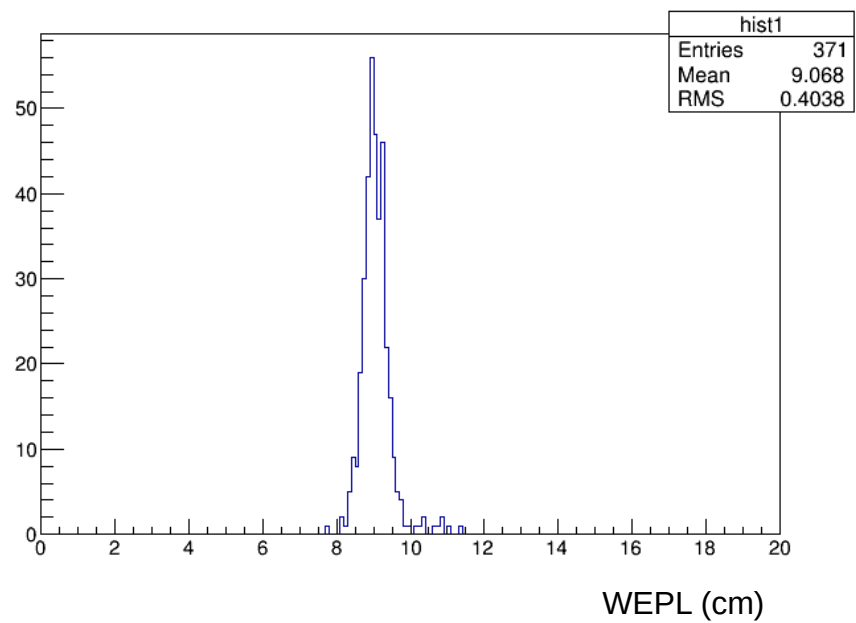
Pixel 1



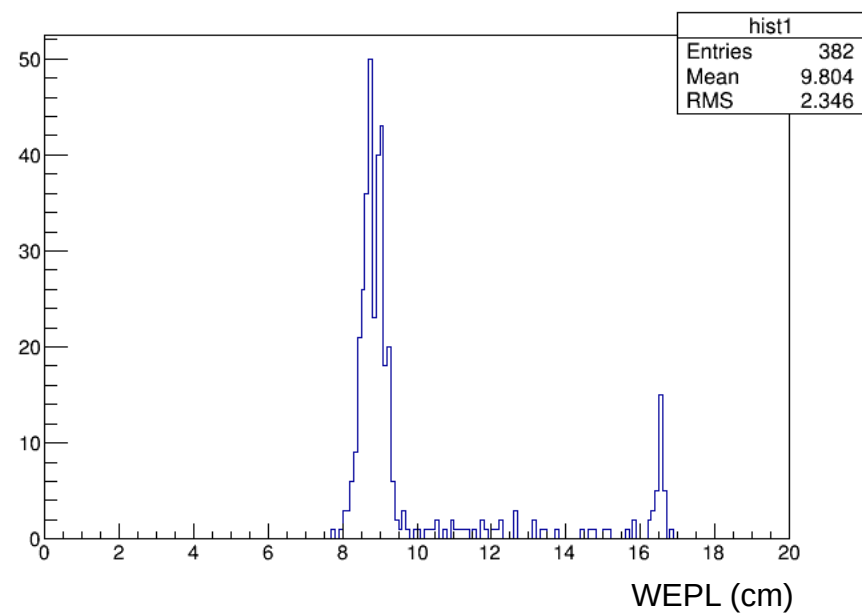
Pixel 2



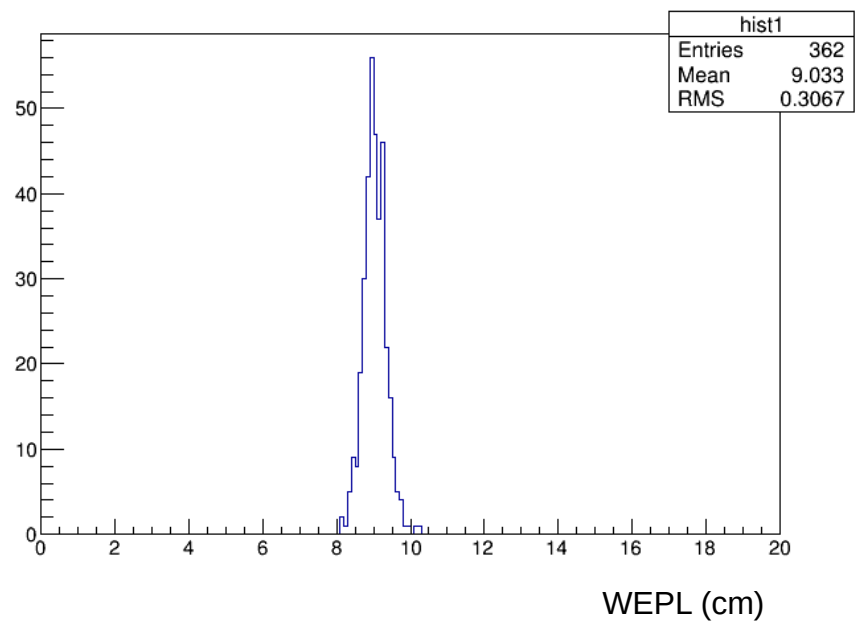
Pixel 1



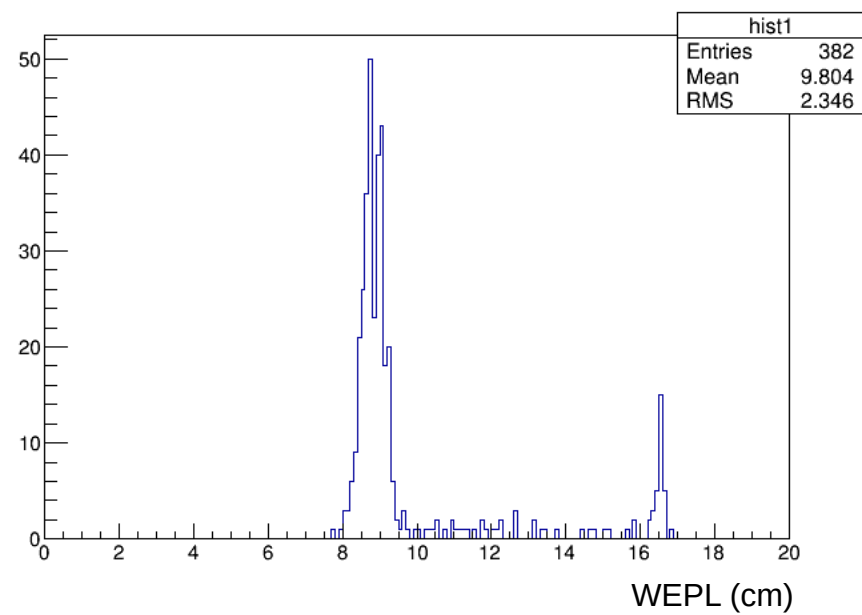
Pixel 2



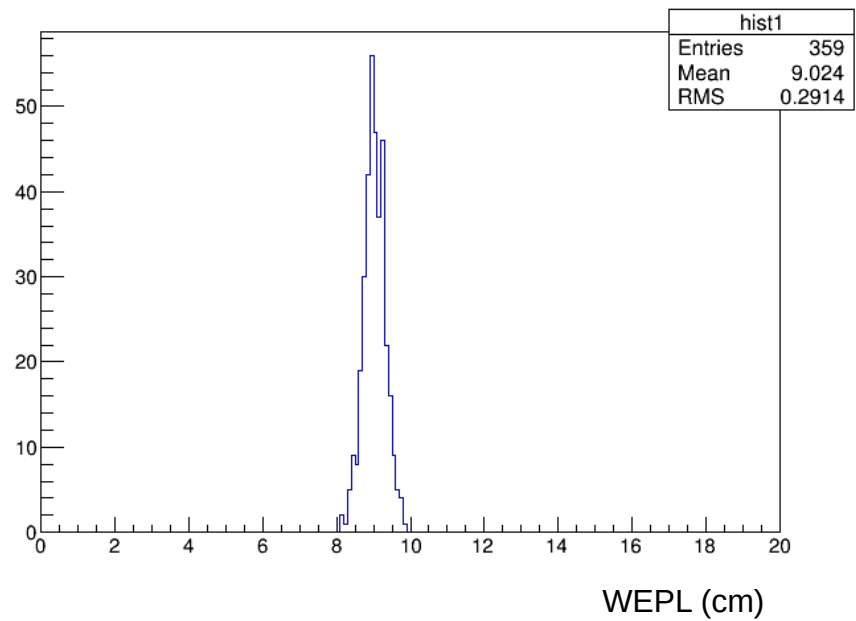
Pixel 1



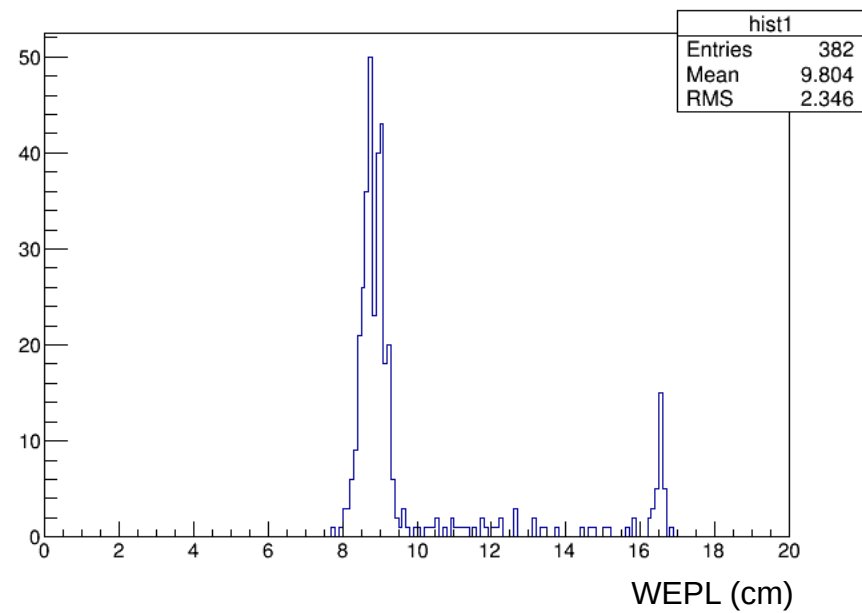
Pixel 2



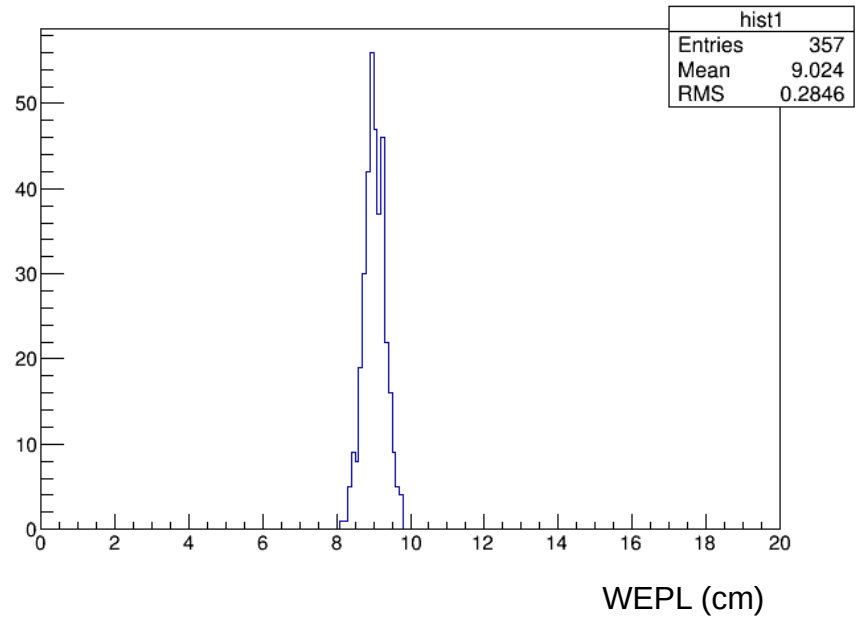
Pixel 1



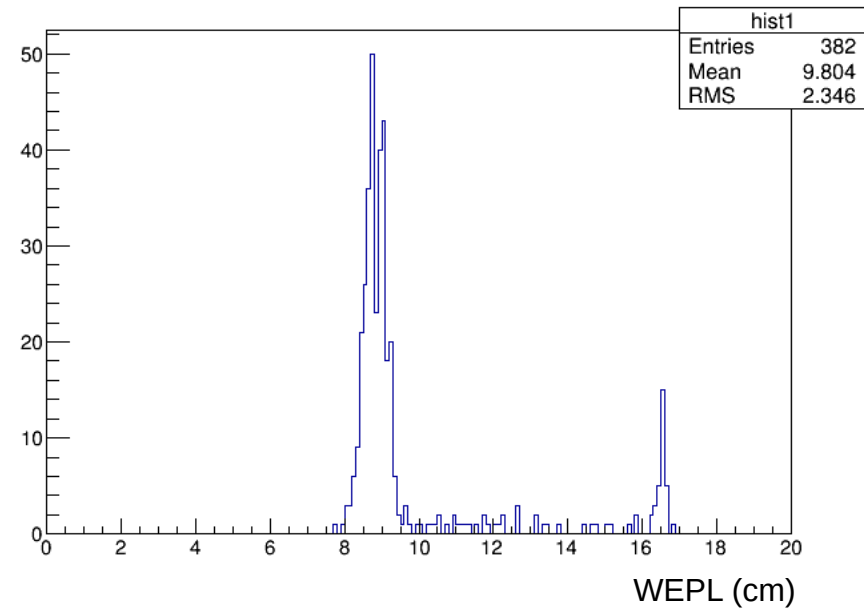
Pixel 2



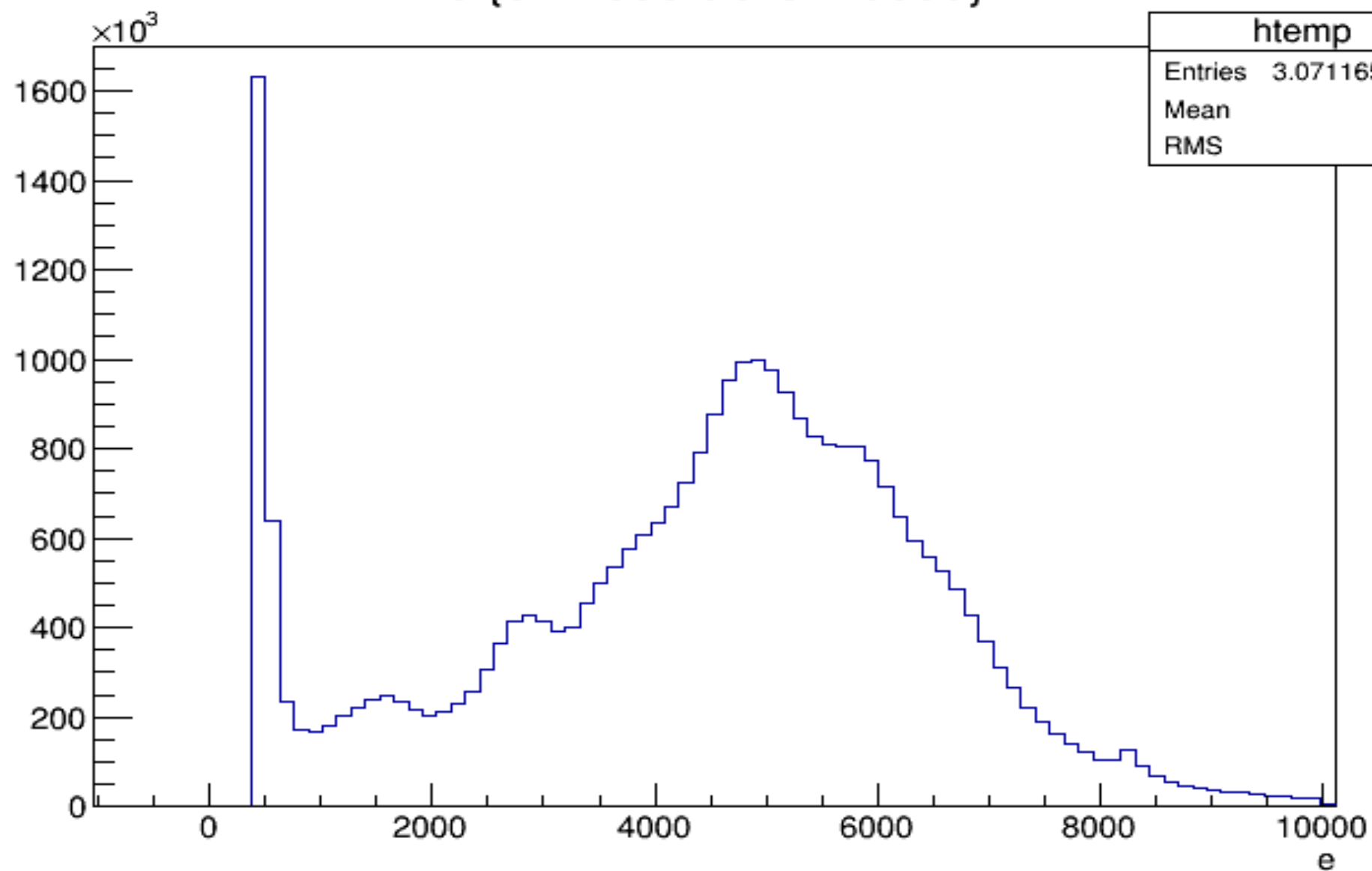
Pixel 1



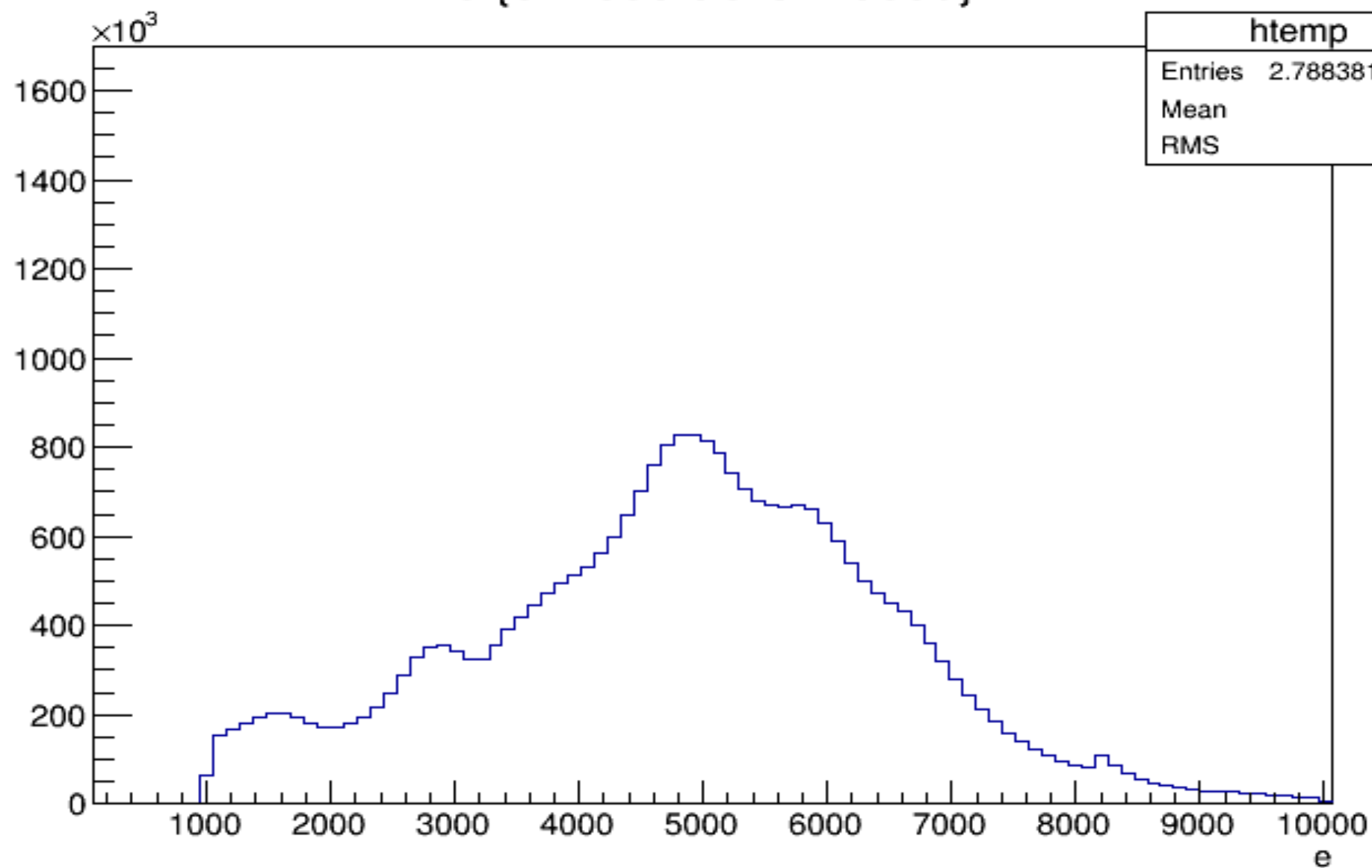
Pixel 2



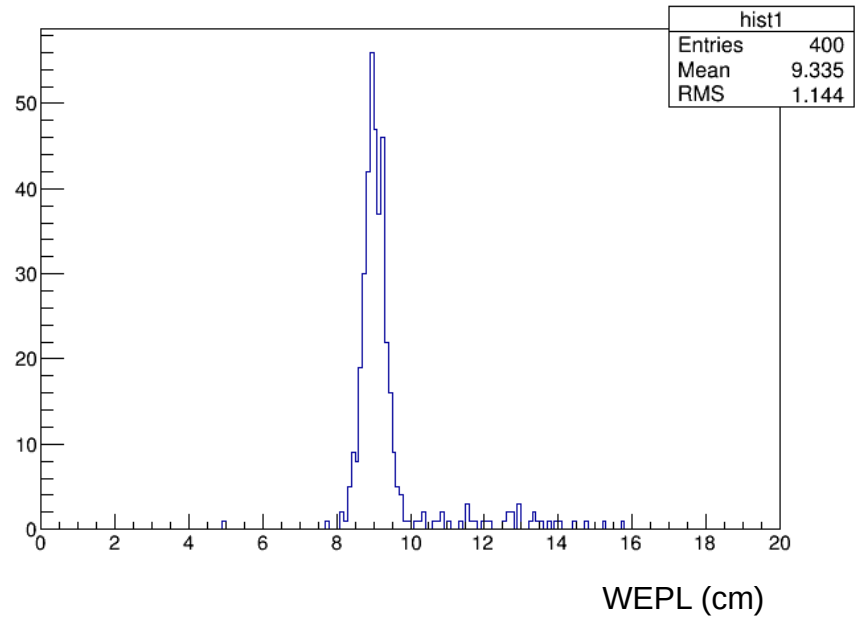
e {e>-1000 && e<10000}



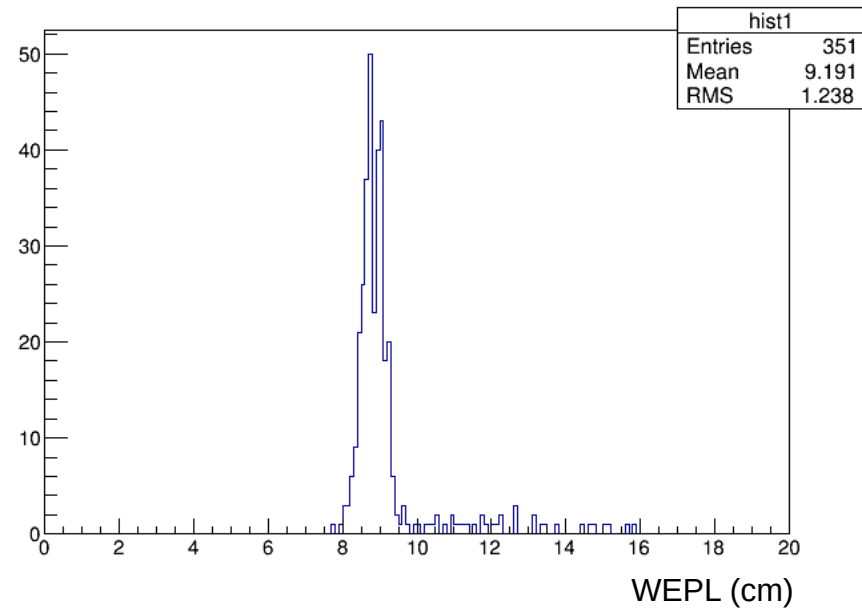
e {e>1000 && e<10000}



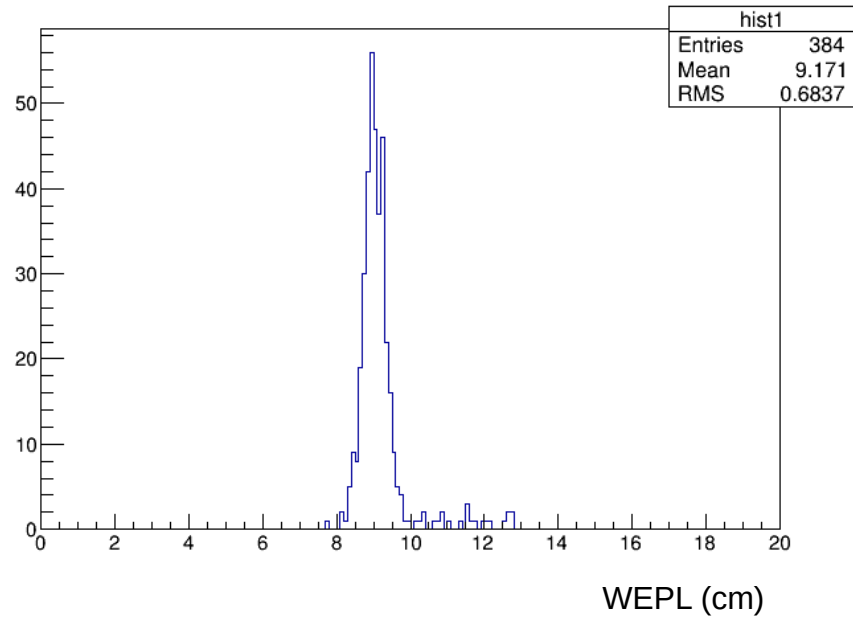
Pixel 1



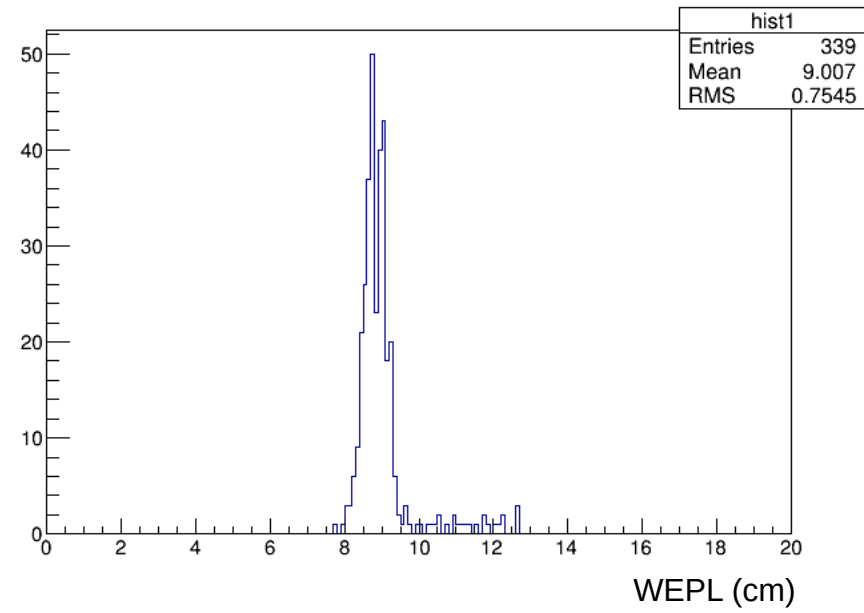
Pixel 2



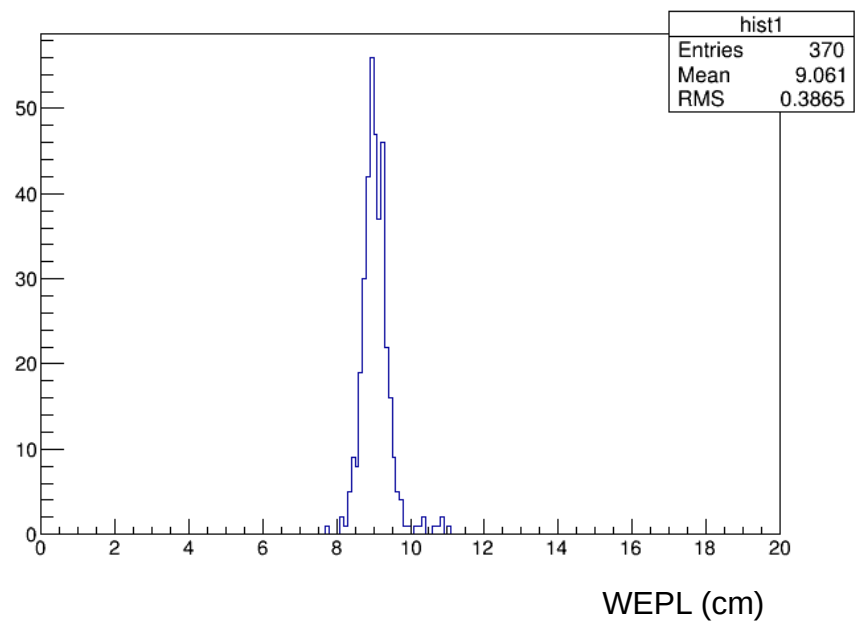
Pixel 1



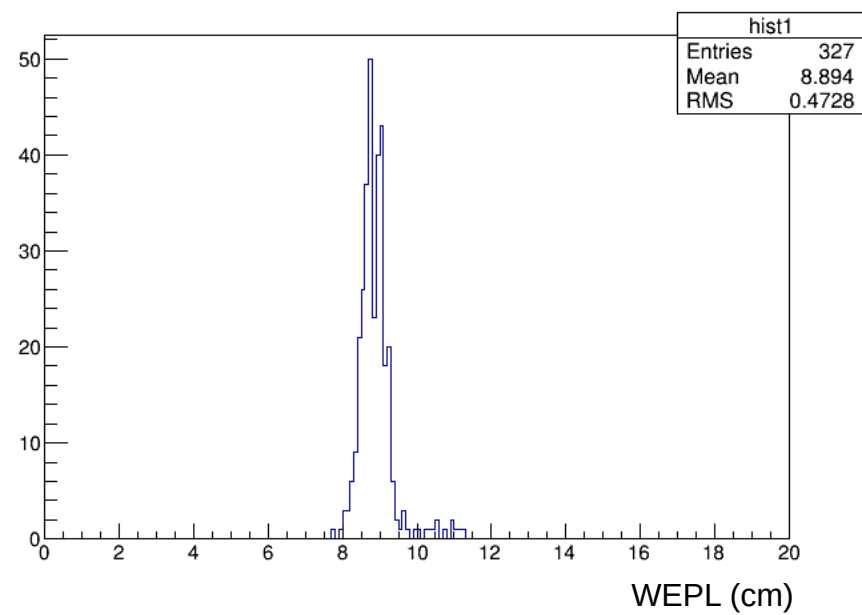
Pixel 2



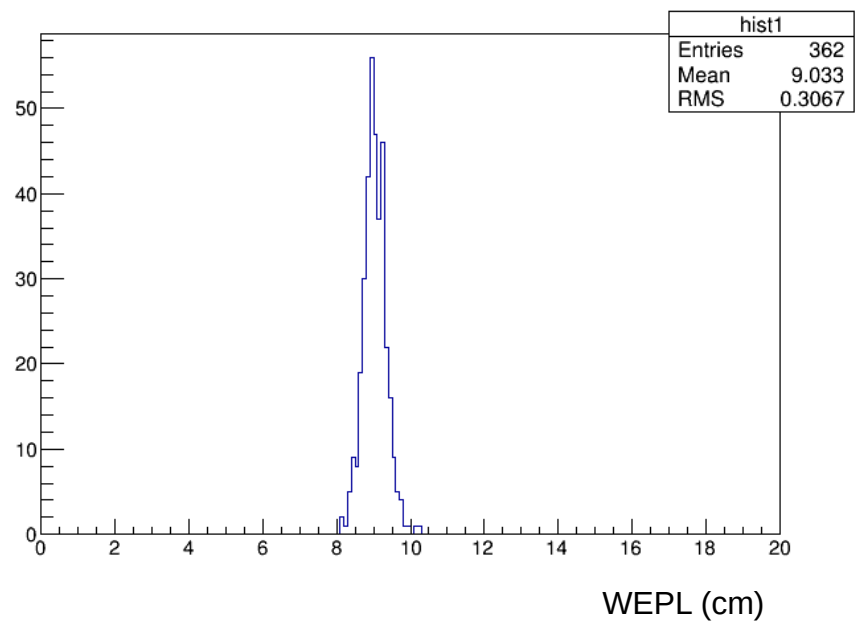
Pixel 1



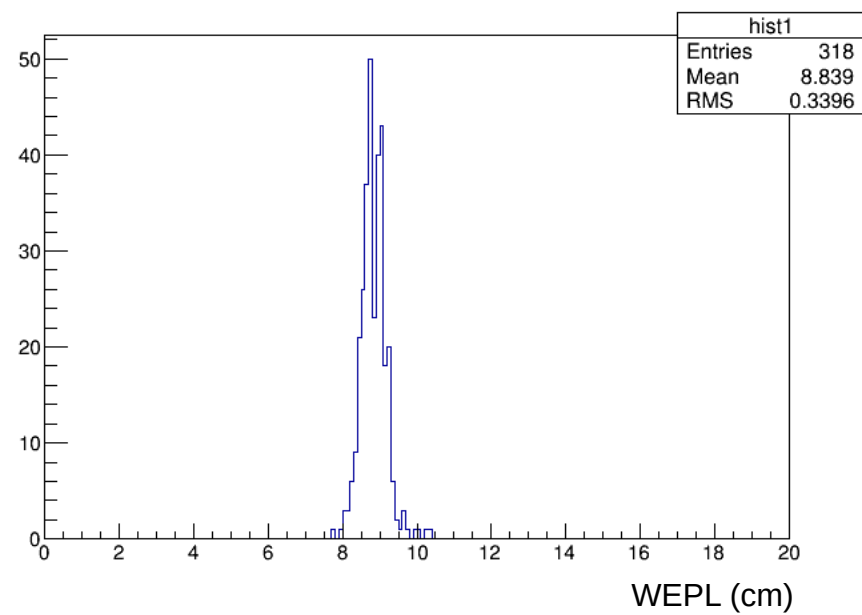
Pixel 2



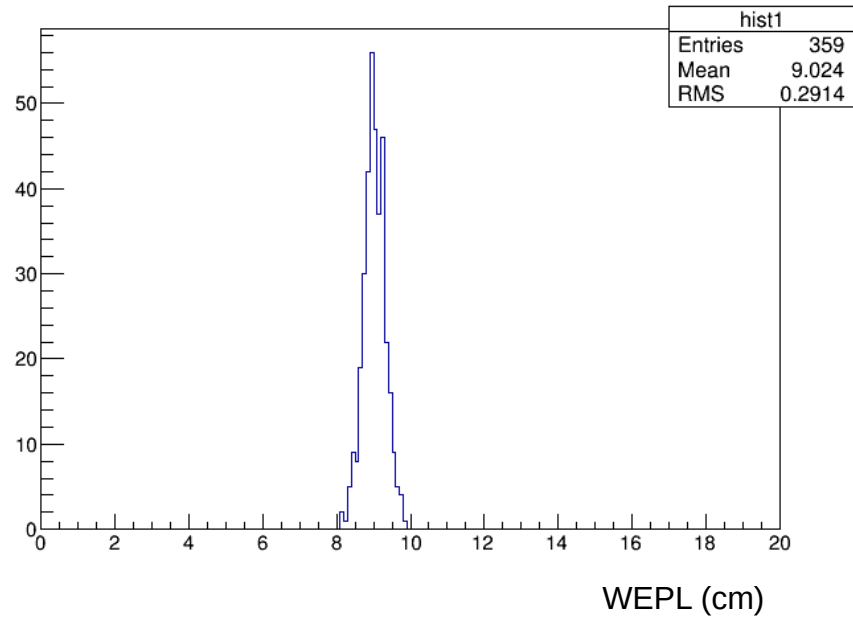
Pixel 1



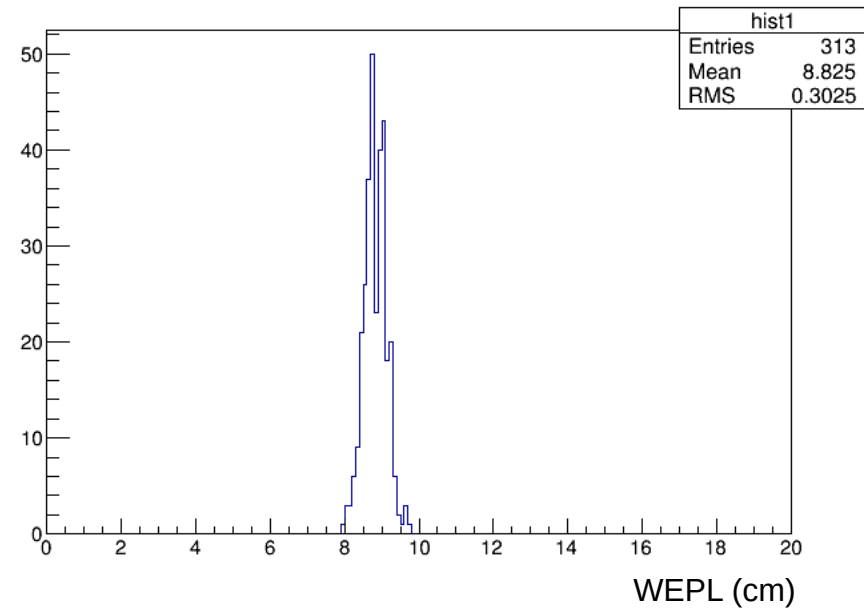
Pixel 2



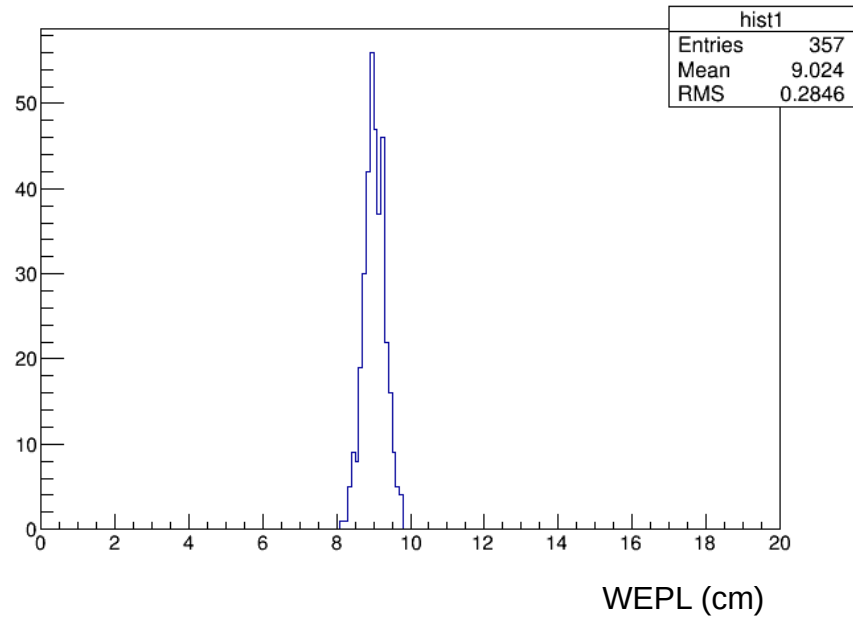
Pixel 1



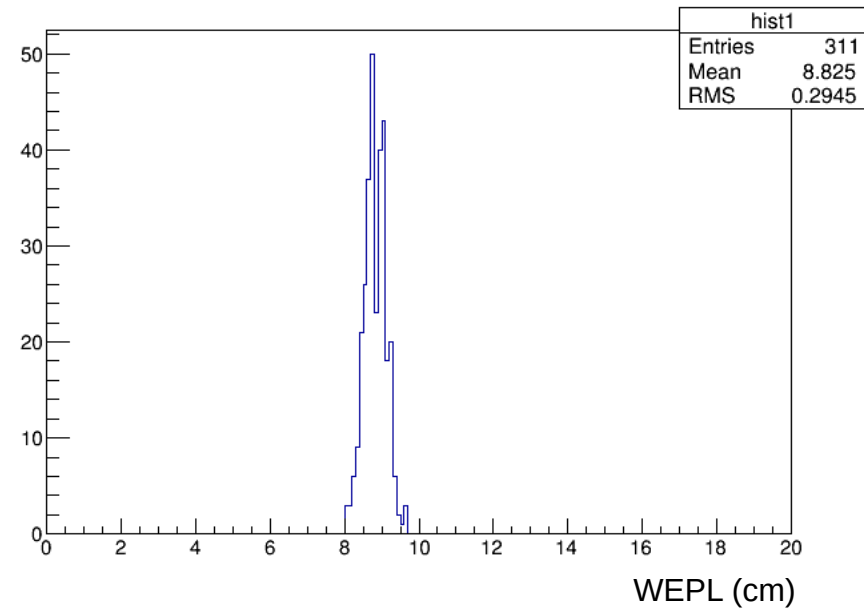
Pixel 2

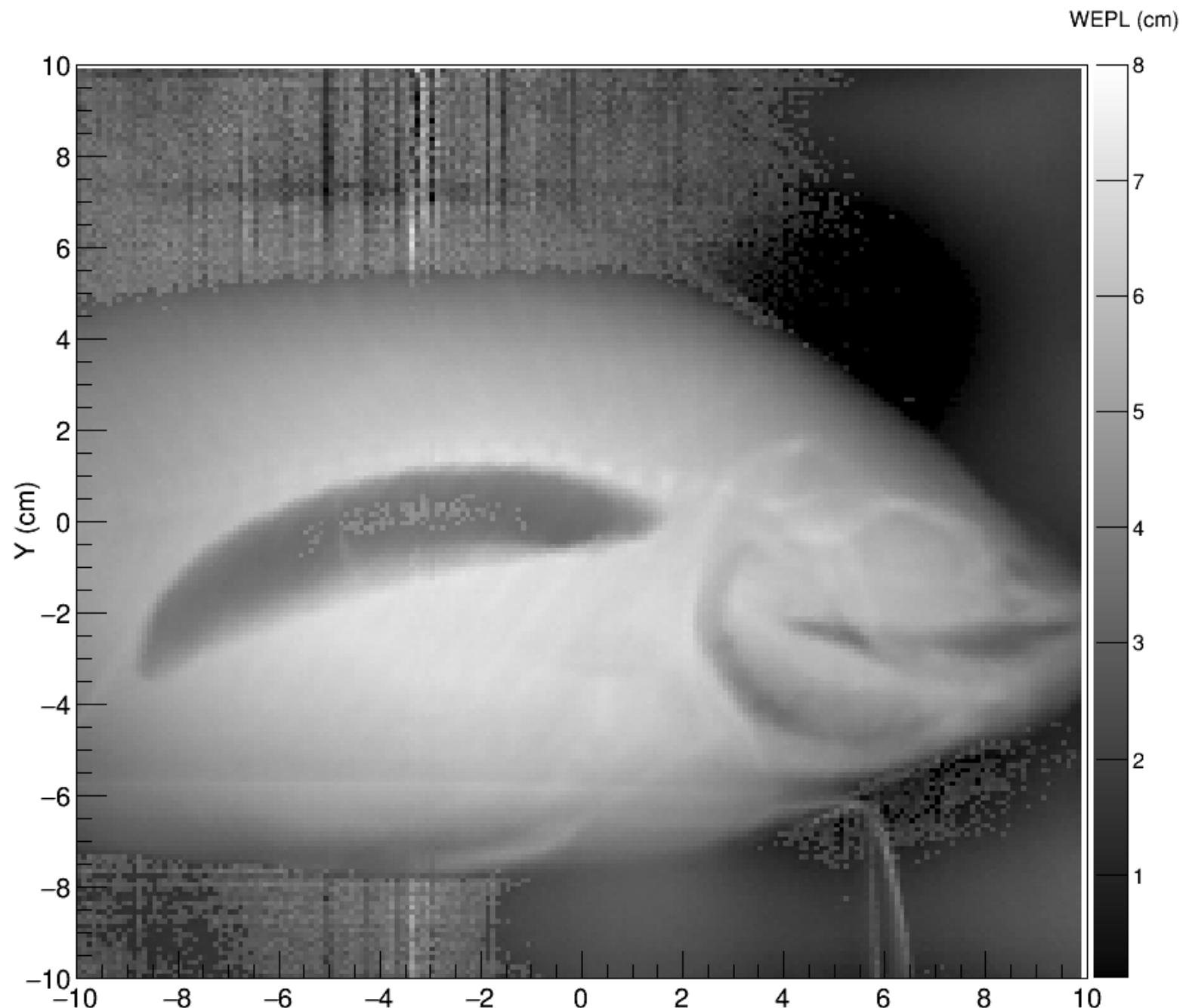


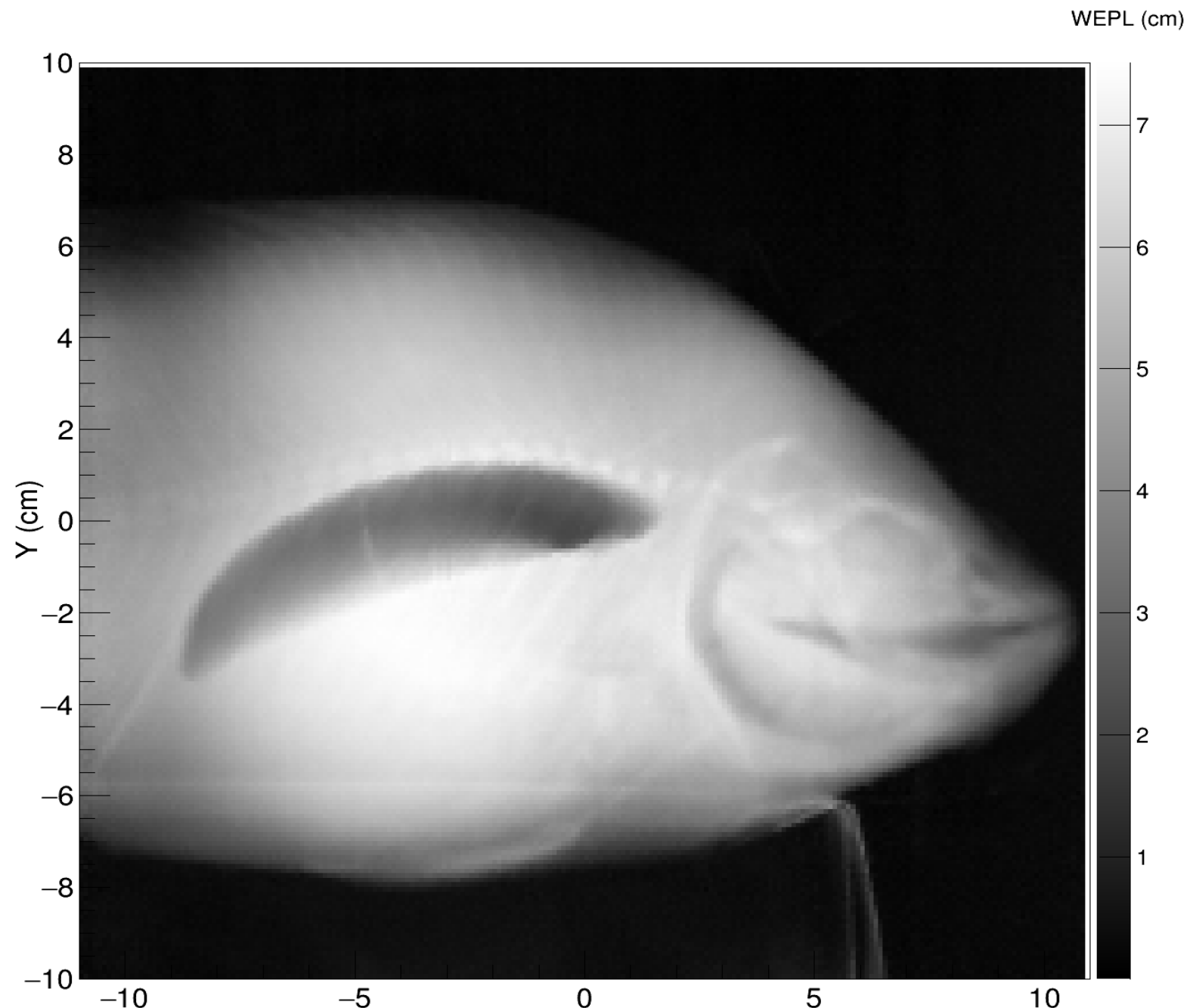
Pixel 1

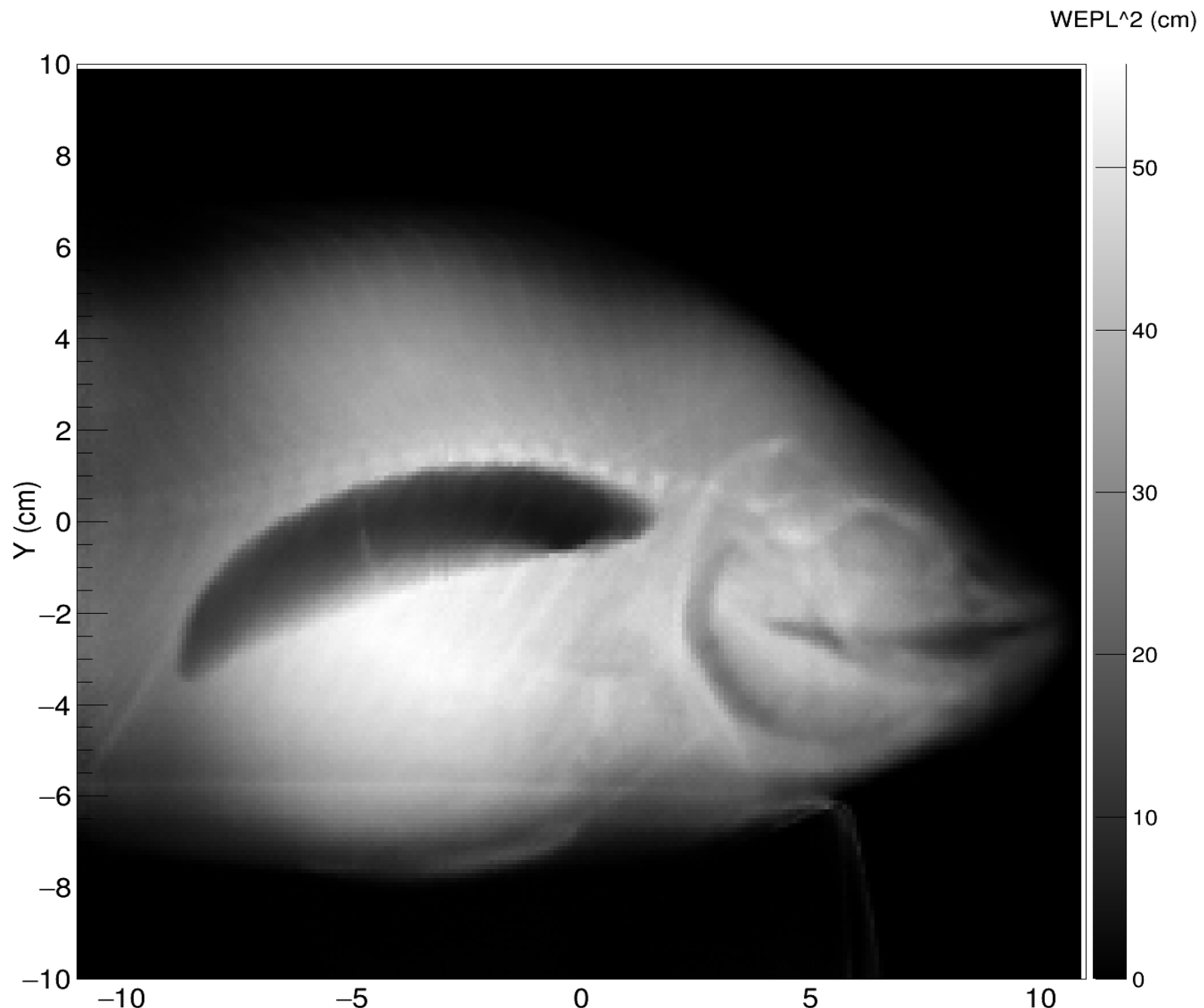


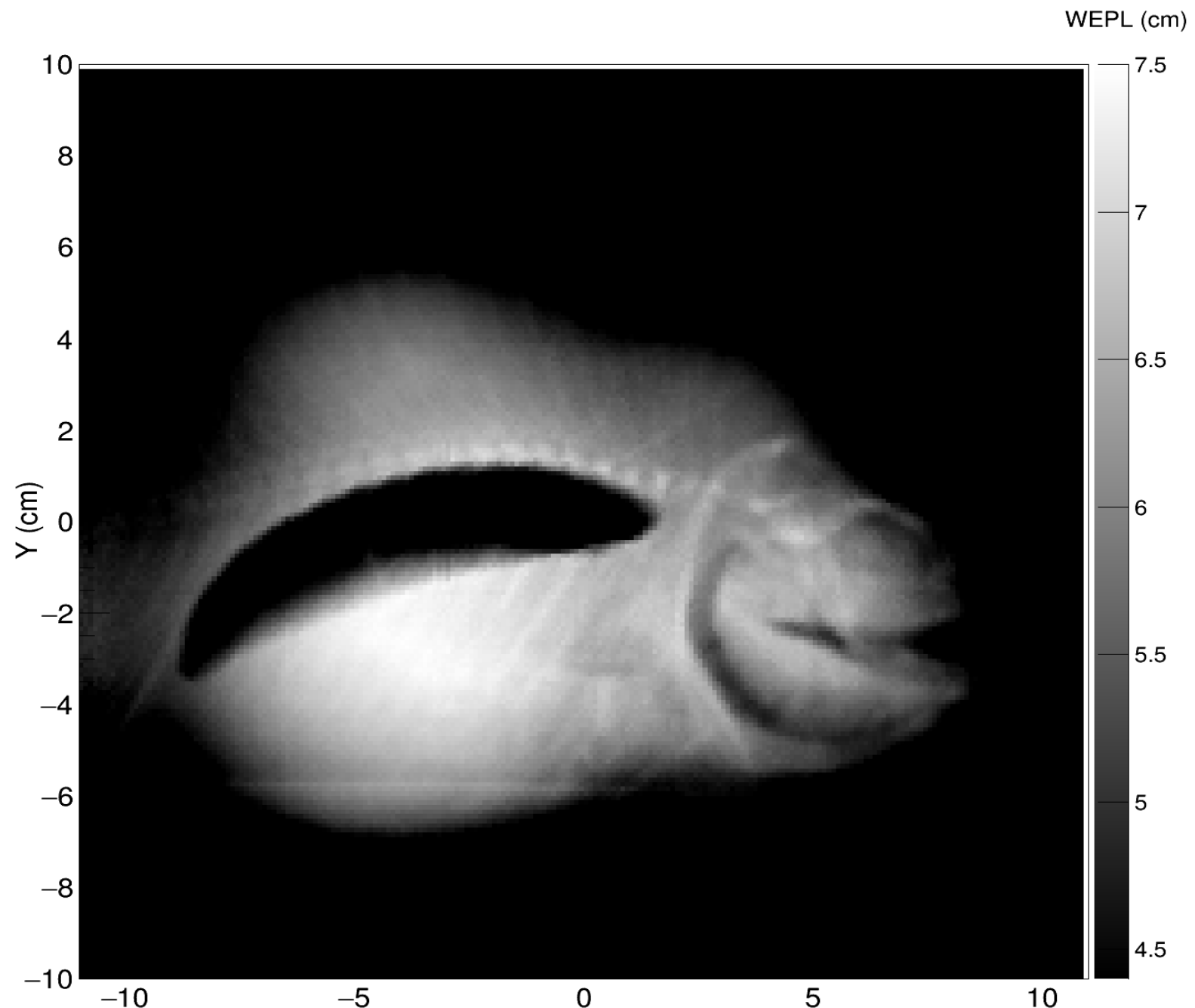
Pixel 2









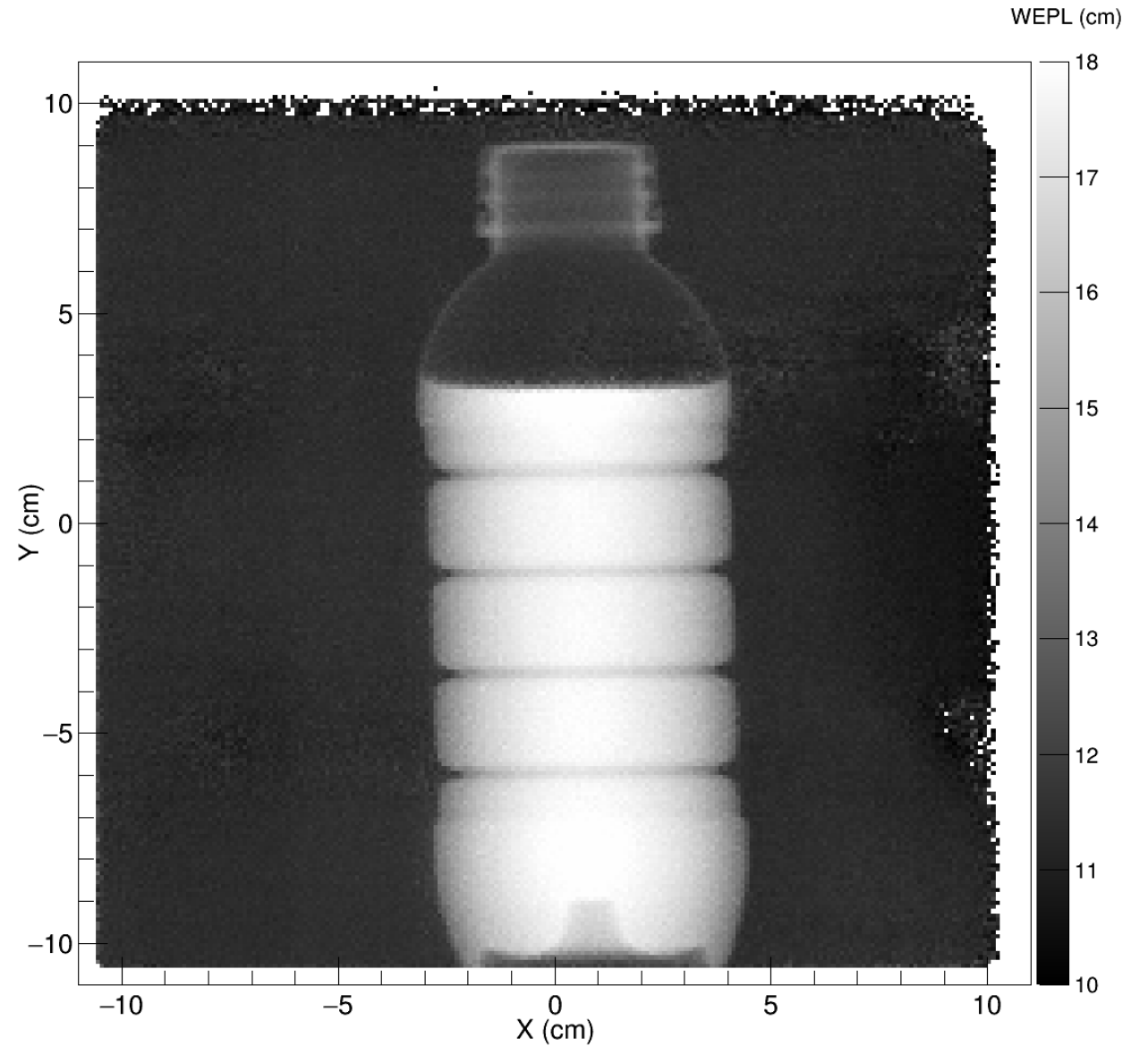


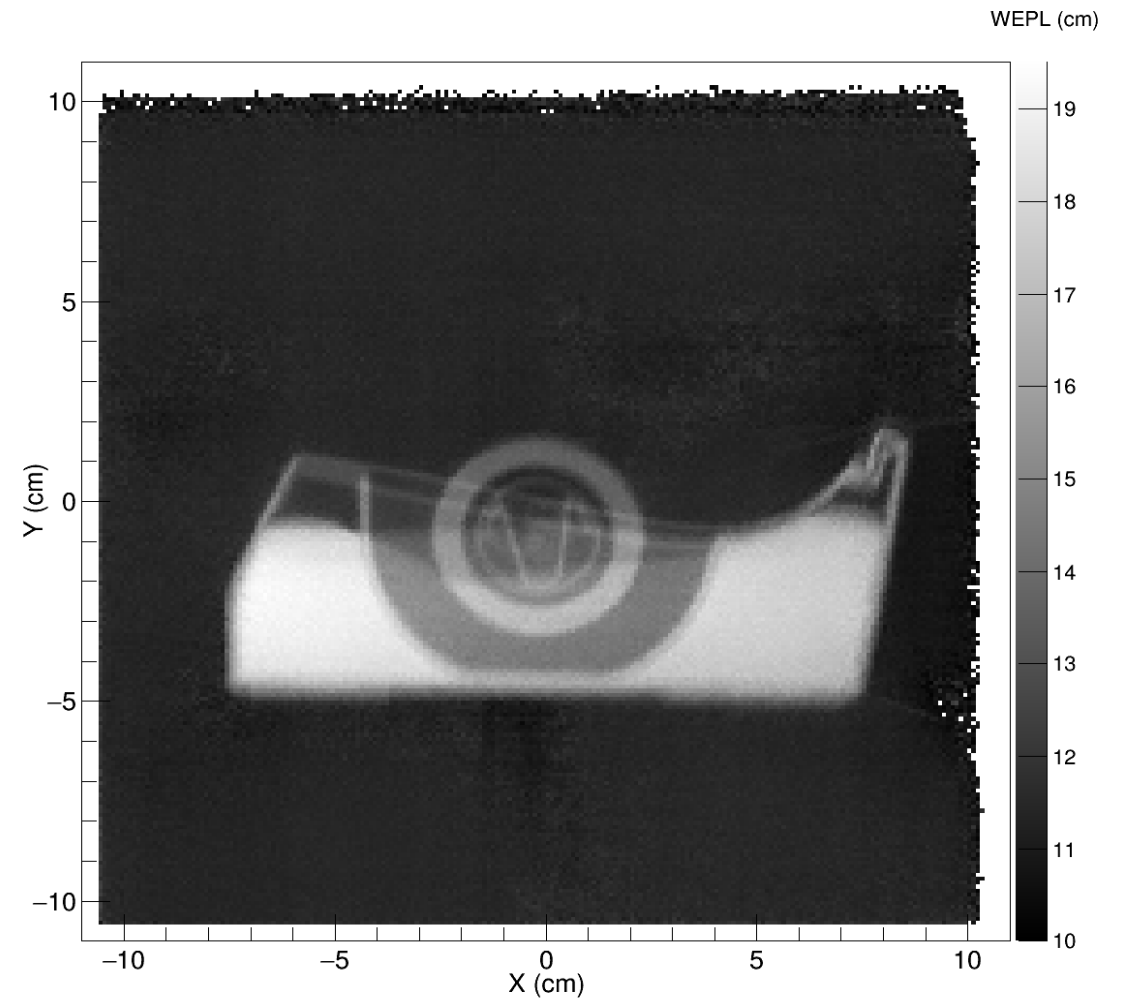
Test with ProtonVDA pRad system and ProNova Knoxville system

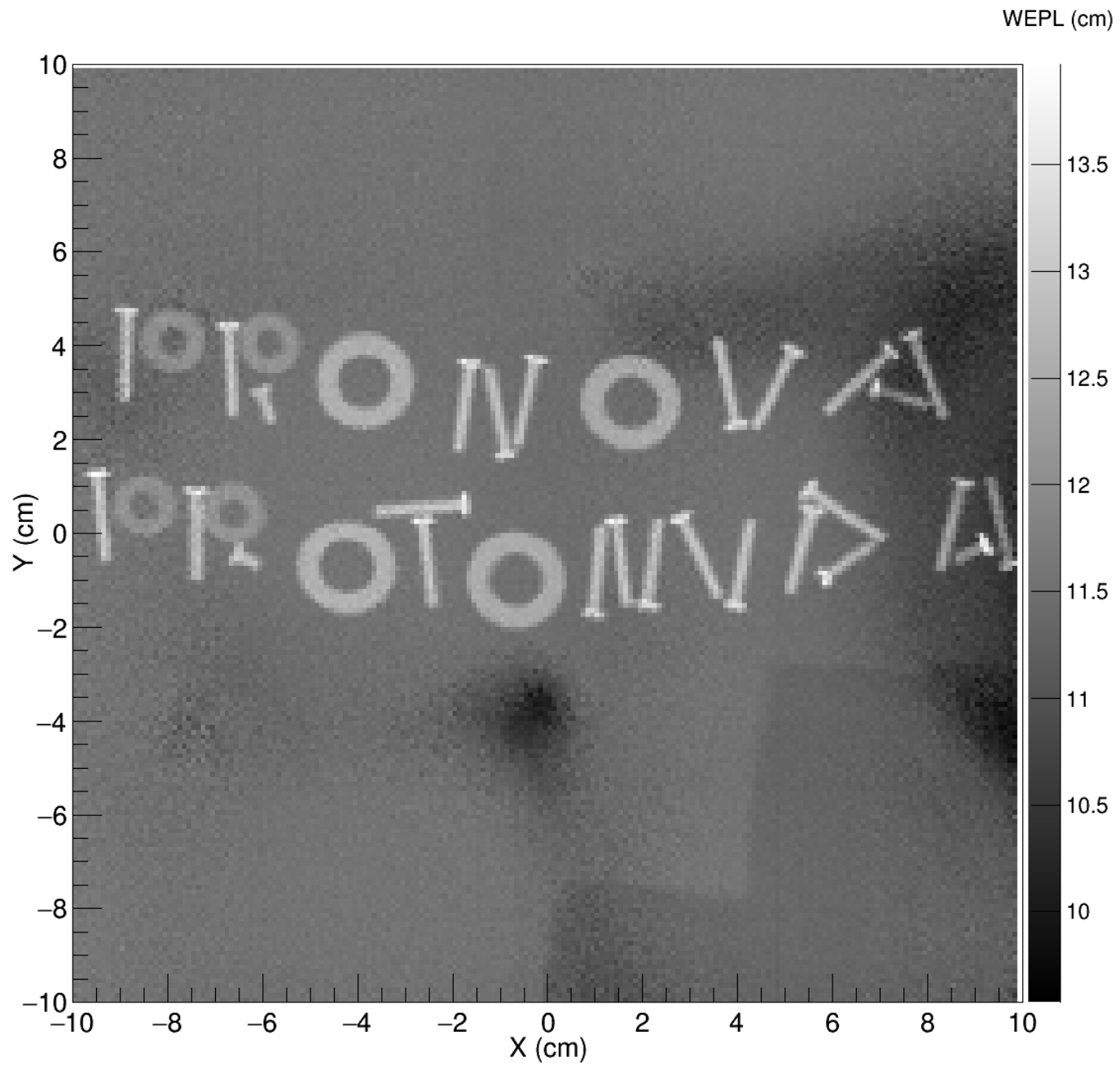
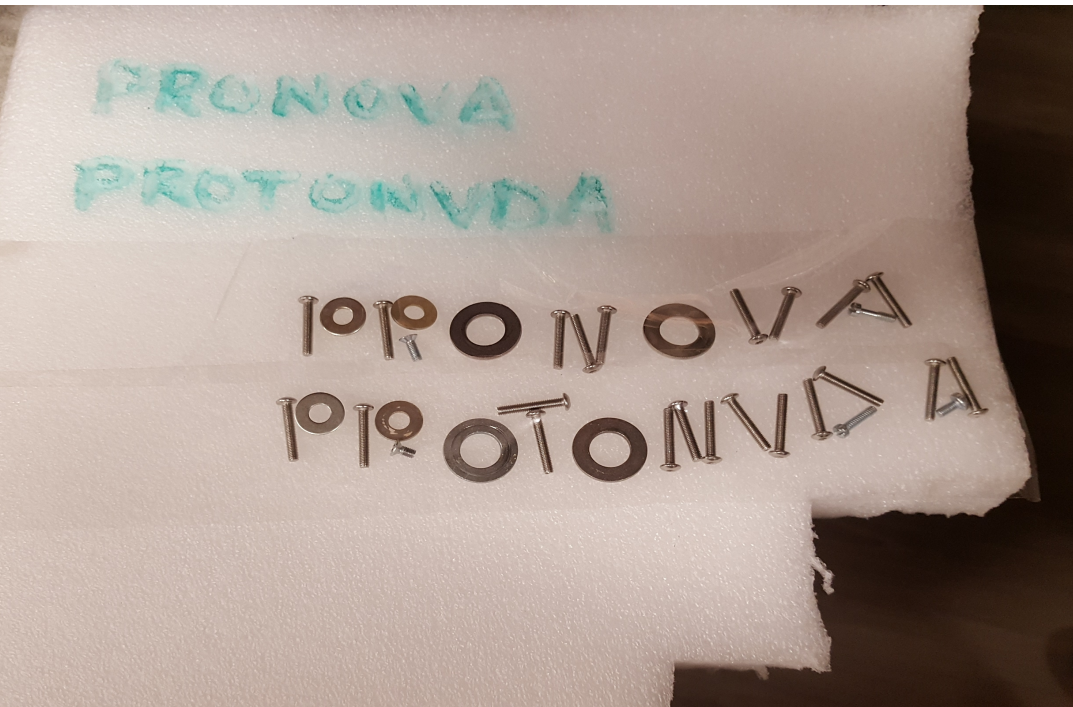
July 26 and 27 2018



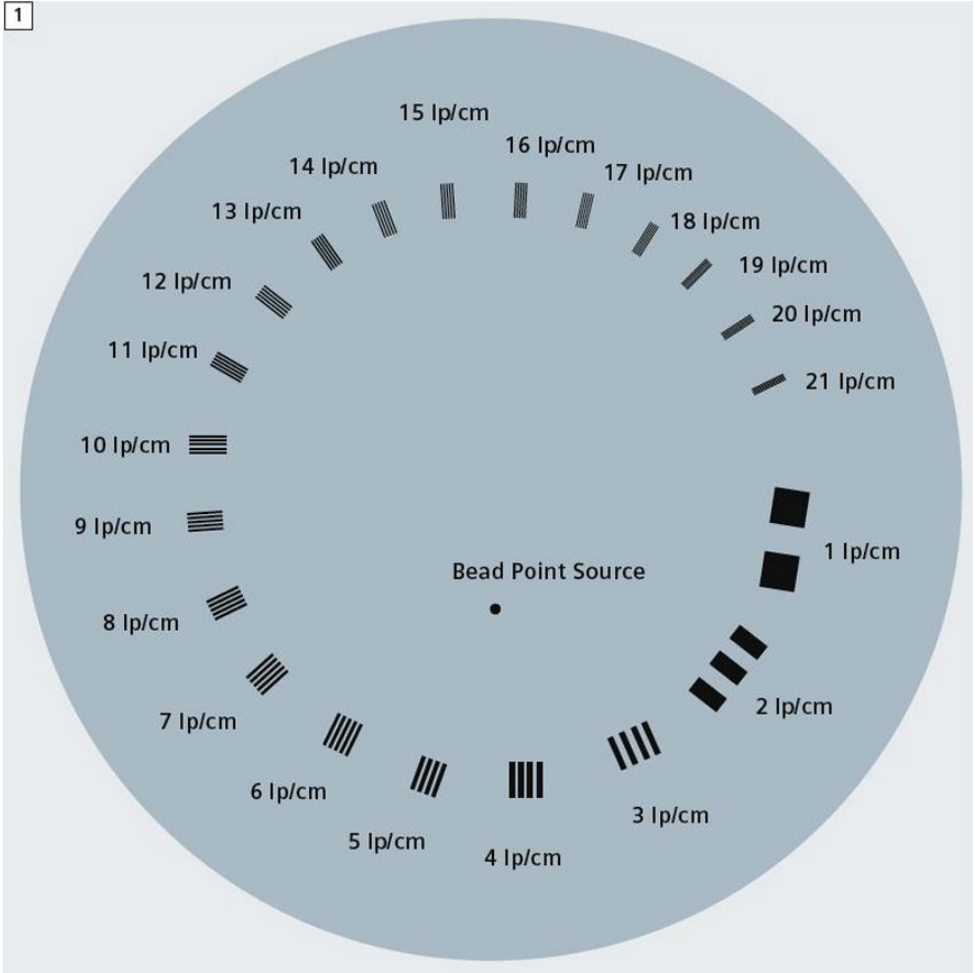
Fast (~ 1 min) online image reconstruction







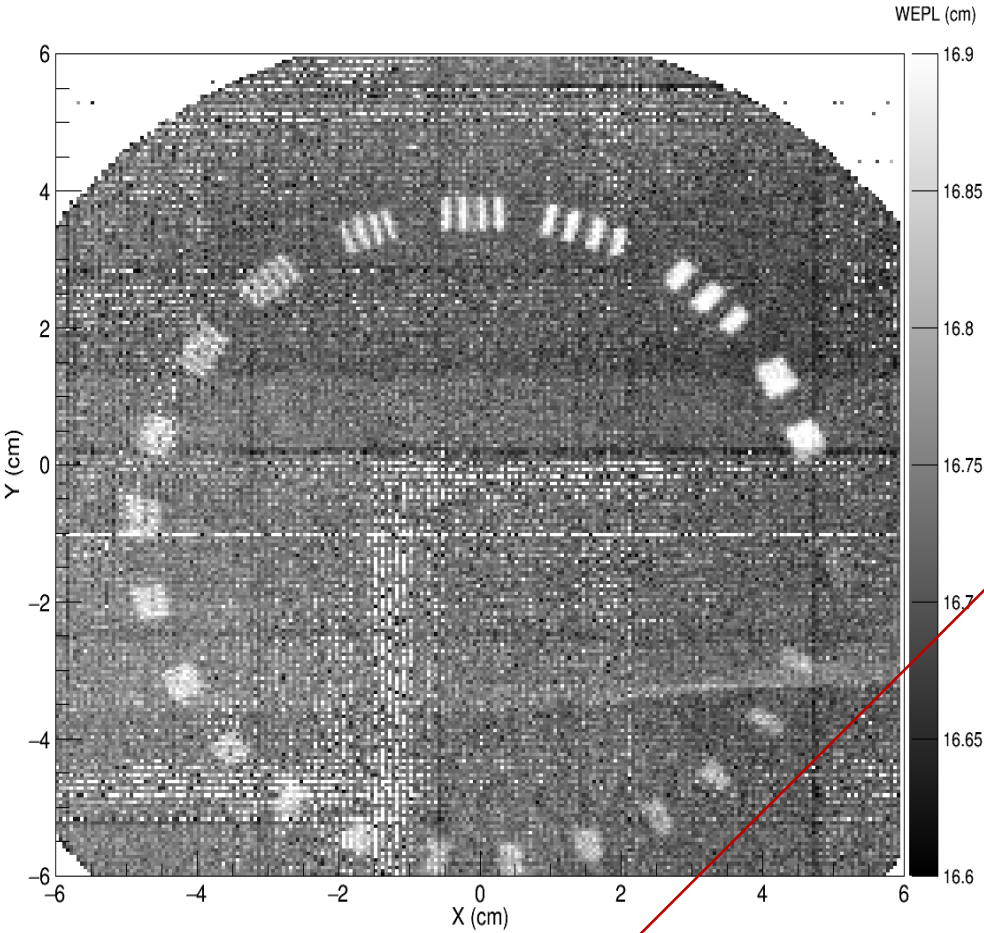
CATPHAN Line Pair Phantom



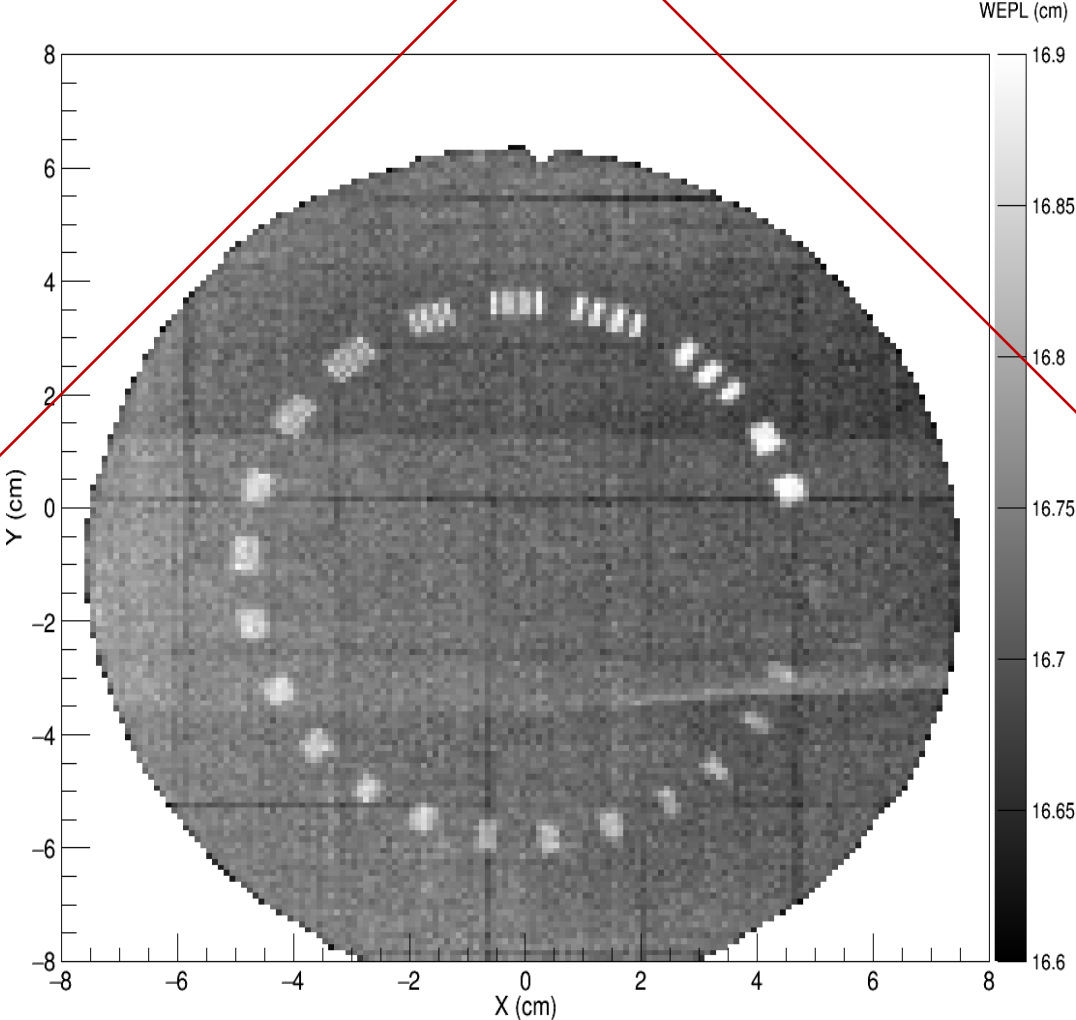
Offline reconstruction

Note: grey scale range only 3 mm!

As to range sensitivity.. We clearly see the 0.2 mm tape supporting the phantom. The density of this tape is slightly above water density.

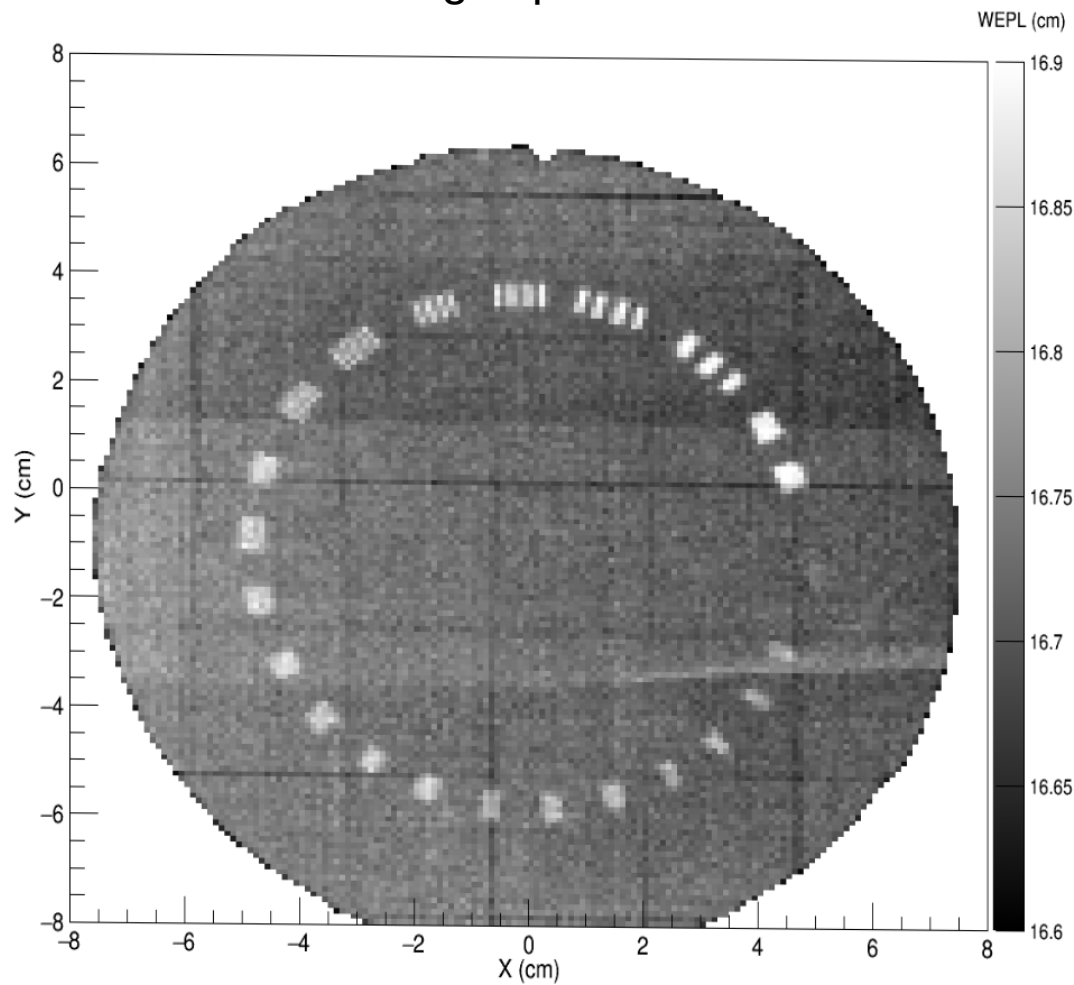


We see 7 line pairs/cm with 0.5 mm pixel size

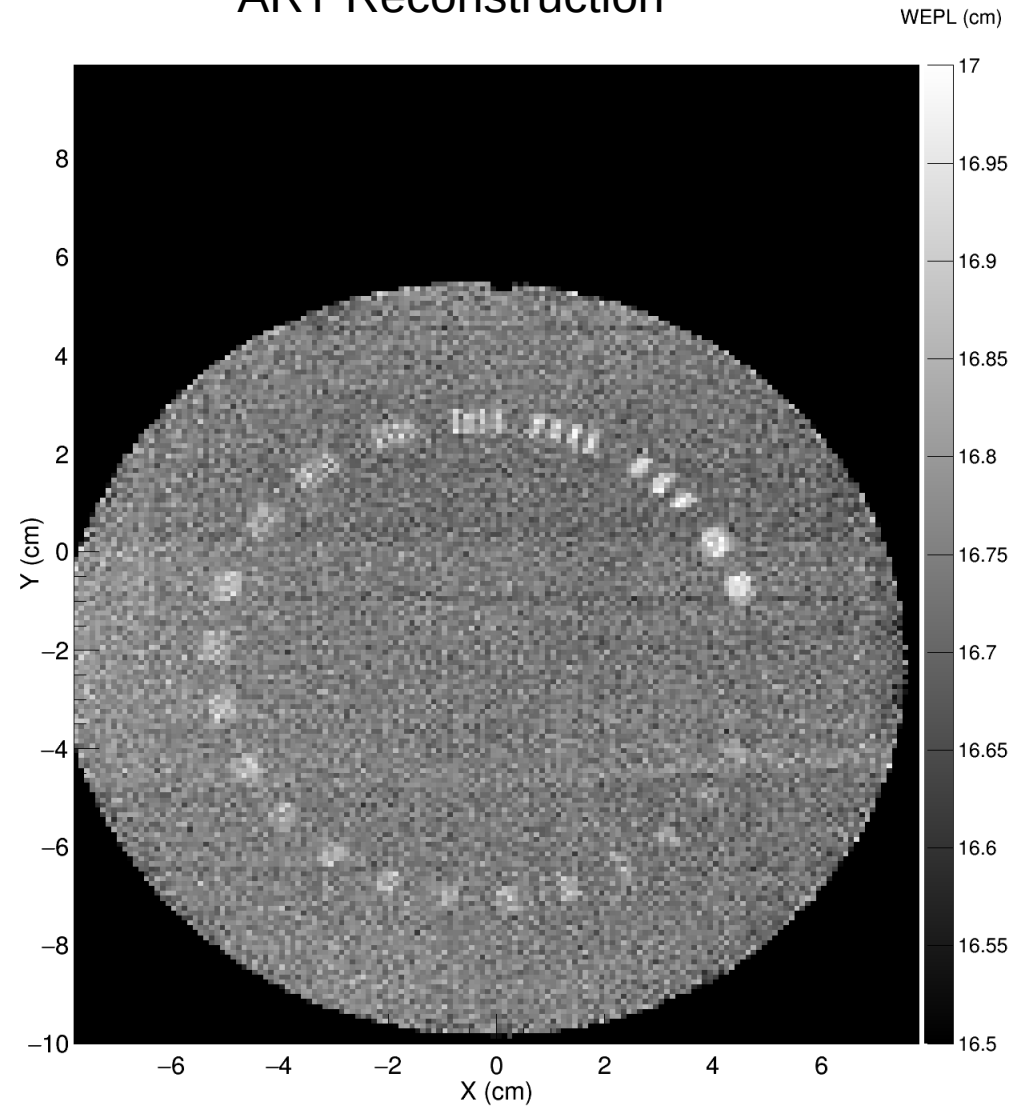


We see 6 line pairs/cm with 1 mm pixel size

2-D Binning in plane of inserts



ART Reconstruction



Loma Linda pCT reconstructed image (offline) of the same line-pair phantom

1014 Giacometti *et al.*: Software platform for simulation of proton CT

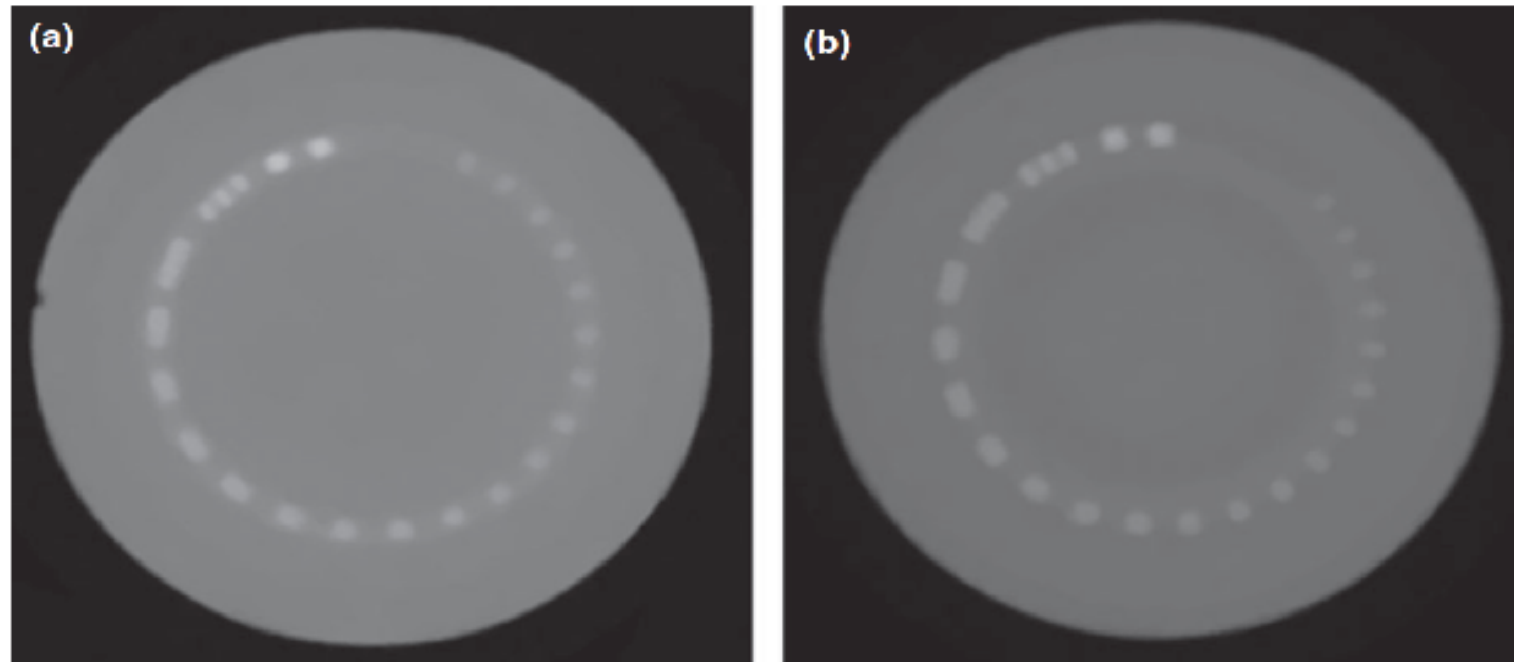
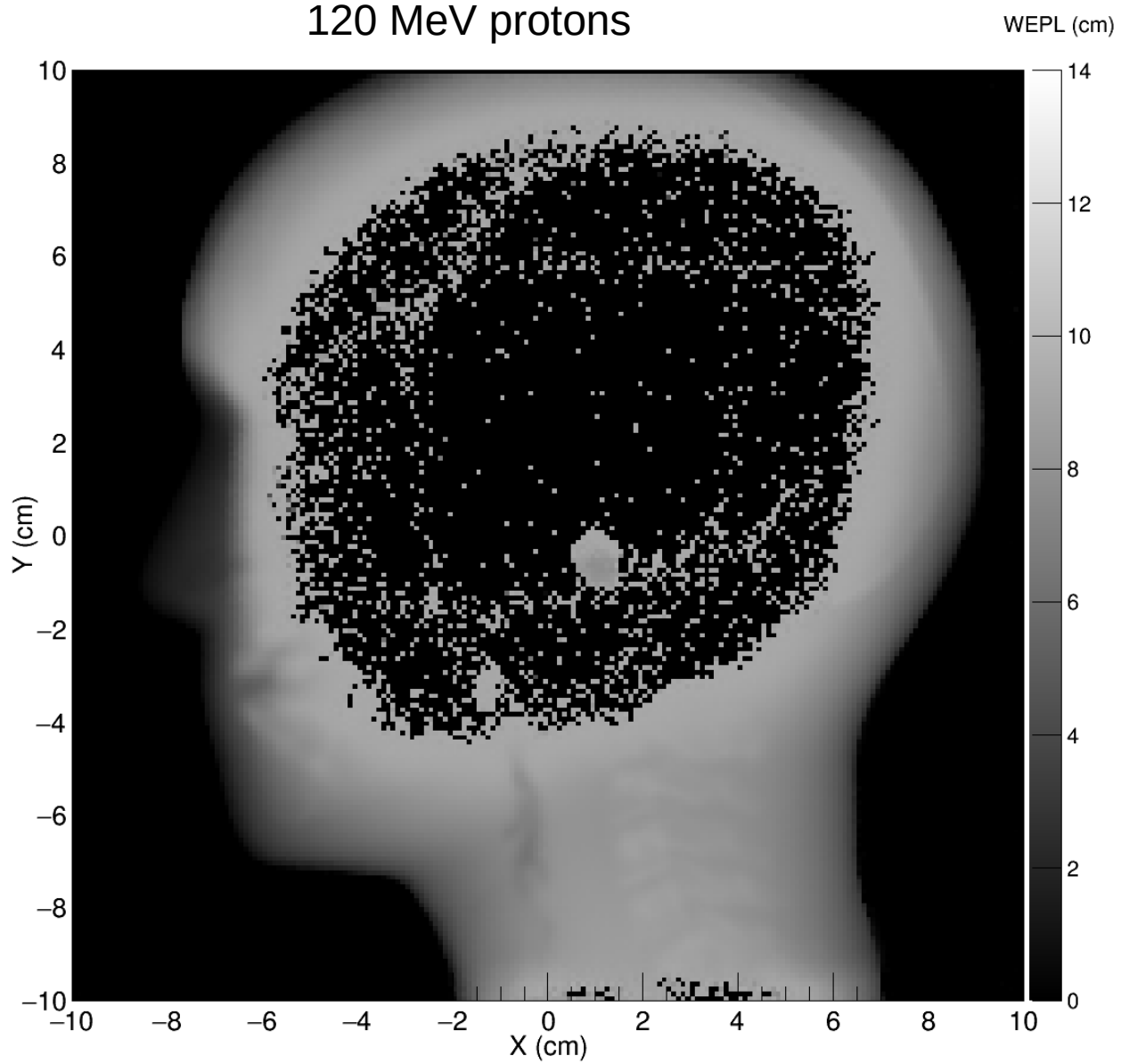


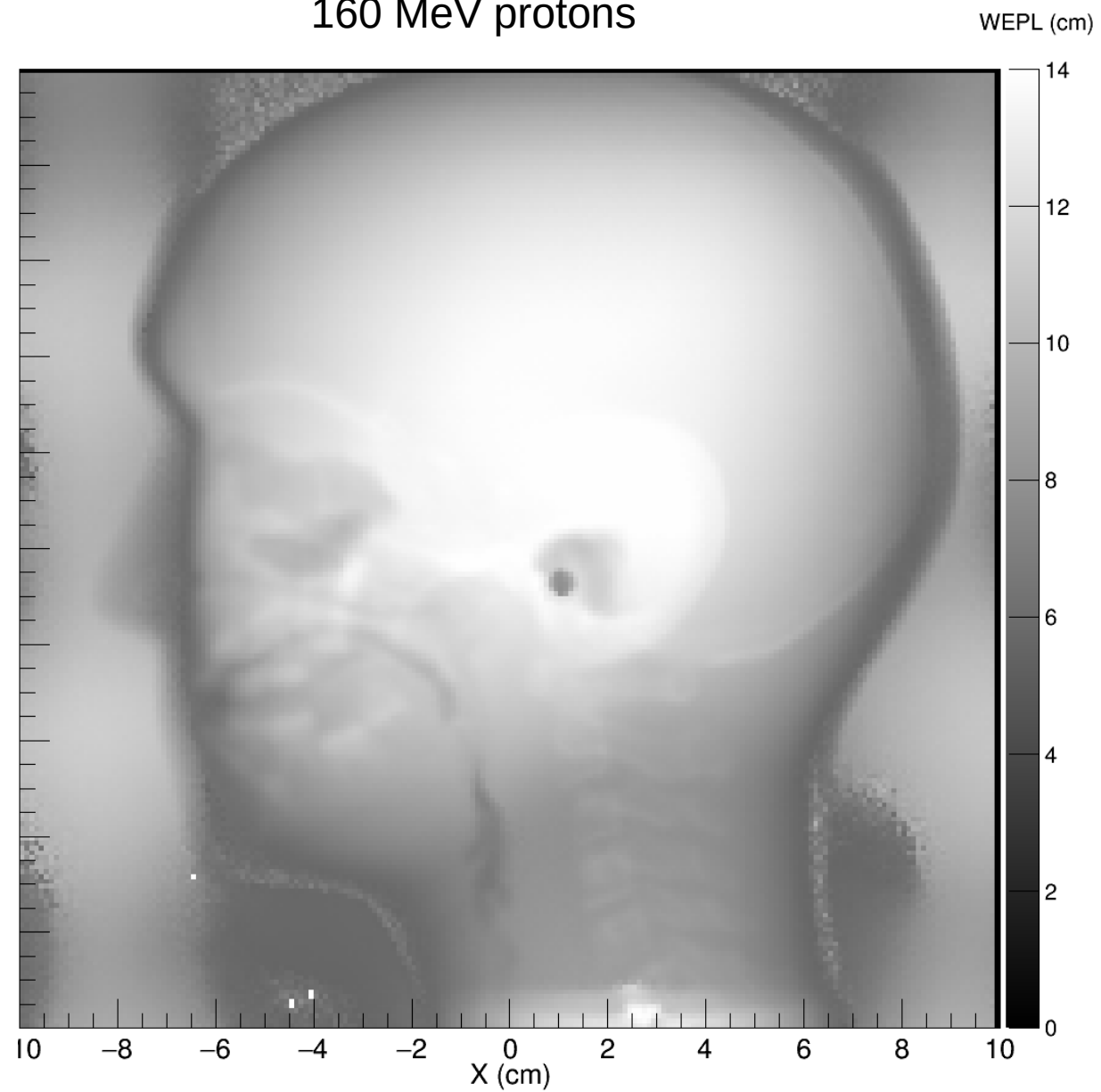
FIG. 13. Catphan[®] 60 Line pair module pCT reconstruction using experimental (a) and simulated (b) data.

Pediatric head phantom

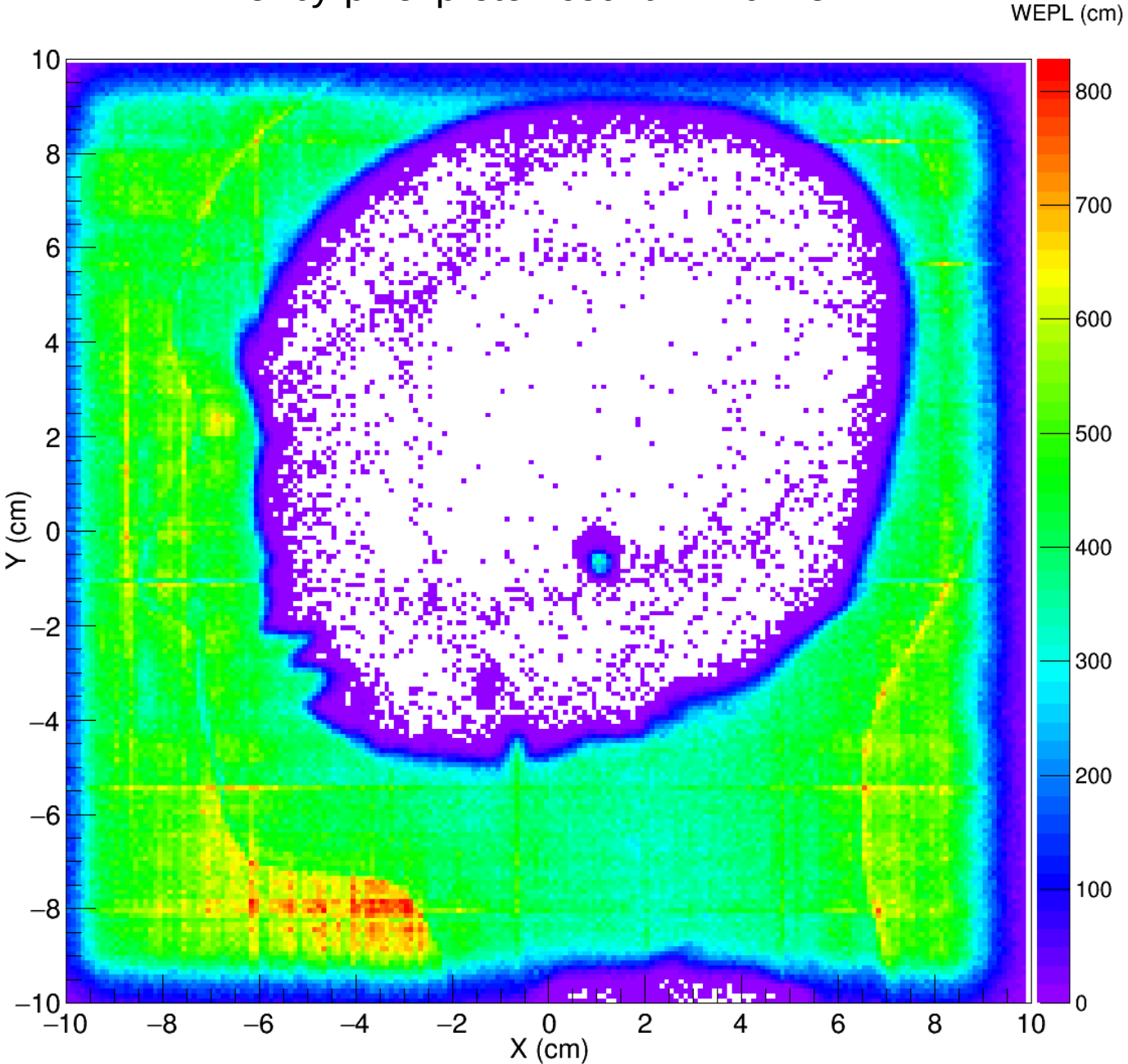
120 MeV protons



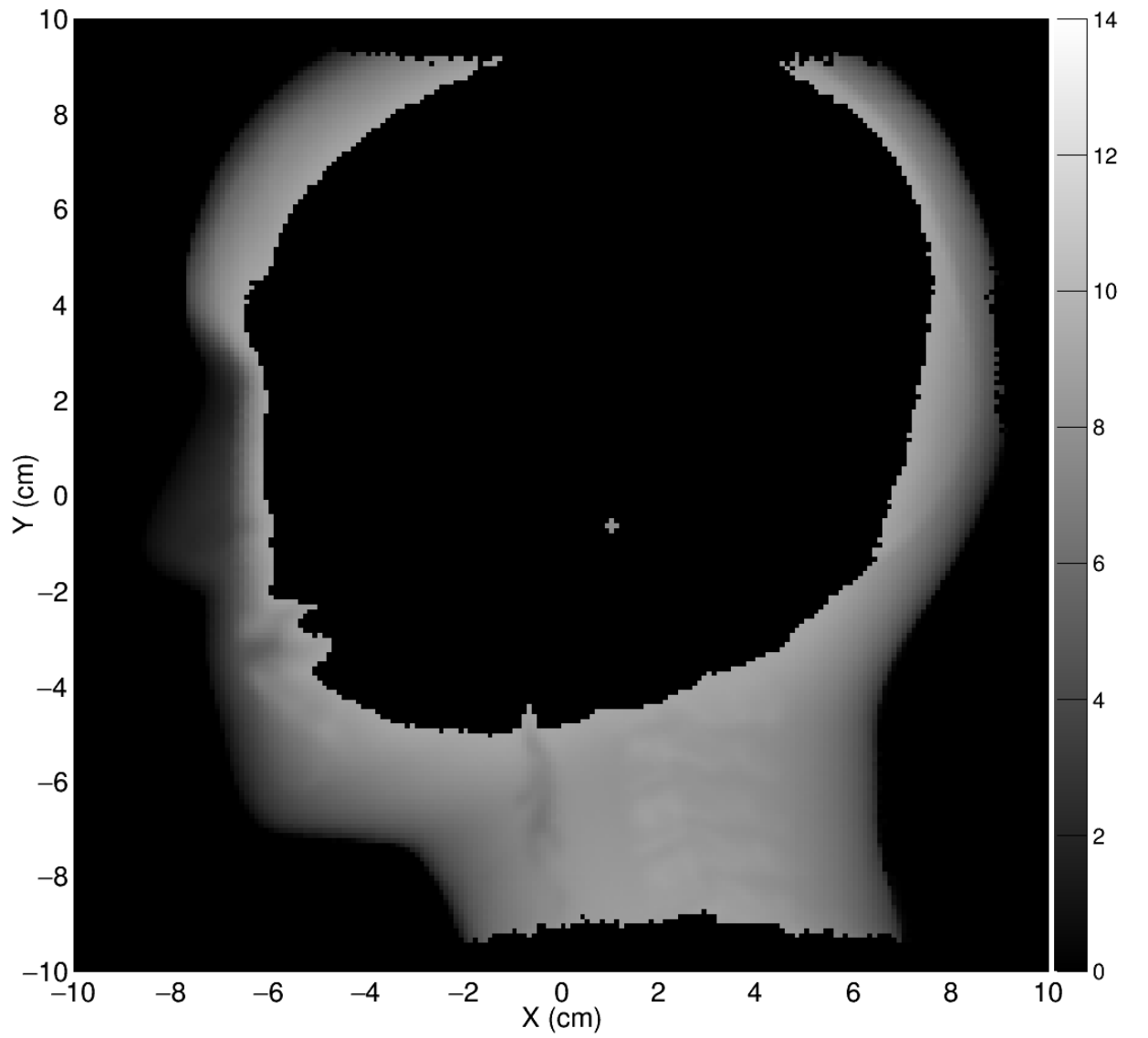
160 MeV protons



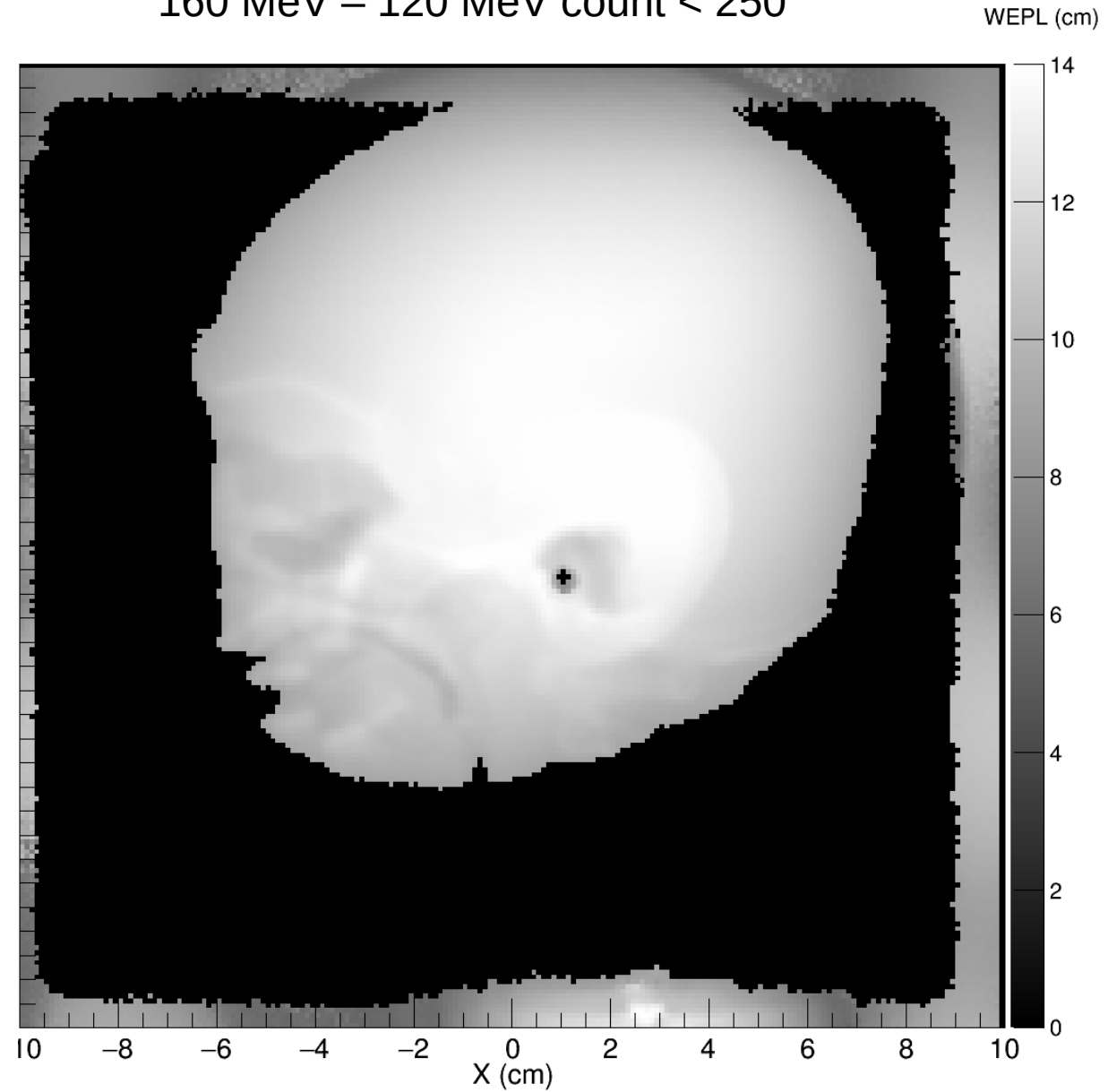
Pixel-by-pixel proton count – 120 MeV

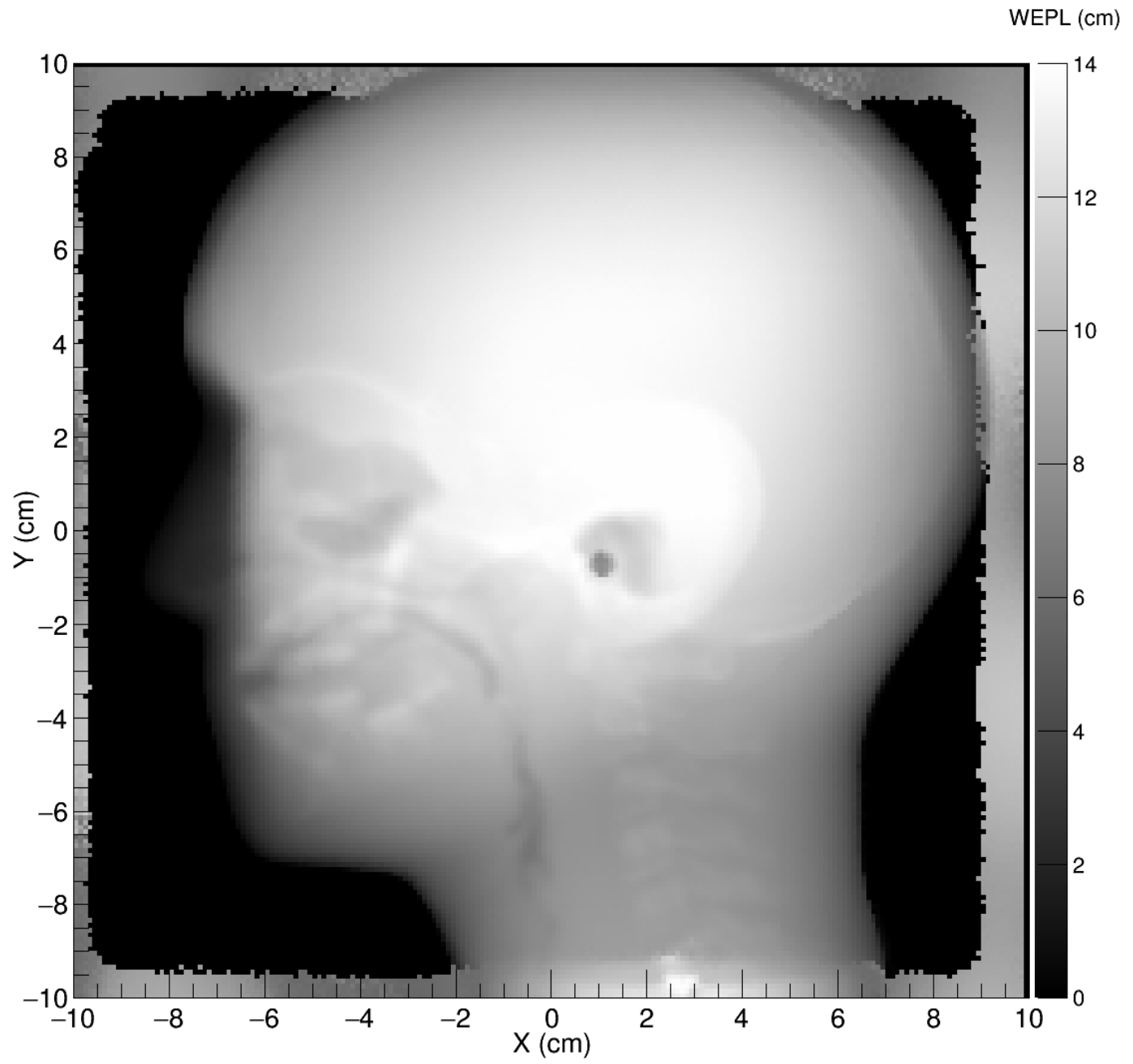


120 MeV – count > 250



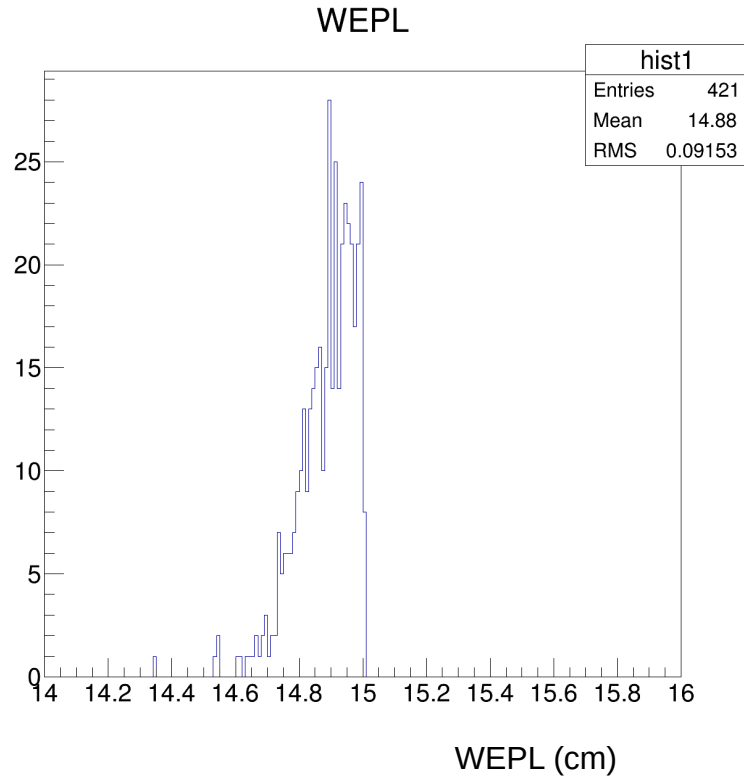
160 MeV – 120 MeV count < 250



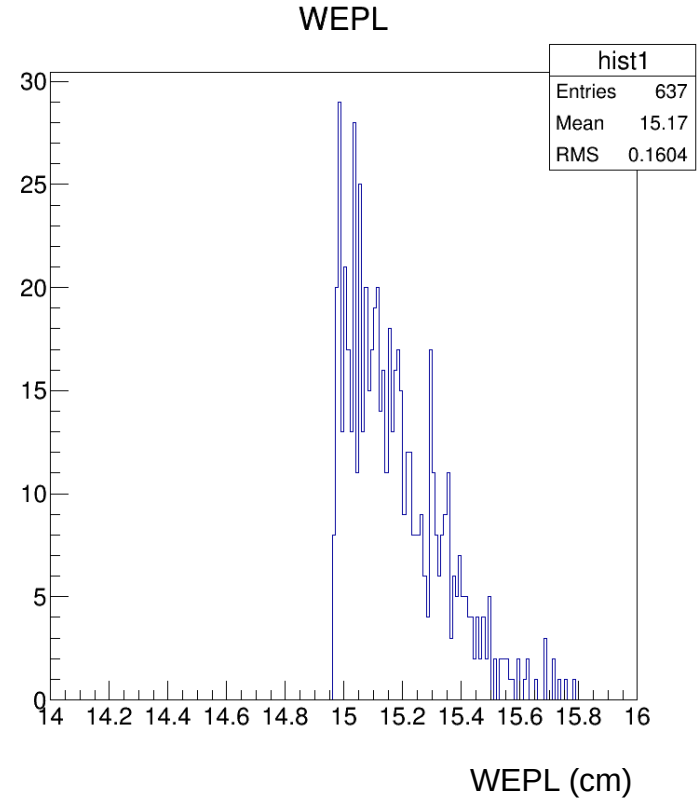


Energy Boundary Artifacts

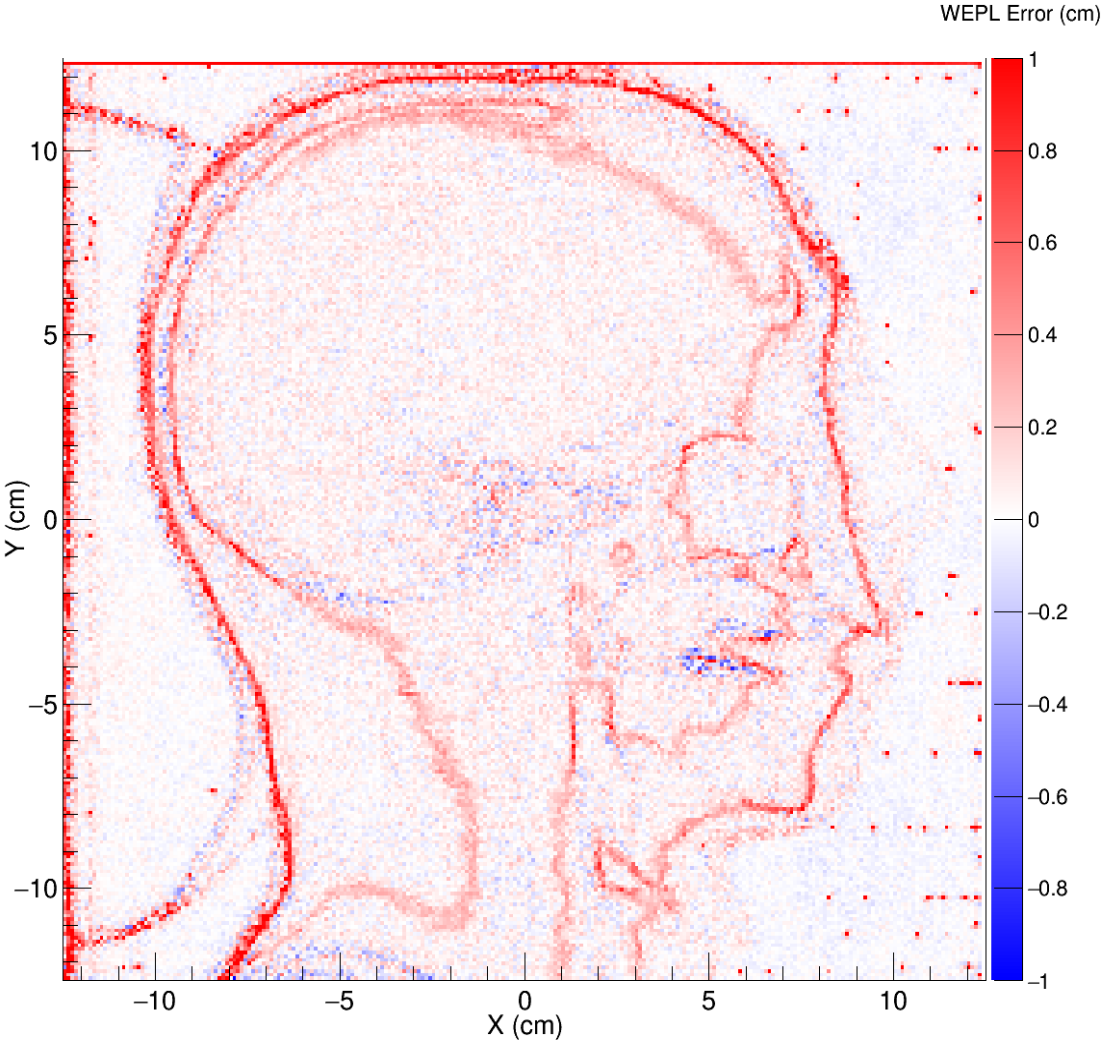
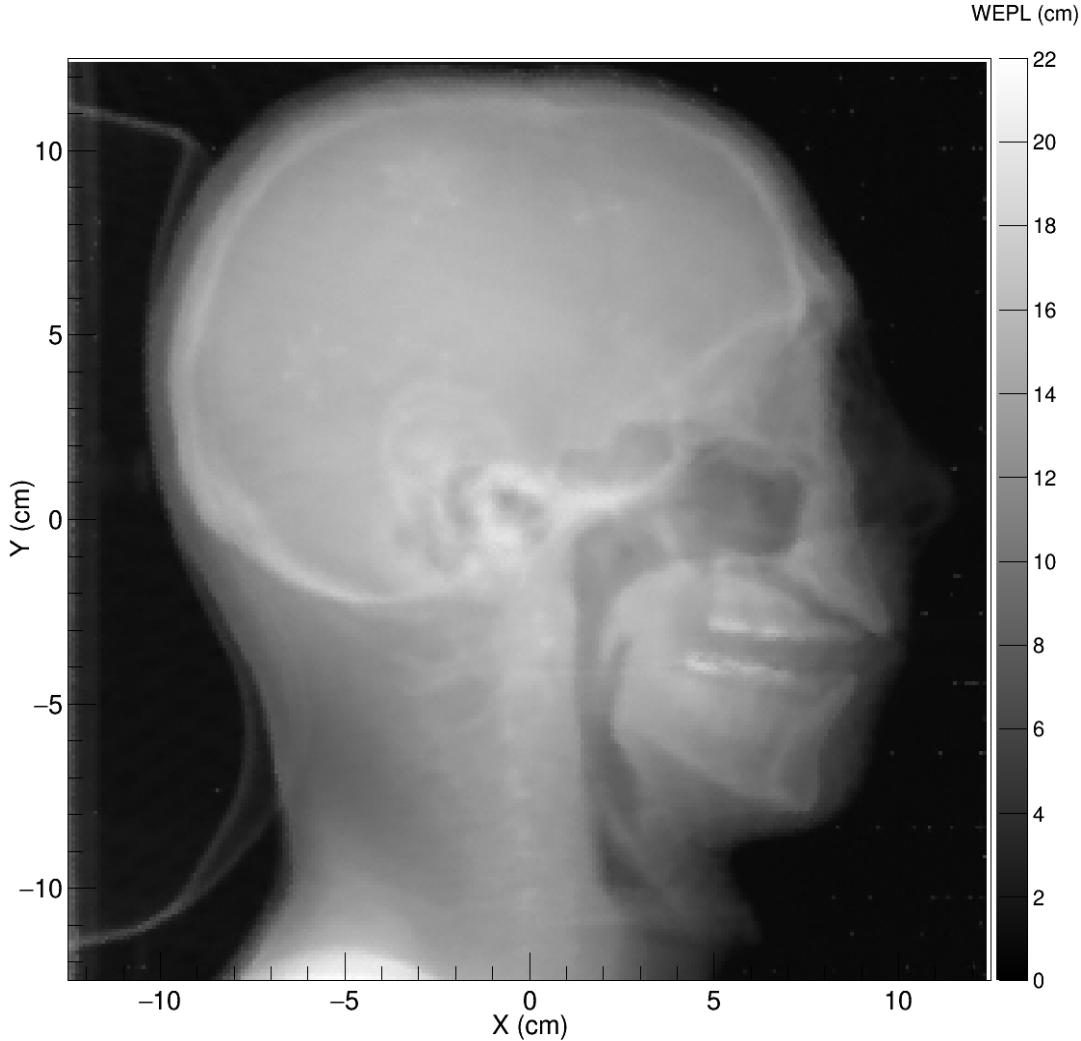
16 cm range beam through 15 cm water:



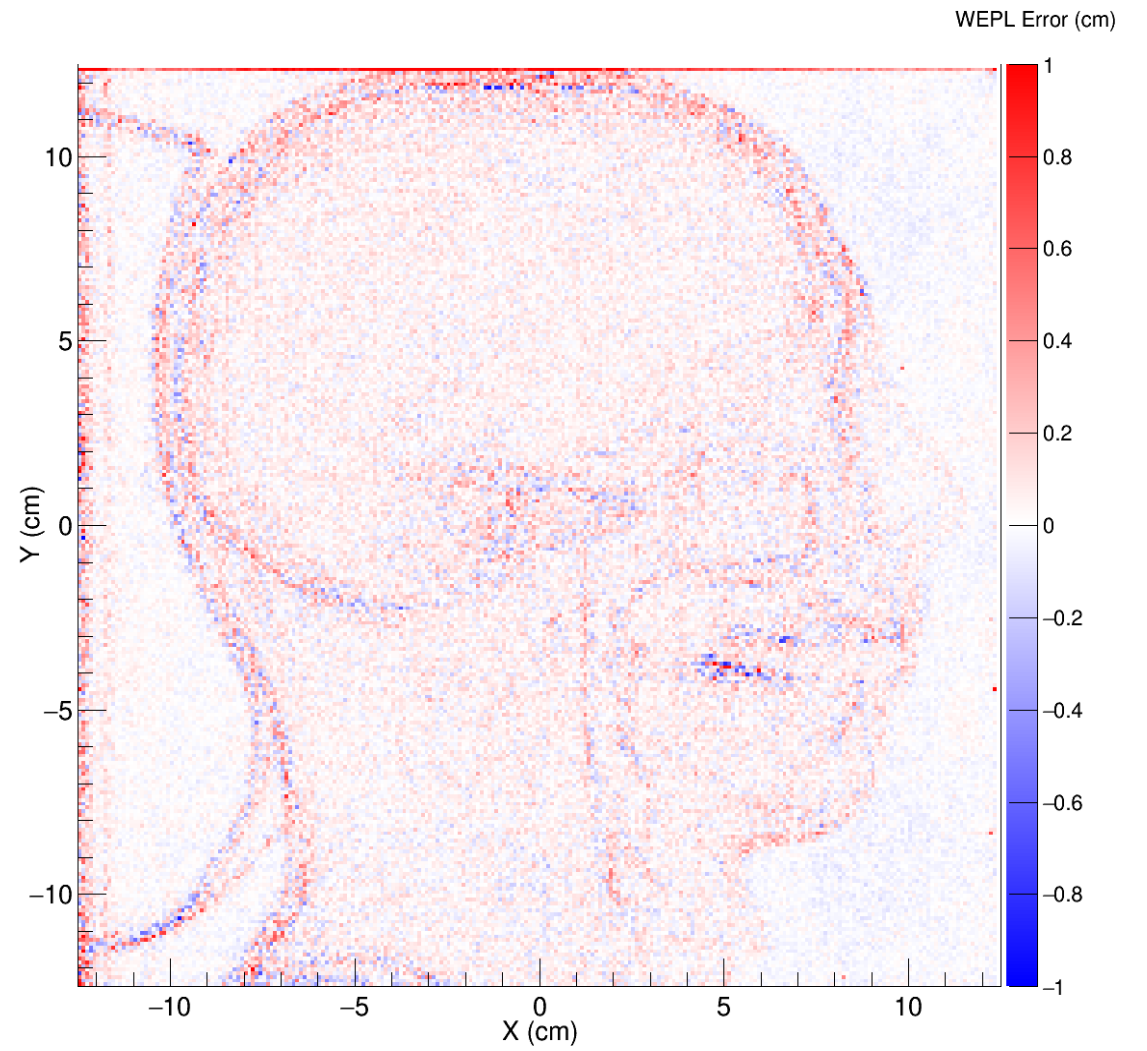
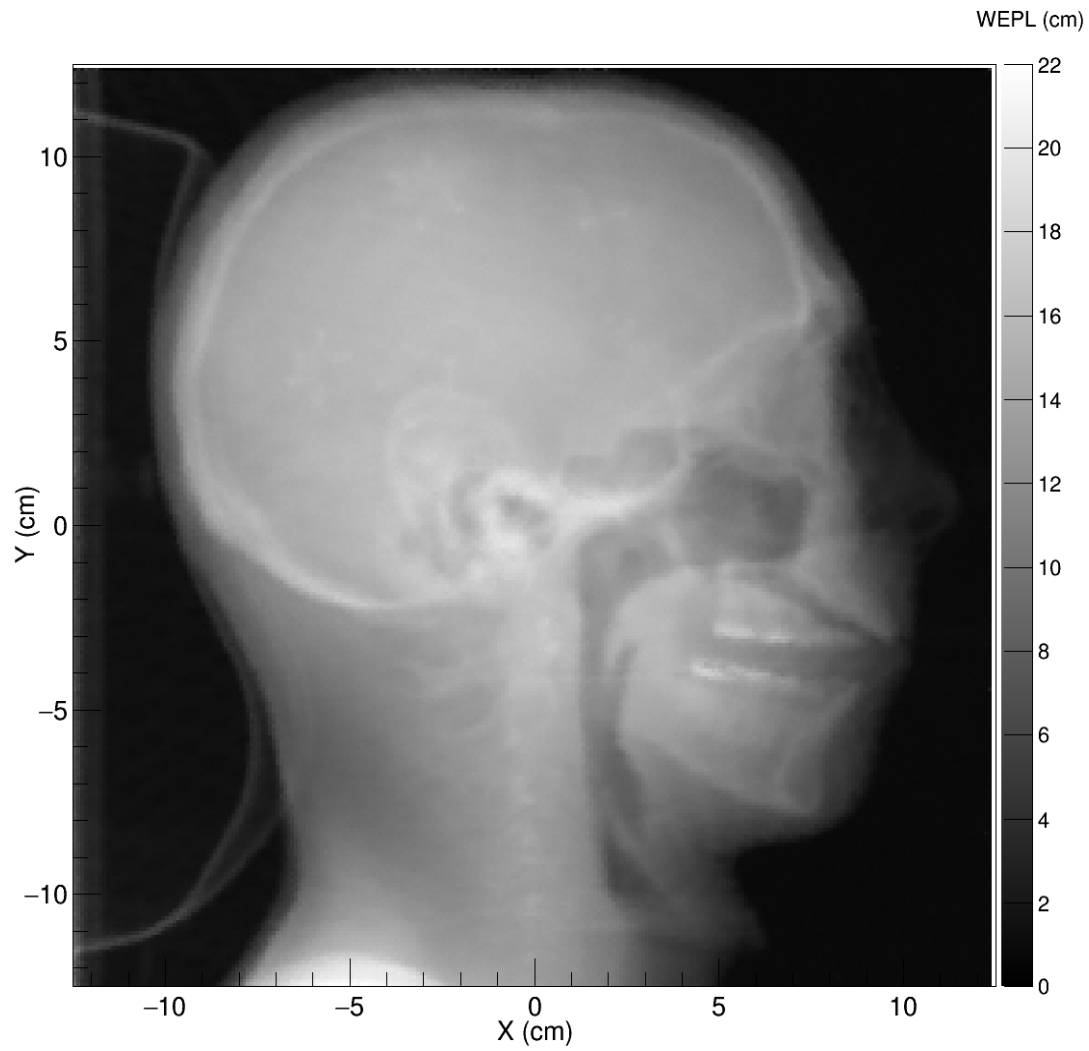
26 cm range beam through 15 cm water:



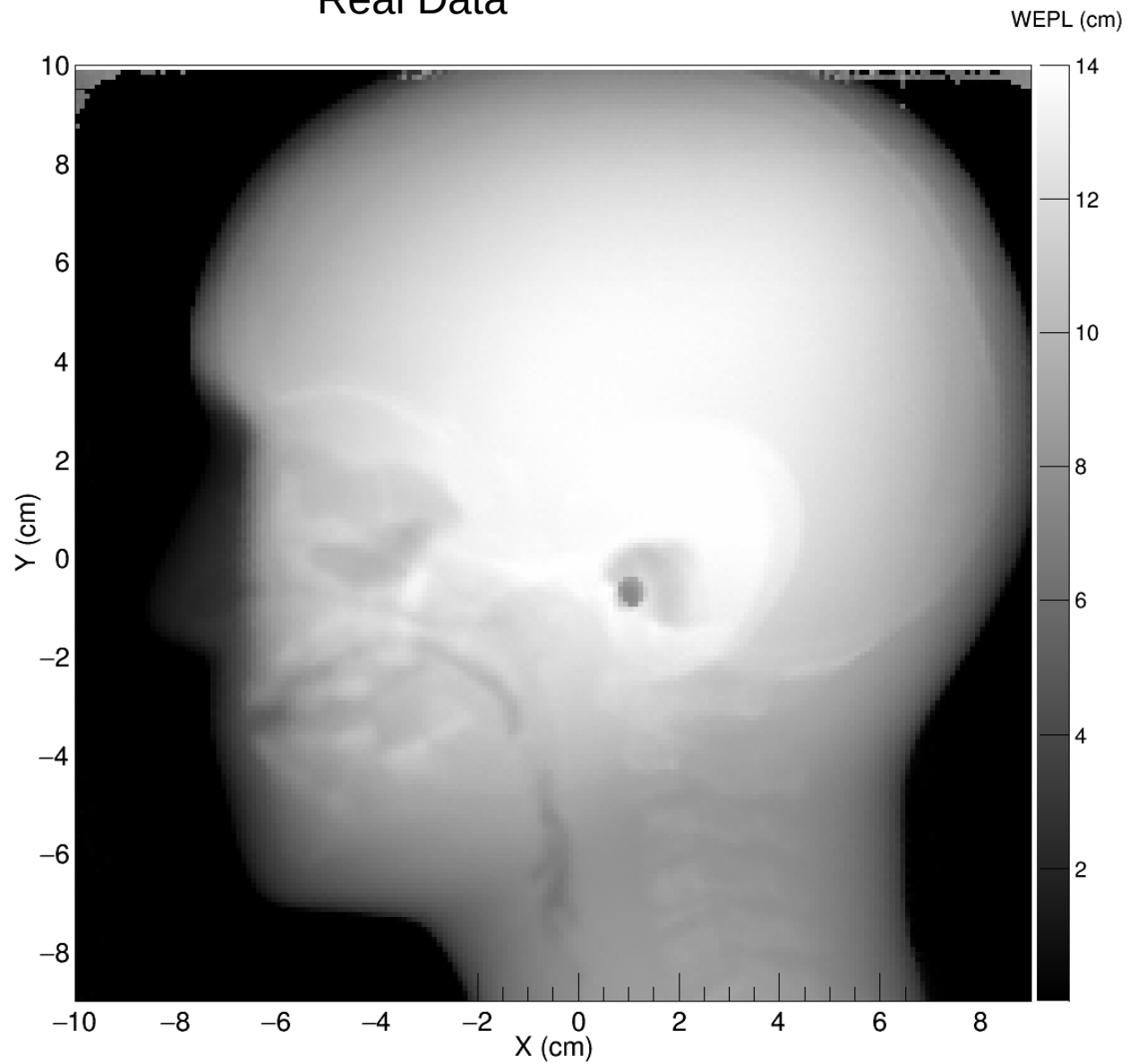
No Energy Boundary Cuts



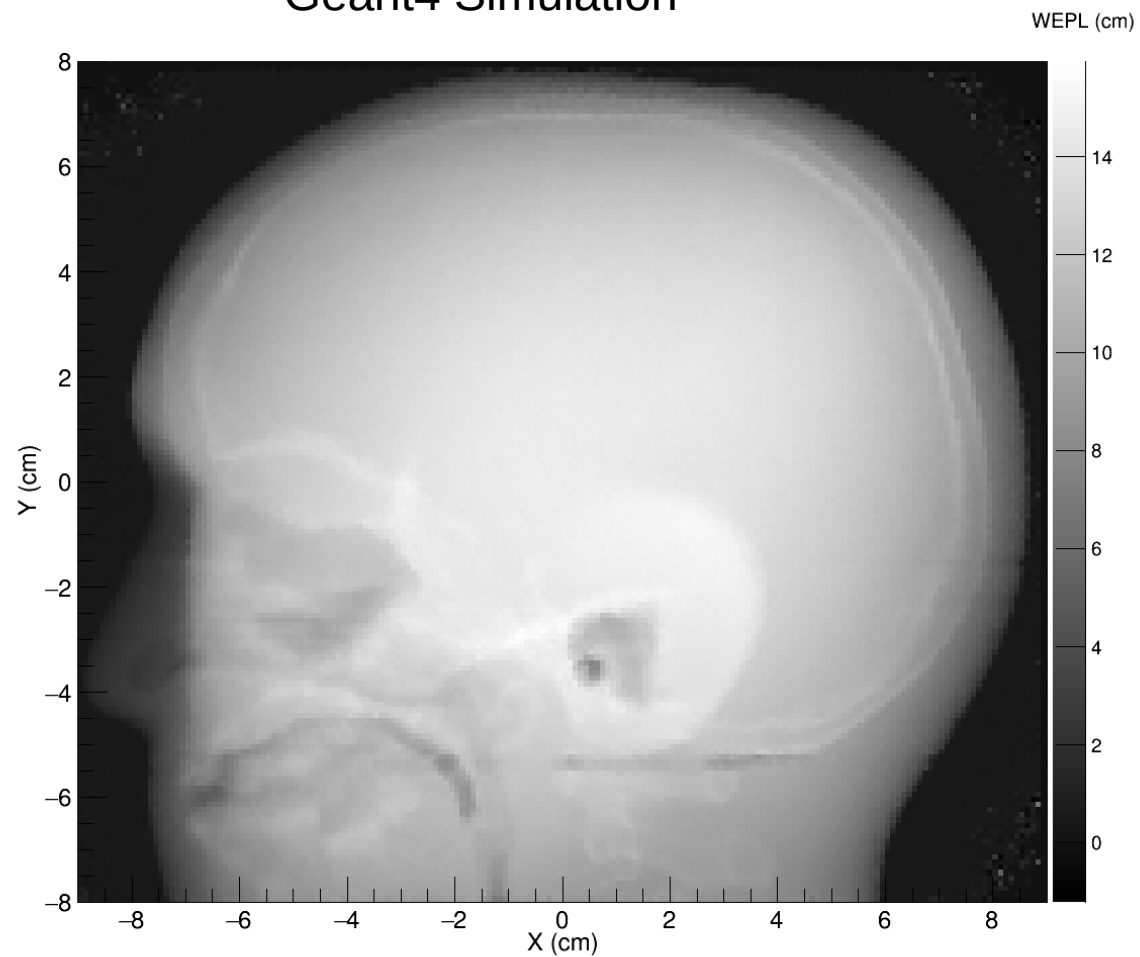
With Energy Boundary Cuts



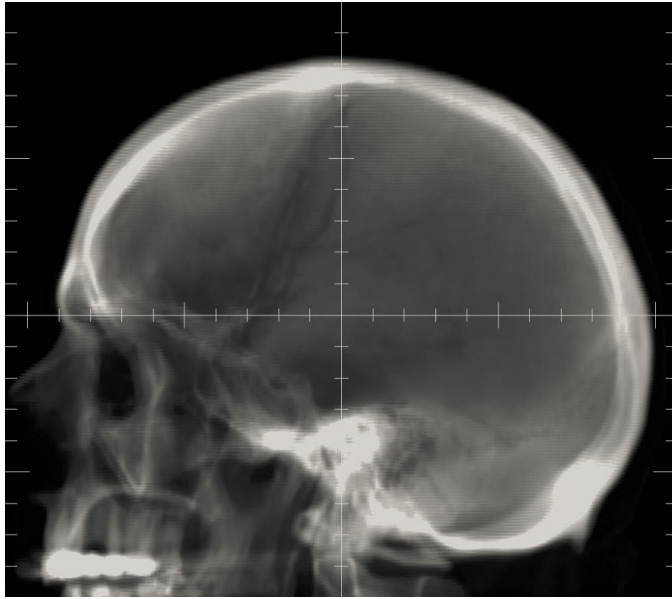
Real Data



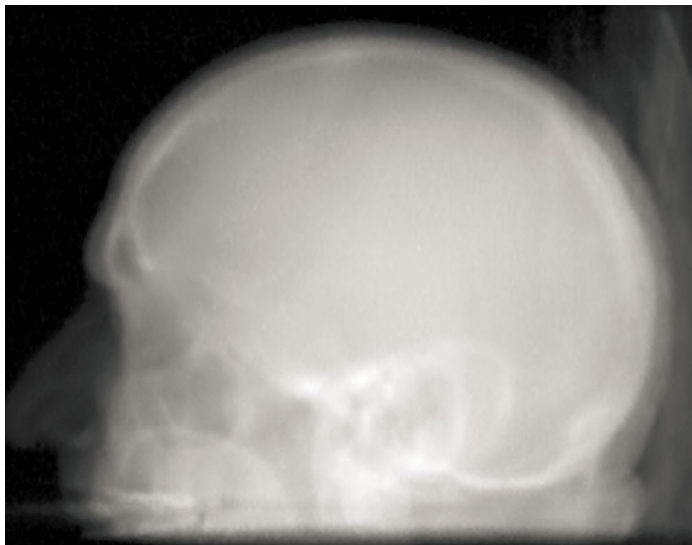
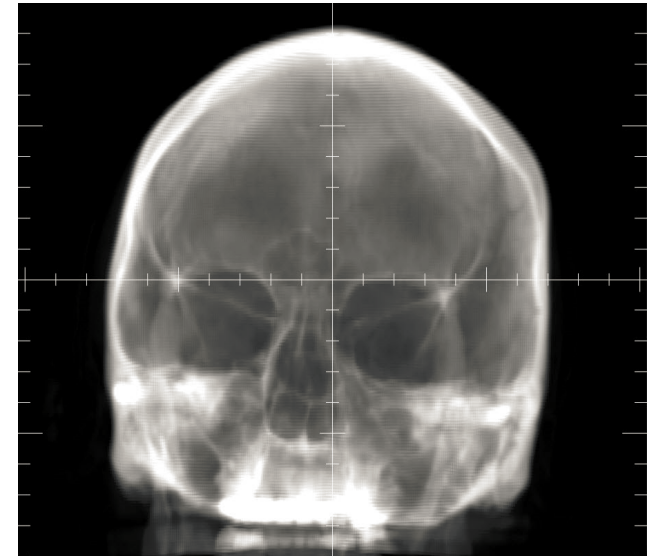
Geant4 Simulation



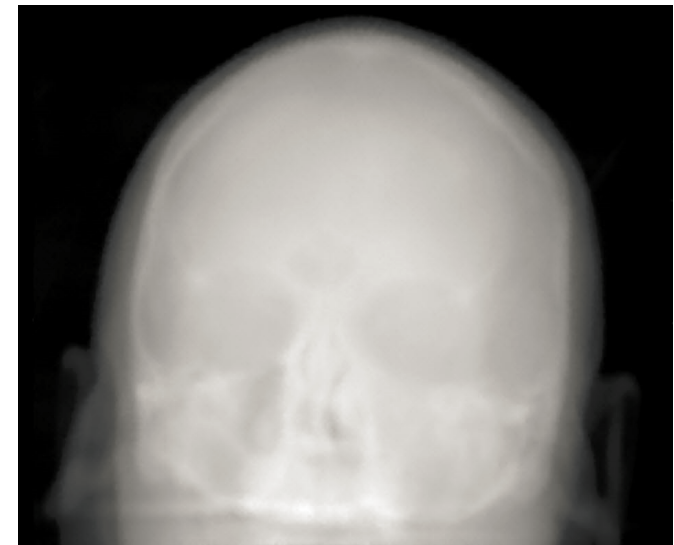
Are these images good enough to align the patient??

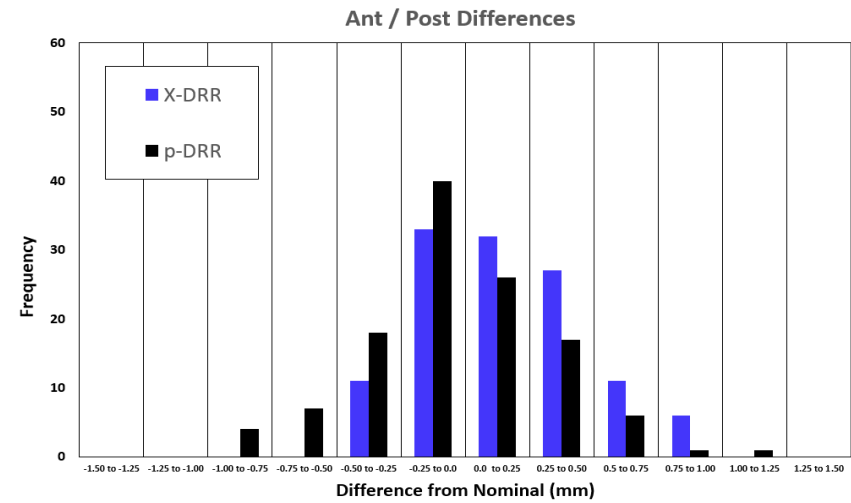
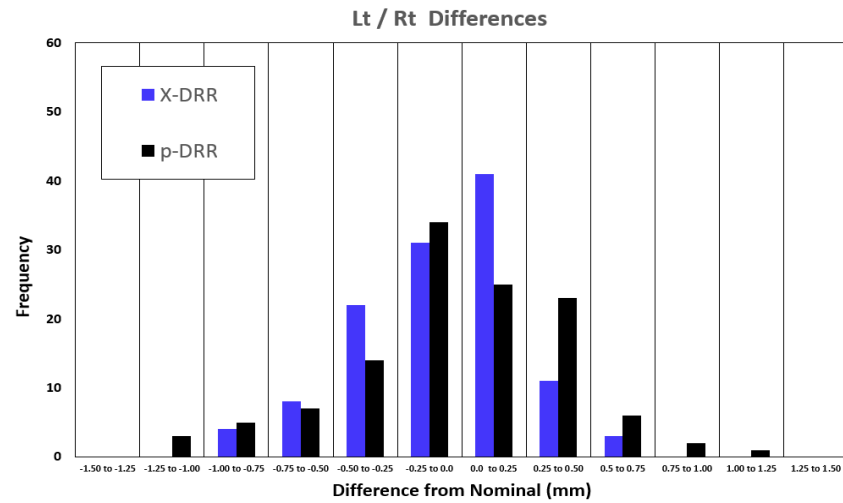


X-DRR



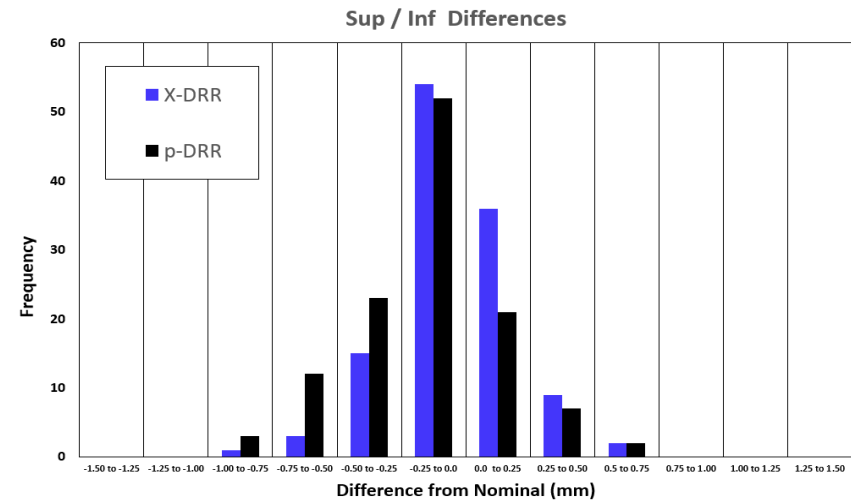
p-DRR

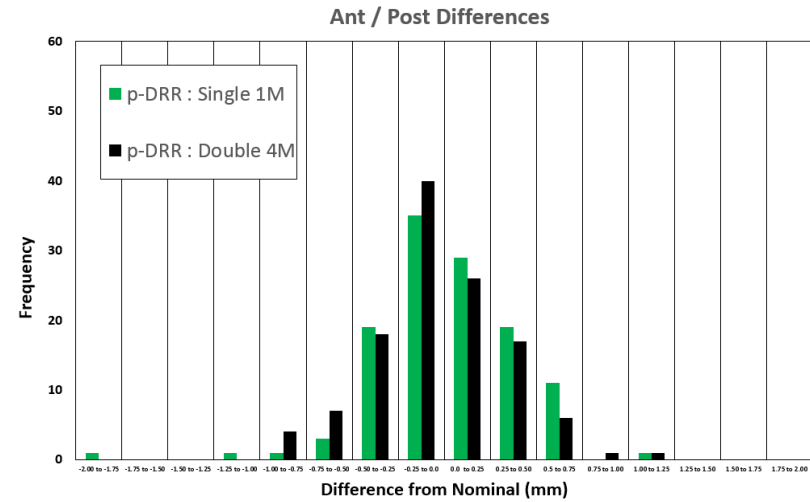
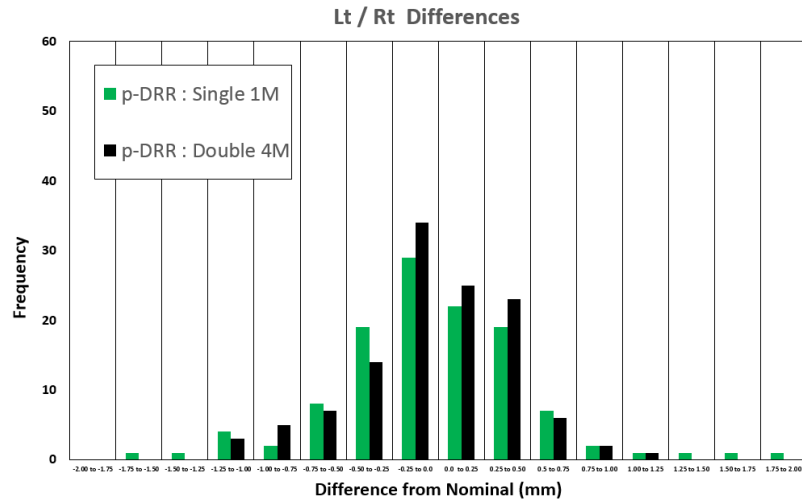




Average Difference (mm)			
	Lt / Rt	Ant/Post	Sup/Inf
X-DRR	-0.1	0.2	0.0
p-DRR	0.0	0.0	-0.1

Standard Deviation (mm)			
	Lt / Rt	Ant/Post	Sup/Inf
X-DRR	0.3	0.3	0.2
p-DRR	0.4	0.4	0.3





Average Difference (mm)			
	Lt / Rt	Ant/Post	Sup/Inf
p-DRR1	-0.1	0.0	-0.1
p-DRR4	0.0	0.0	-0.1

Standard Deviation (mm)			
	Lt / Rt	Ant/Post	Sup/Inf
p-DRR1	0.6	0.4	0.4
p-DRR4	0.4	0.4	0.3

