

# **RDH-WP11-FRED**

**A. Schiavi - Sapienza Univ. di Roma**

RDH-IRPT meeting - 01/02/**2017**  
Dip. SBAl - Sapienza - Roma

# CNAO

S. Molinelli

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# FRED

- **Angelo Schiavi** - Sapienza-Roma / INFN
- Vincenzo Patera - Sapienza-Roma / INFN
- Martina Senzacqua - Post-grad.
- Stefano Pioli - INFN-LNF
- Giuseppe Battistoni - INFN-MI

# Milestone per il 2016

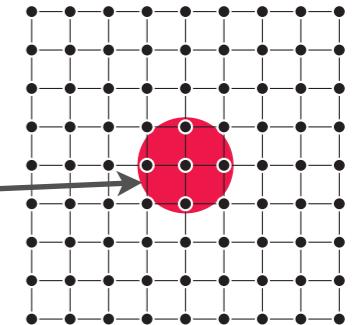
testare e validare su 5 trattamenti a protoni (testa-collo e pelvi) il ricalcolo della dose con Fred e eseguire confronto con TPS-CNAO e full-MC

- modello nucleare protoni e deutoni
- fine-tuning profili longitudinali in acqua
- fine-tuning profili laterali
- ricalcolo in acqua dei cubi per il QA protocol
- ricalcolo QA verification plan (acqua)
- ricalcolo piano trattamento protoni

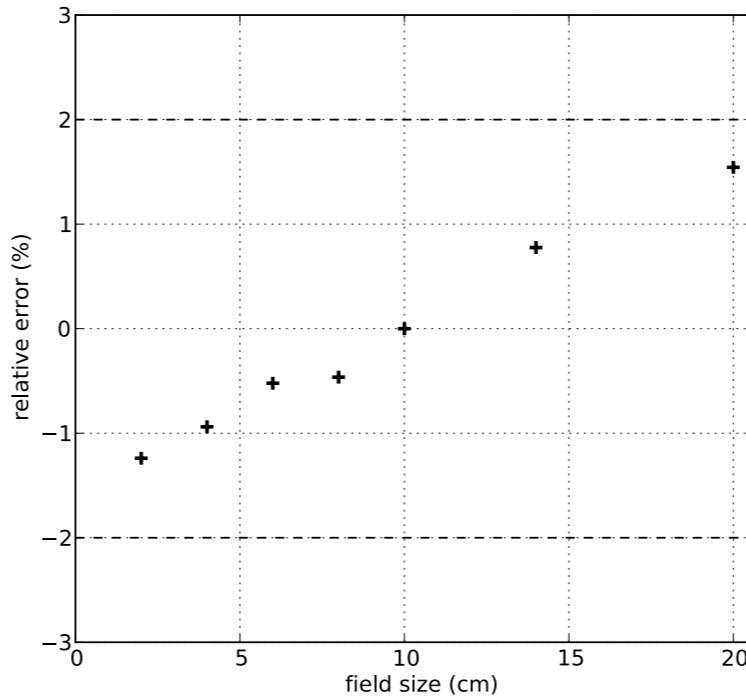
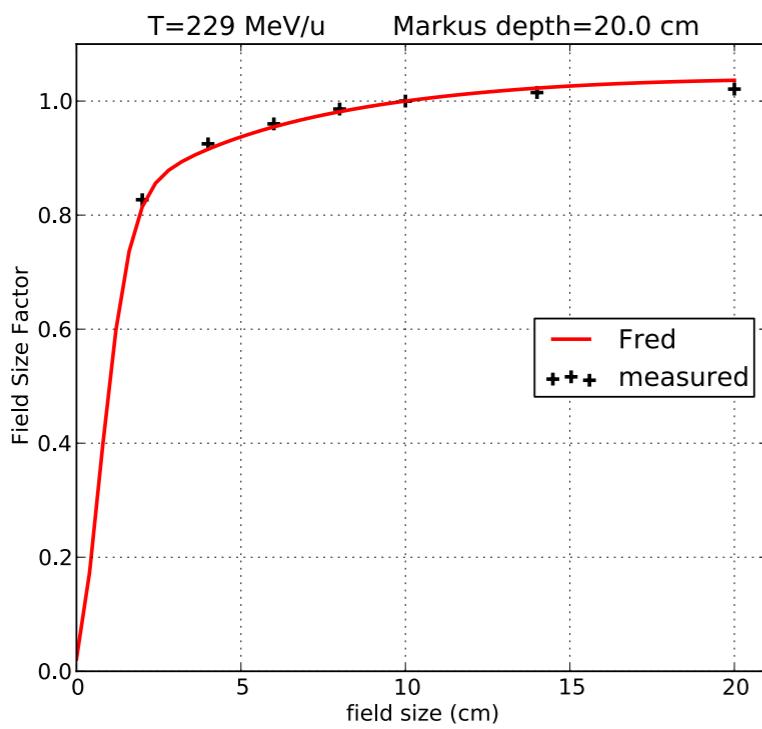
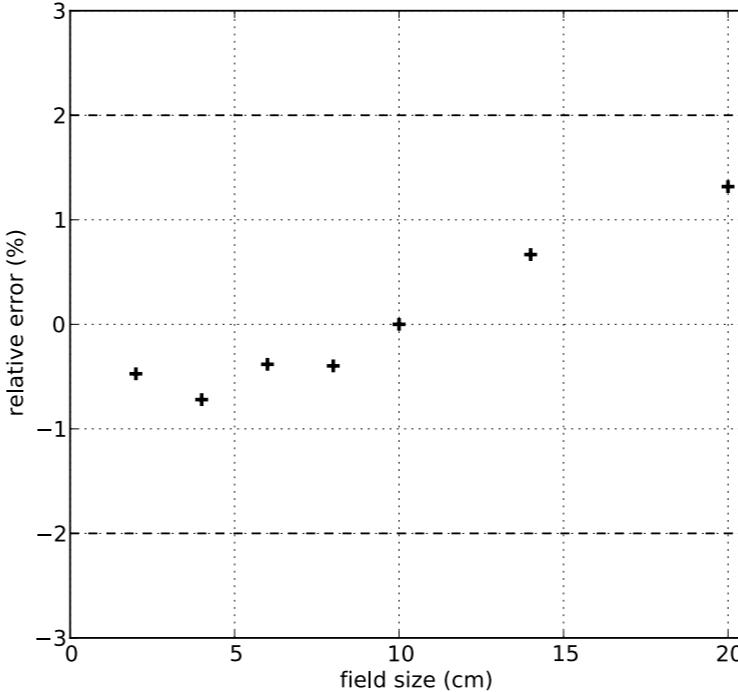
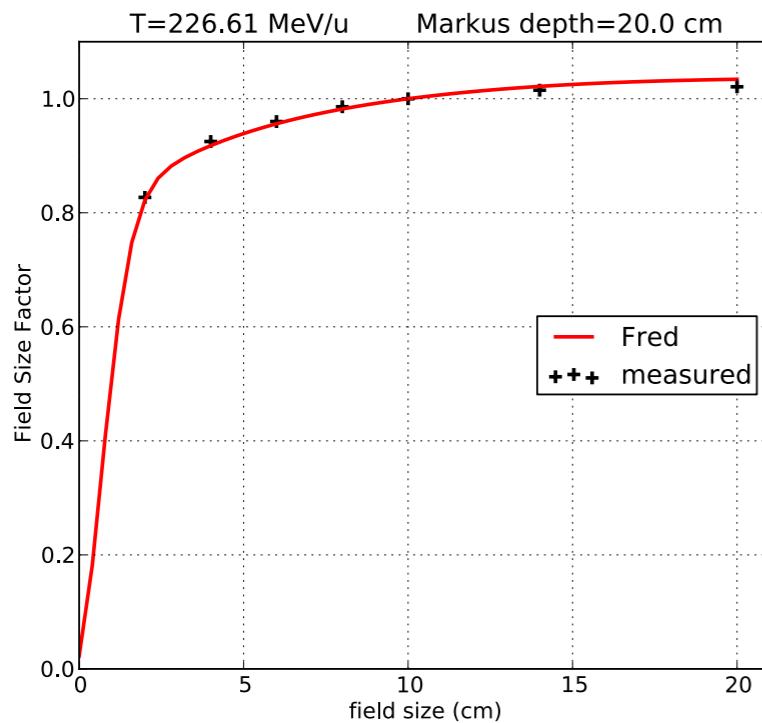
situazione un anno fa

# Field Size Factor @ CNAO

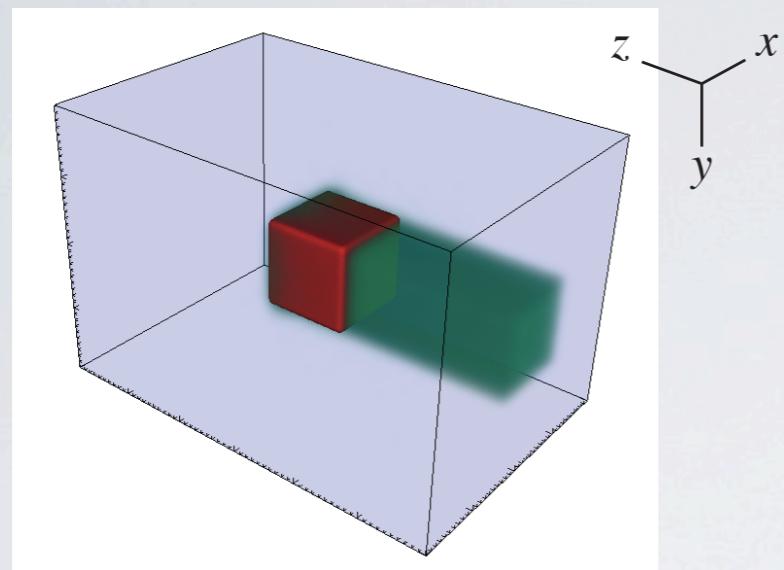
detector



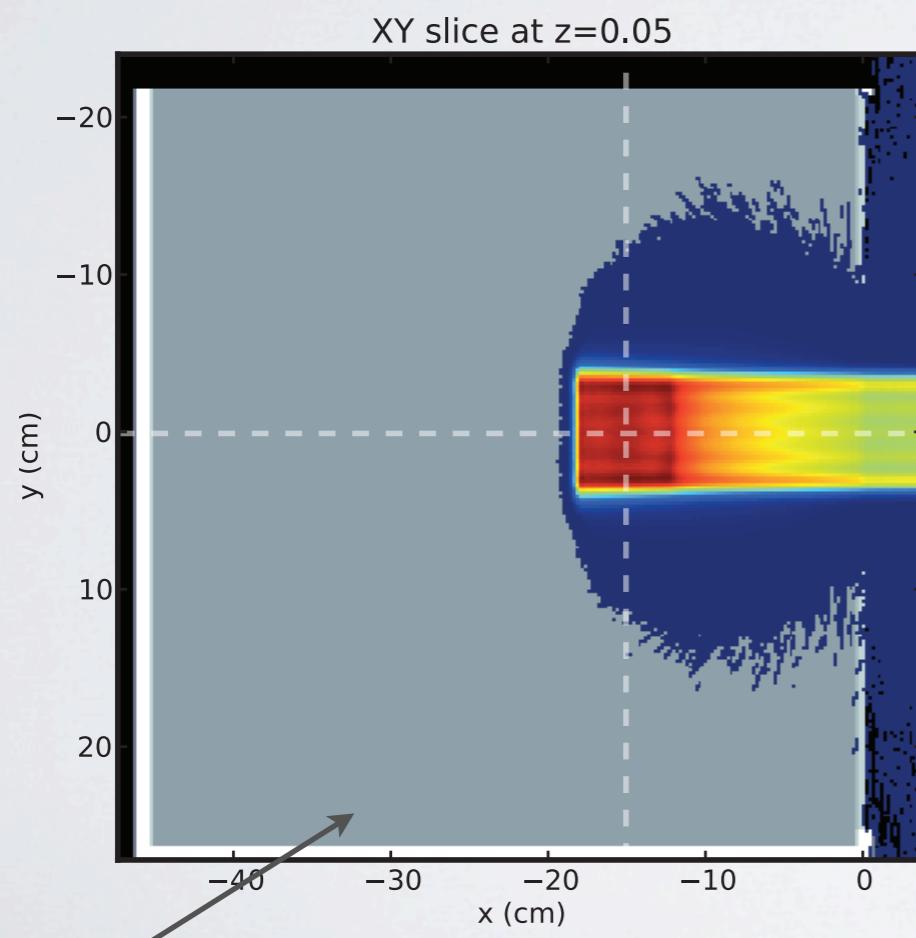
scan spots



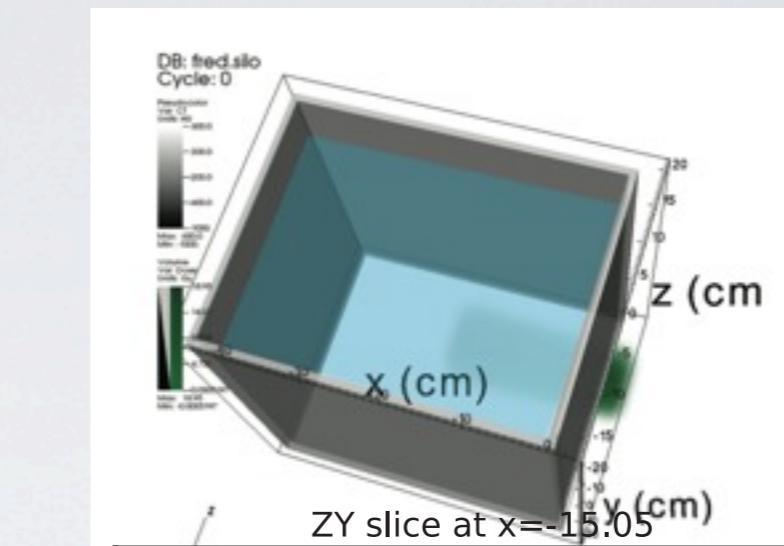
# SOBP QA - I



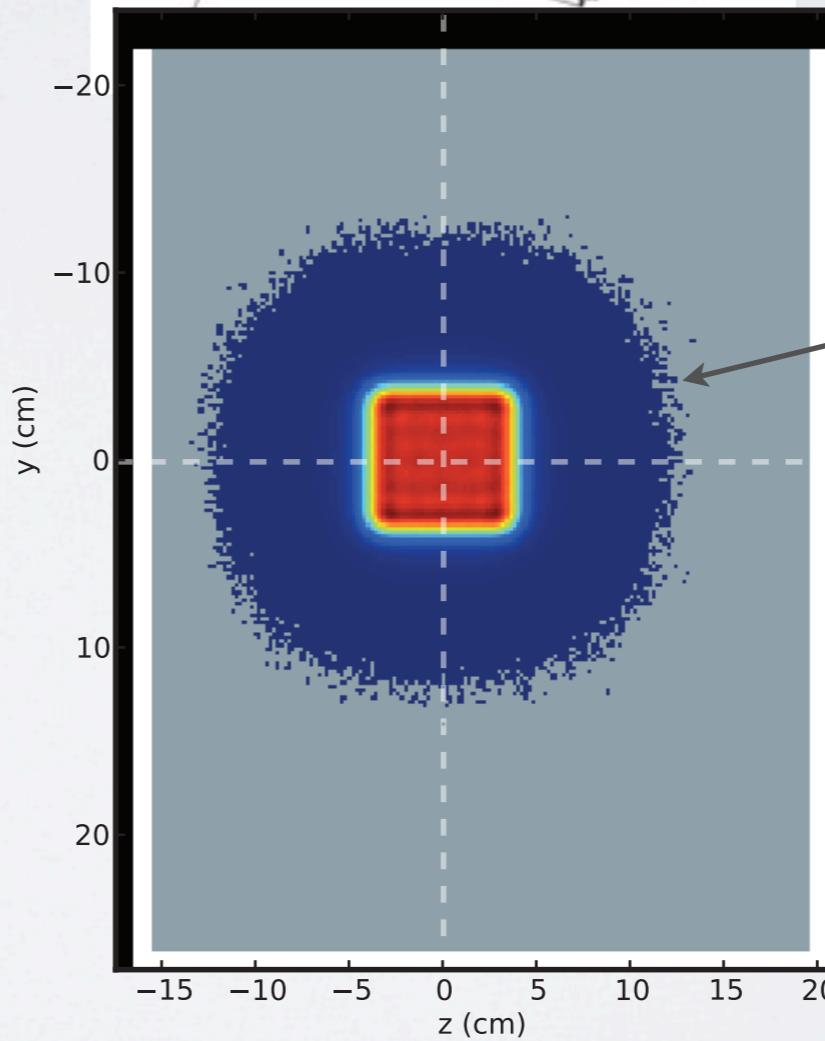
a)



b)



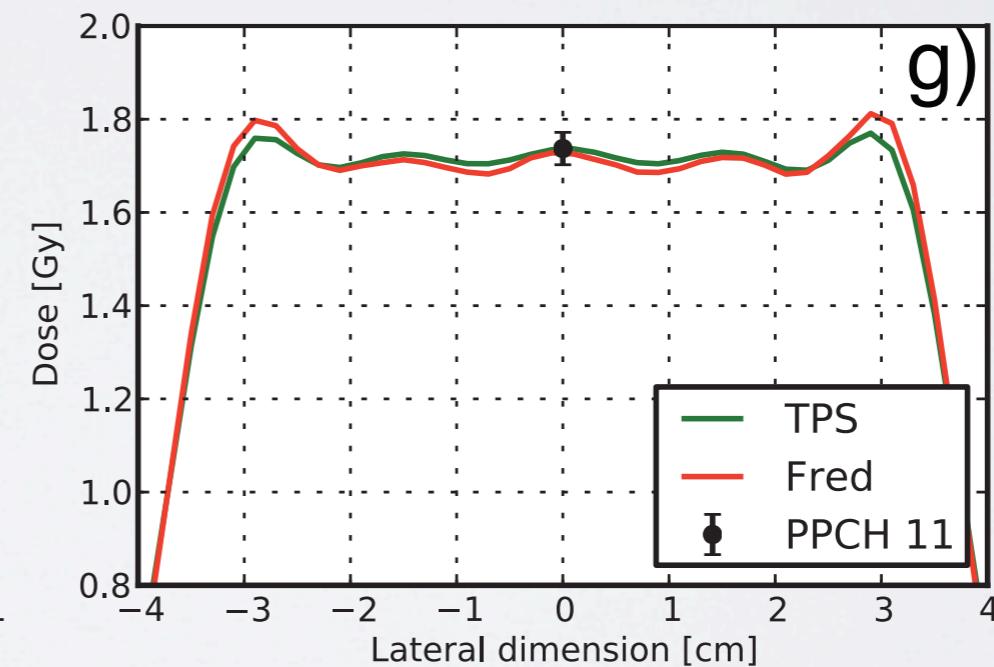
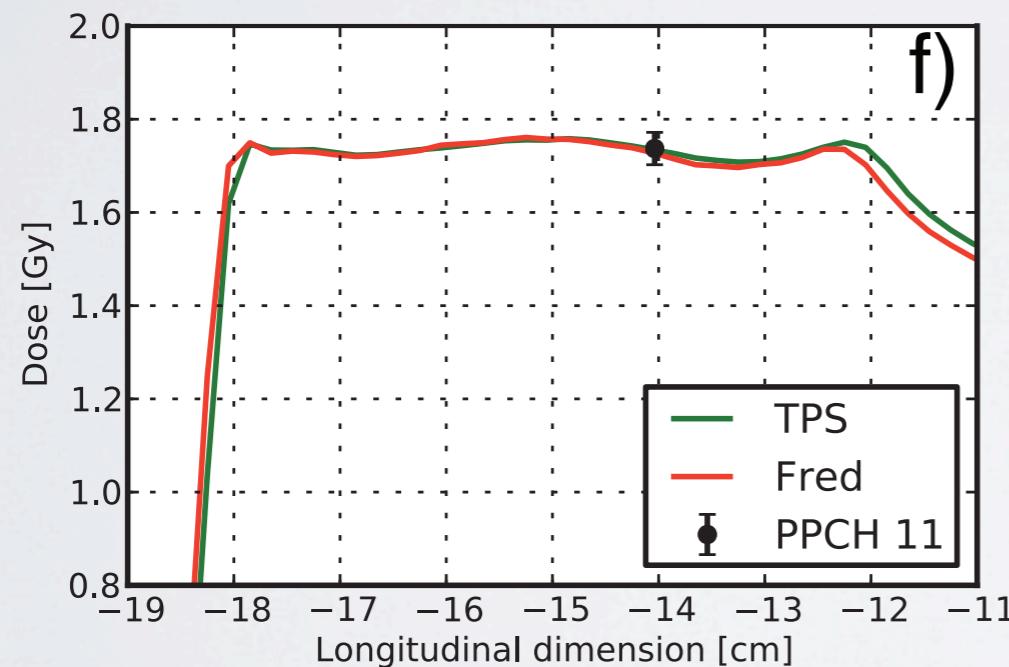
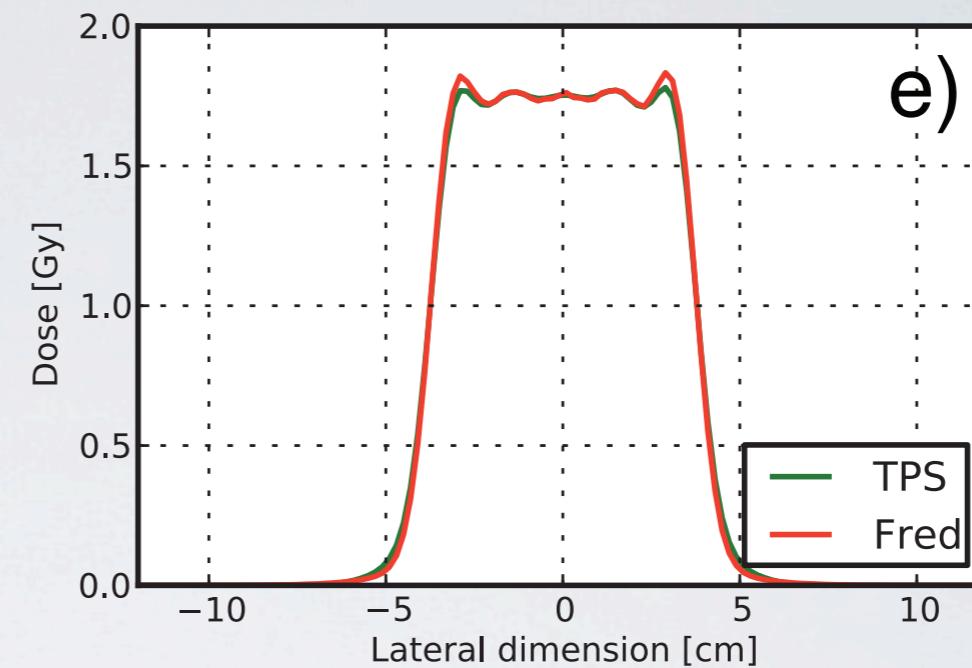
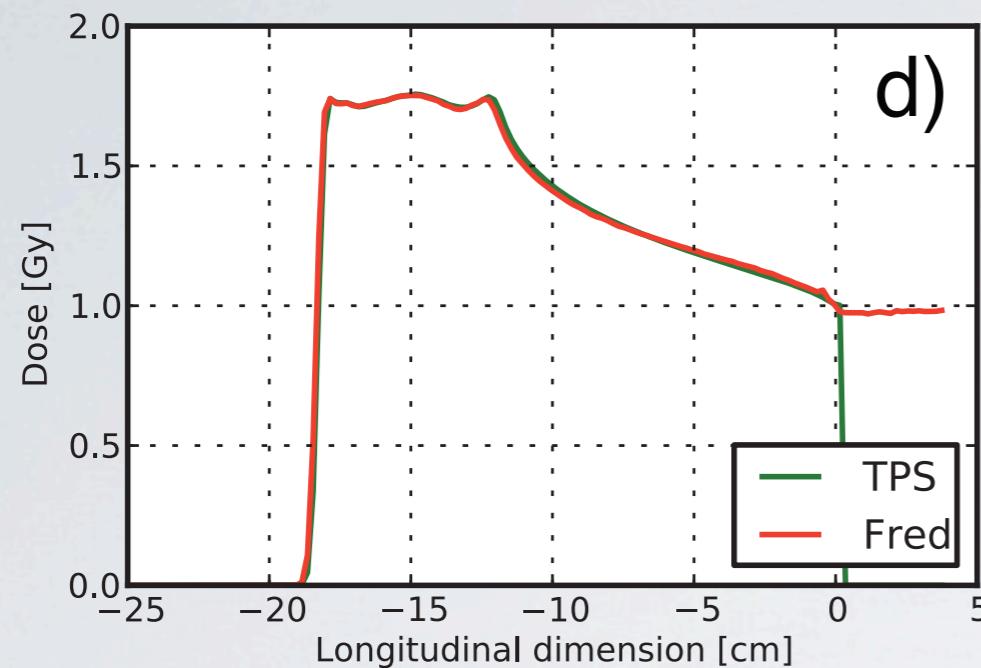
c)



Fred  
dose  
map

PTW MP3-P water phantom

# SOBP QA - 2



dose profiles and measured data point

# SOBP QA - 3

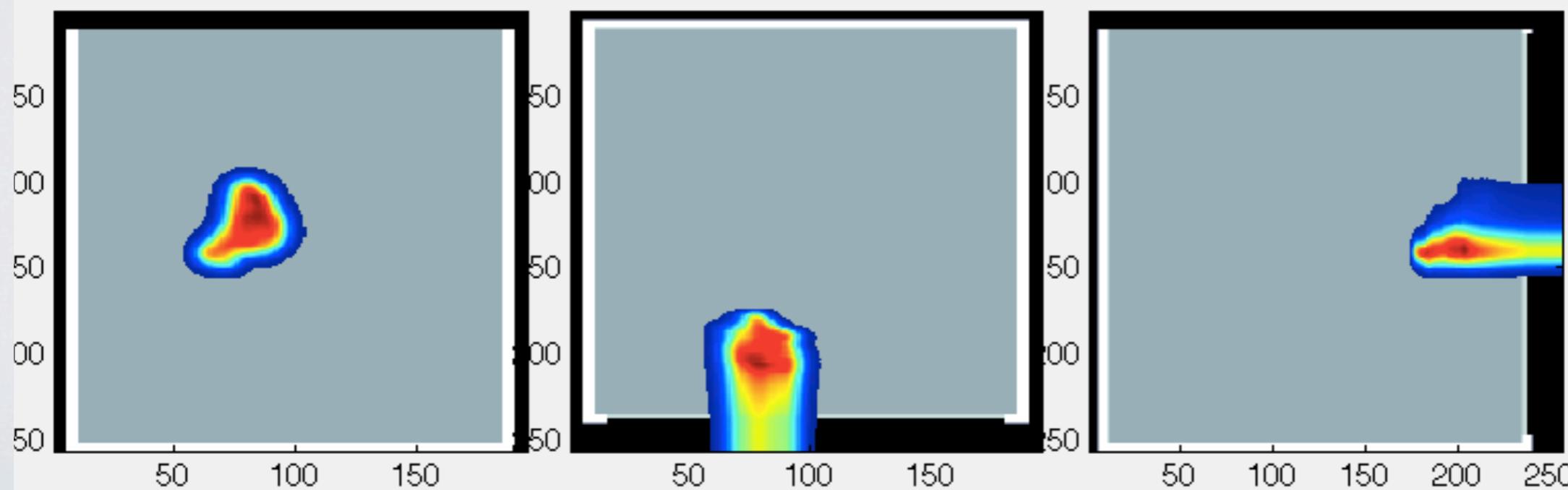
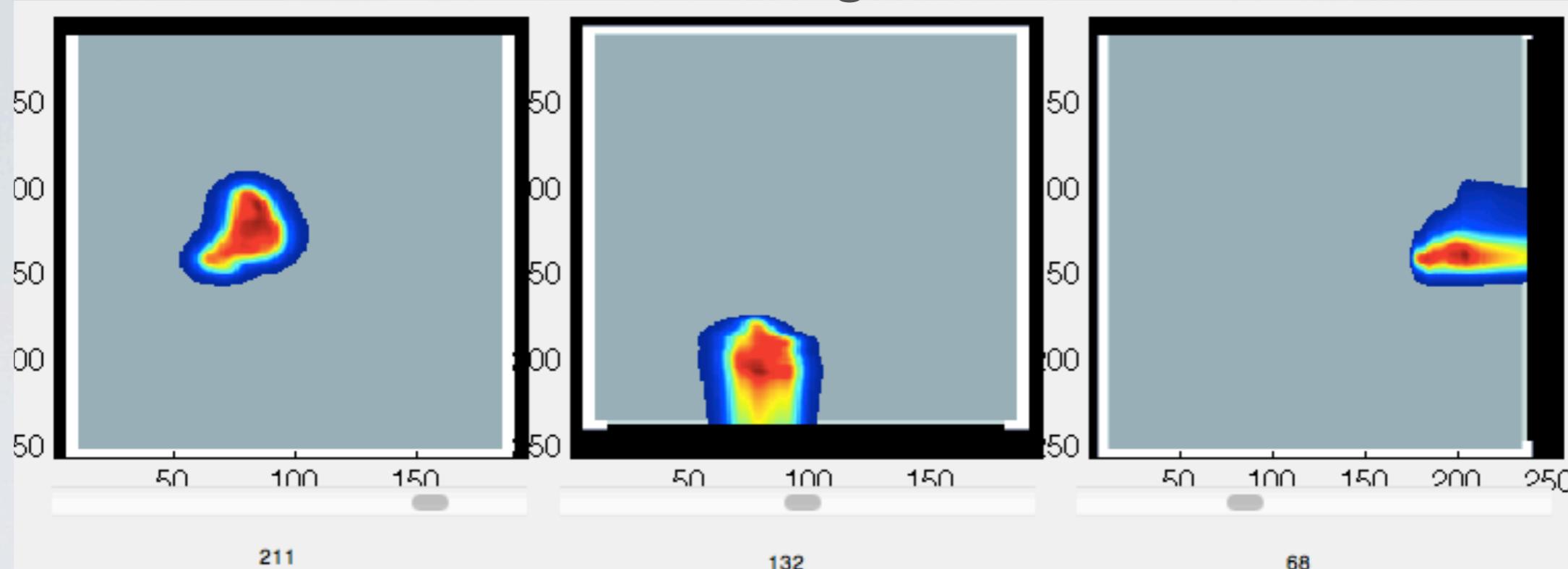
cube	<i>N</i>	zero shift				best shift			
		TPS		FRED		TPS		FRED	
		mean dose diff	stdev	mean dose diff	stdev	mean dose diff	stdev	mean dose diff	stdev
<b>6x6x6 cm<sup>3</sup> 9 cm depth</b>	11	-1.89 %	0.72%	1.27% 0.65%		-1.60% 1.08%		-0.56% 1.32%	
<b>6x6x6 cm<sup>3</sup> 15 cm depth</b>	11	-0.44%	0.55%	0.06% 0.57%		0.0% 0.62%		0.0% 0.58%	
<b>6x6x6 cm<sup>3</sup> 21 cm depth</b>	11	-0.16%	0.47%	0.41% 0.85%		0.0% 0.38%		0.06% 1.13%	
<b>3x3x3 cm<sup>3</sup> 21 cm depth</b>	10	-0.32%	1.78%	0.65% 2.03%		0.0% 1.27%		0.0% 1.48%	
<b>6x6x6 cm<sup>3</sup> 27 cm depth</b>	11	-0.29%	0.48%	1.57% 0.60%		-0.15% 0.88%		1.4% 0.8%	

protocol acceptance threshold = 5 %

# Patient verification plan

gamma-index 99.6% @ 2mm/2%

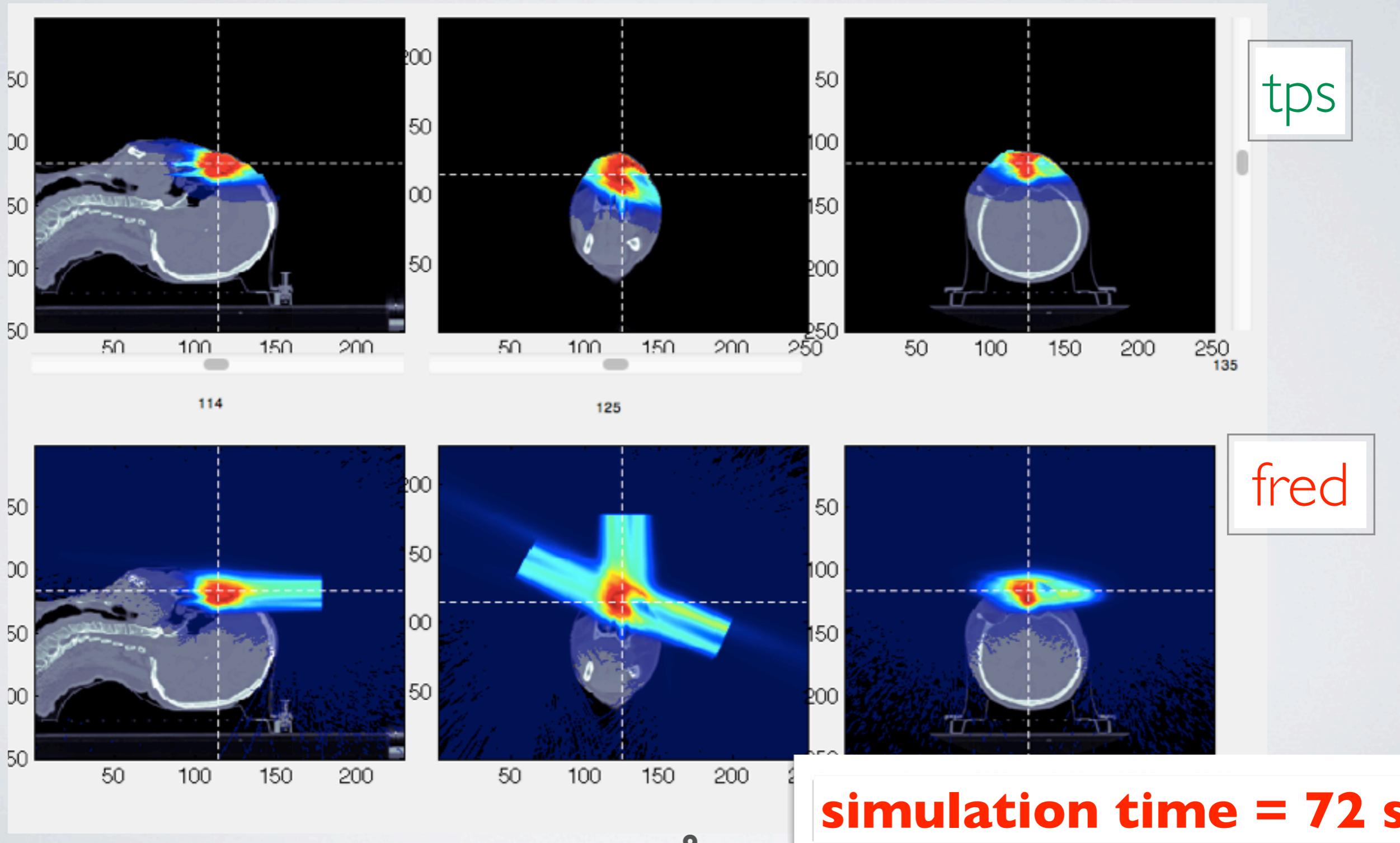
gamma-index 96.7% @ 1mm/1%



# Patient recalculation plan

recalculation at 1% = 700  
million primary protons

gamma-index 97% @ 2mm/2%  
gamma-index 92% @ 1mm/1%



# Hardware and Performance

		Threads	primary/s	$\mu\text{s}/\text{primary}$
CPU <sup>a</sup>	full-MC *	1	0.75 k	1340
	FRED	1	15 k	68
	FRED	16	50 k	20
	FRED	32	80 k	12.5
GPU	FRED	1 GPU <sup>1</sup>	500 k	2
	FRED	2 GPU <sup>2</sup>	2000 k	0.5
	FRED	4 GPU <sup>3</sup>	20000 k	0.05

Table A1: Computing times for different hardware architectures.

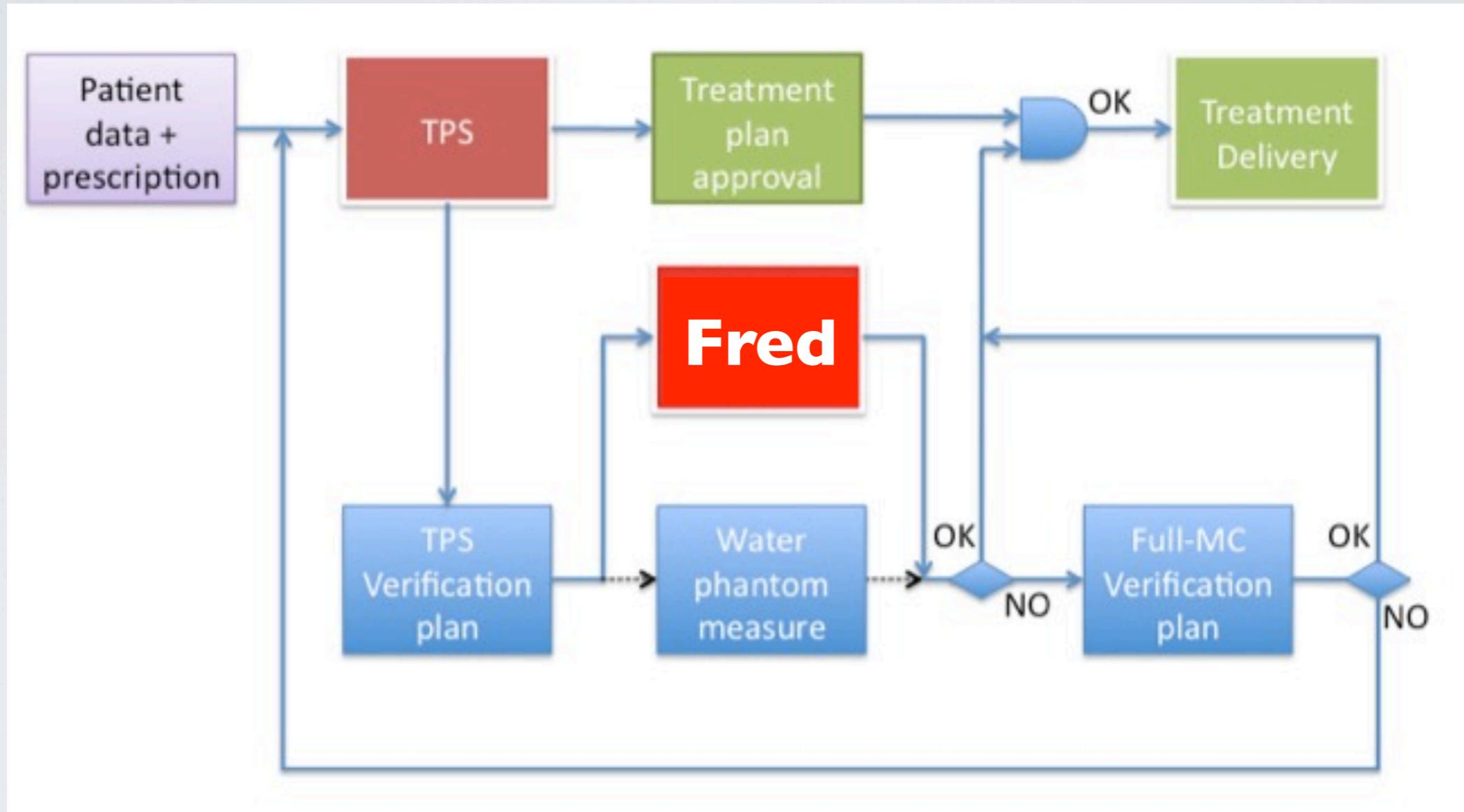
<sup>a</sup> motherboard with two Intel® Xeon E5-2687 8-Core CPU at 3,1GHz

<sup>1</sup> LAPTOP: Apple® MacBook Pro with one AMD® Radeon R9 M370X.

<sup>2</sup> DESKTOP: Apple® Mac Pro with two AMD® FirePro D300.

<sup>3</sup> WORKSTATION: Linux box with four NVIDIA® GTX 980.

# Proton Patient QA



# **Consuntivo attività Fred - WP II e prospettive**

- completato il modello “protoni in acqua”
  - fine-tuning e confronto con TPS
  - accordo Fred-TPS ampiamente entro le soglie del protocollo QA per i SOBP
  - ricalcolo piani di verifica in acqua in meno di 2 min
  - ricalcolo fast-MC piano paziente
- 
- commissioning per il proton patient QA
  - curve di calibrazione CNAO per paziente
  - validazione ricalcolo paziente e dose biologica