

Data reduction and analysis of SPS data



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FOR UA9 COLLABORATION

OUTLINE

2

- I. Synchronization
- II. Alignment procedures
- III. Qualitative analysis of dispersive area scans
- IV. Conclusions

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3

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SYNCHRONIZATION

4

Problems:

- All the UA9's devices are logged in different ways:
 - Different acquisition time.
- Logging of machine parameter completely uncorrelated from all other acquisitions.



Solution:

- ✓ Take all the different files and make like a “tetris”:
 - I. Choose the time range in which all the files have data.
 - II. Synchronization of the all data by Unix Timestamp.

SYNCHRONIZATION

5

What the synchronization program does:

- ✓ Starts from the higher initial Timestamp.
- ✓ Writes every parameter in a ROOT file every second.
- ✓ When data are not present, they are replaced with the previous acquisition.
- ✓ Stops at the lower final Timestamp.

File 1	File 2
12:40:00	
12:40:01	12:40:01
	12:40:02
12:40:03	12:40:03

ROOT file

12:40:01	12:40:01	12:40:01
12:40:02	12:40:01	12:40:02
12:40:03	12:40:03	12:40:03

After that we have a synchronized ROOT file containing:

- Acquisition time.
- Acquisitions of all the detectors (Scintillators, GEM, BLM,...)
- Positions of all mobile devices (Crystal, Collimator, Absorber,...)
- All the SPS parameters (Beam Intesity, Tune, Orbit,...)

**We have the complete knowledge of what happened in SPS and in UA9 apparatus:
we can make all the correlation plots that we need!**

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6

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What we have:

Position measured from garage position

What we want:

Relative position from the beam



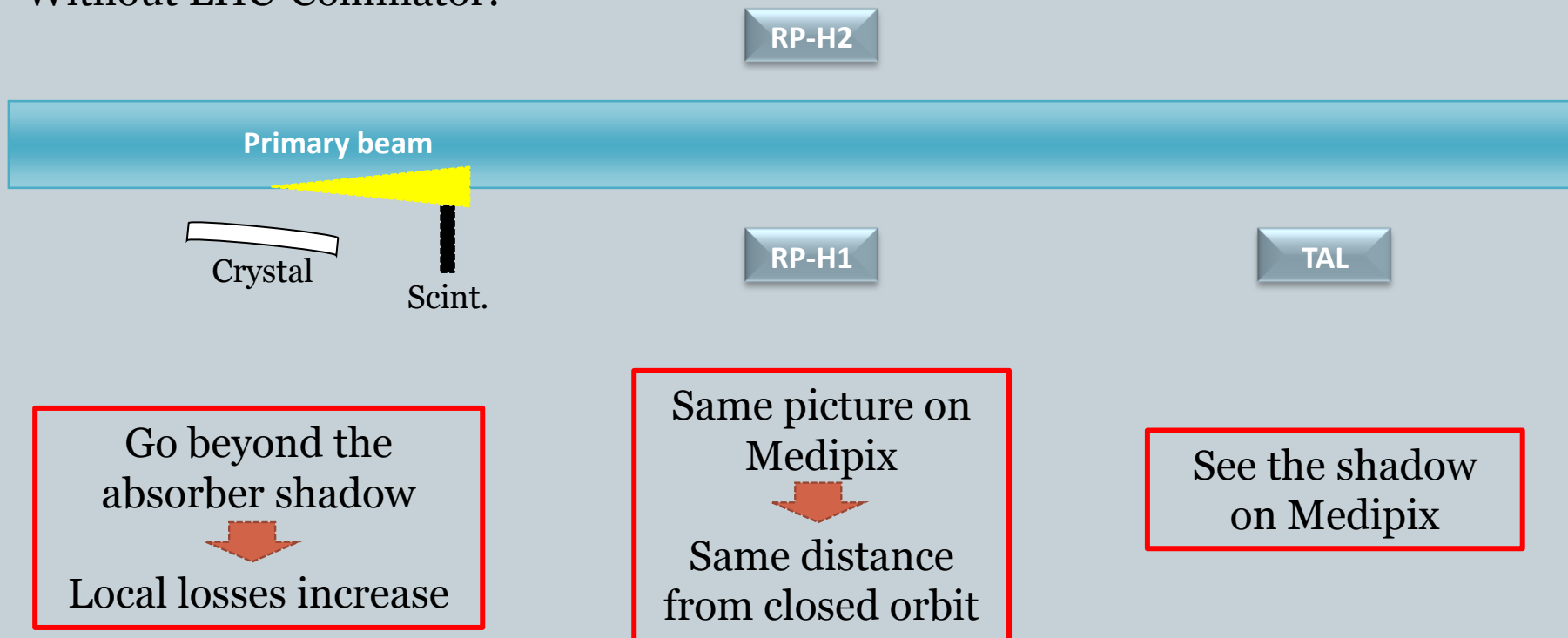
Needed a alignment
with respect to the beam

ALIGNMENT

7

Two different procedures depending on the presence or not of the LHC-Collimator.

Without LHC-Collimator:

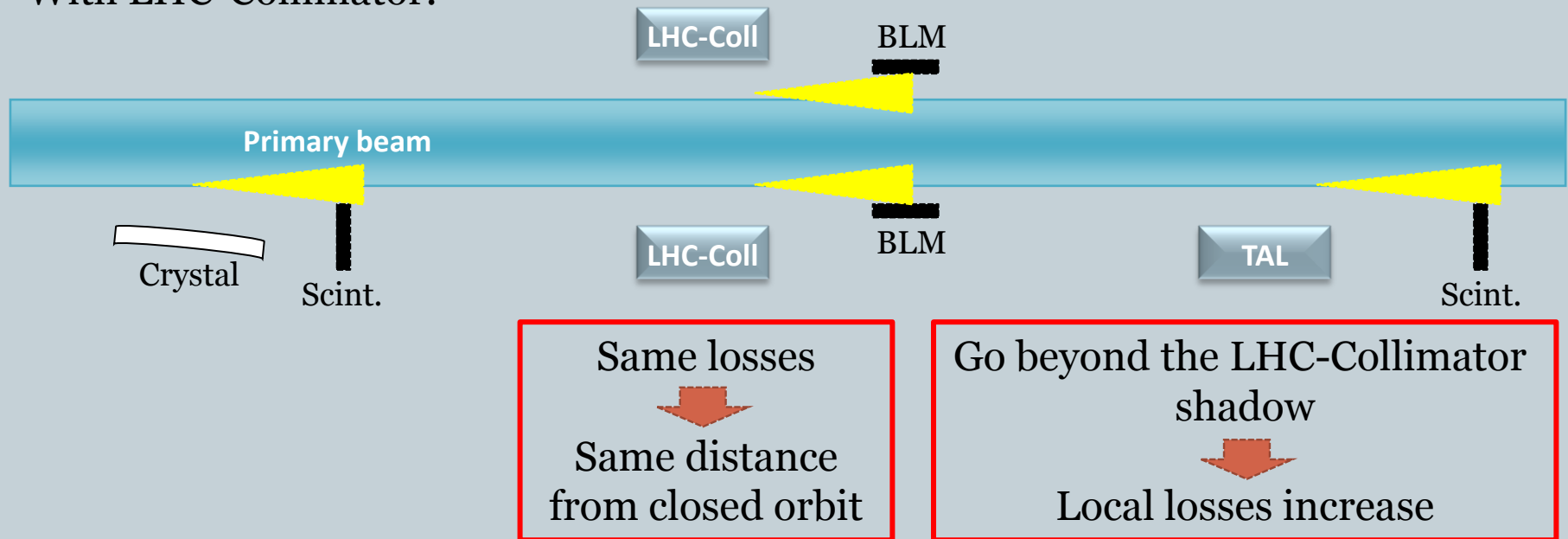


ALIGNMENT

8

Two different procedures due to the presence or not of the LHC-Collimator.

With LHC-Collimator:



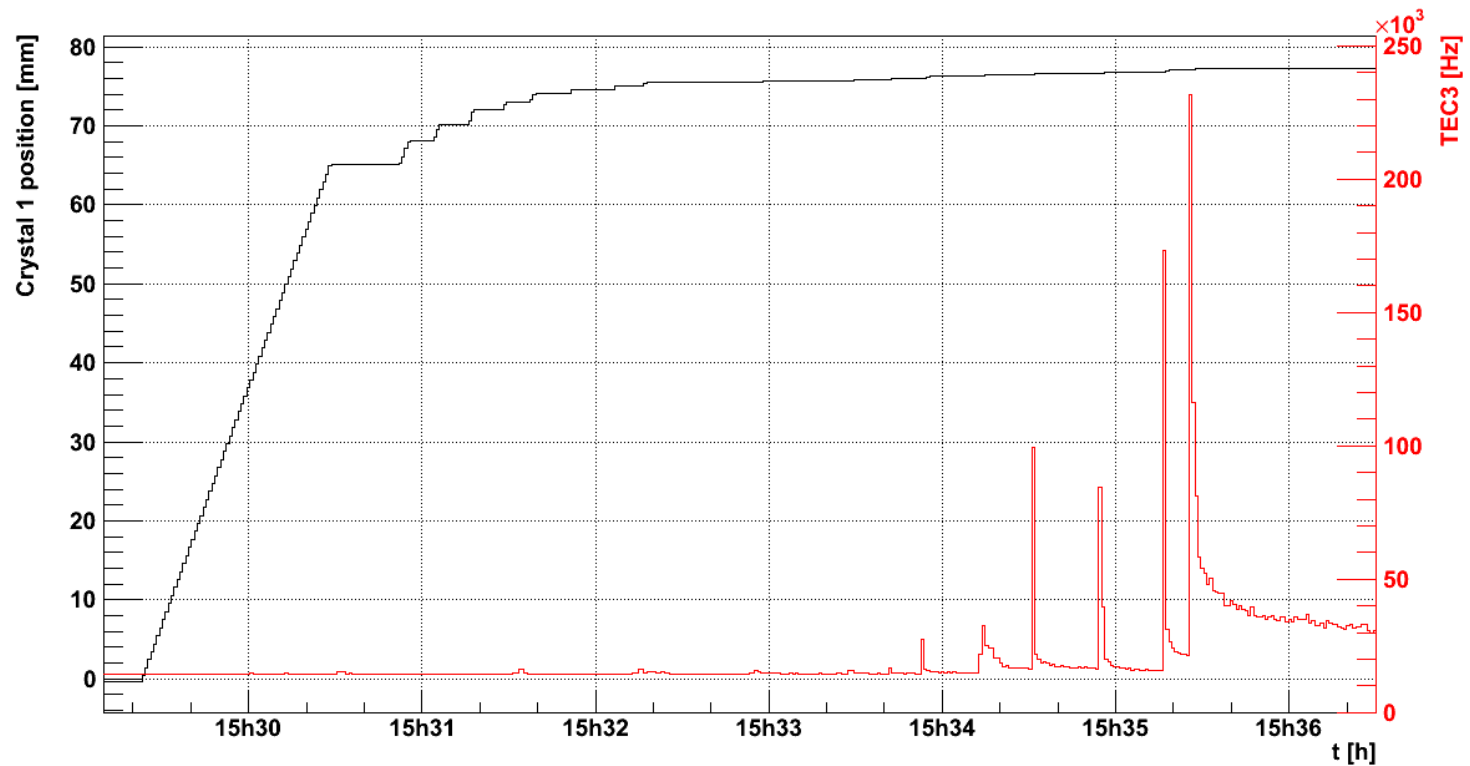
Basic configuration after the alignment:

- Insert crystal 0.5mm inside respect the alignment position.
- Retract absorber of channeled beam of 1.5mm respect the alignment position.

ALIGNMENT

9

Example of crystal alignment with LHC-Collimator:



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10

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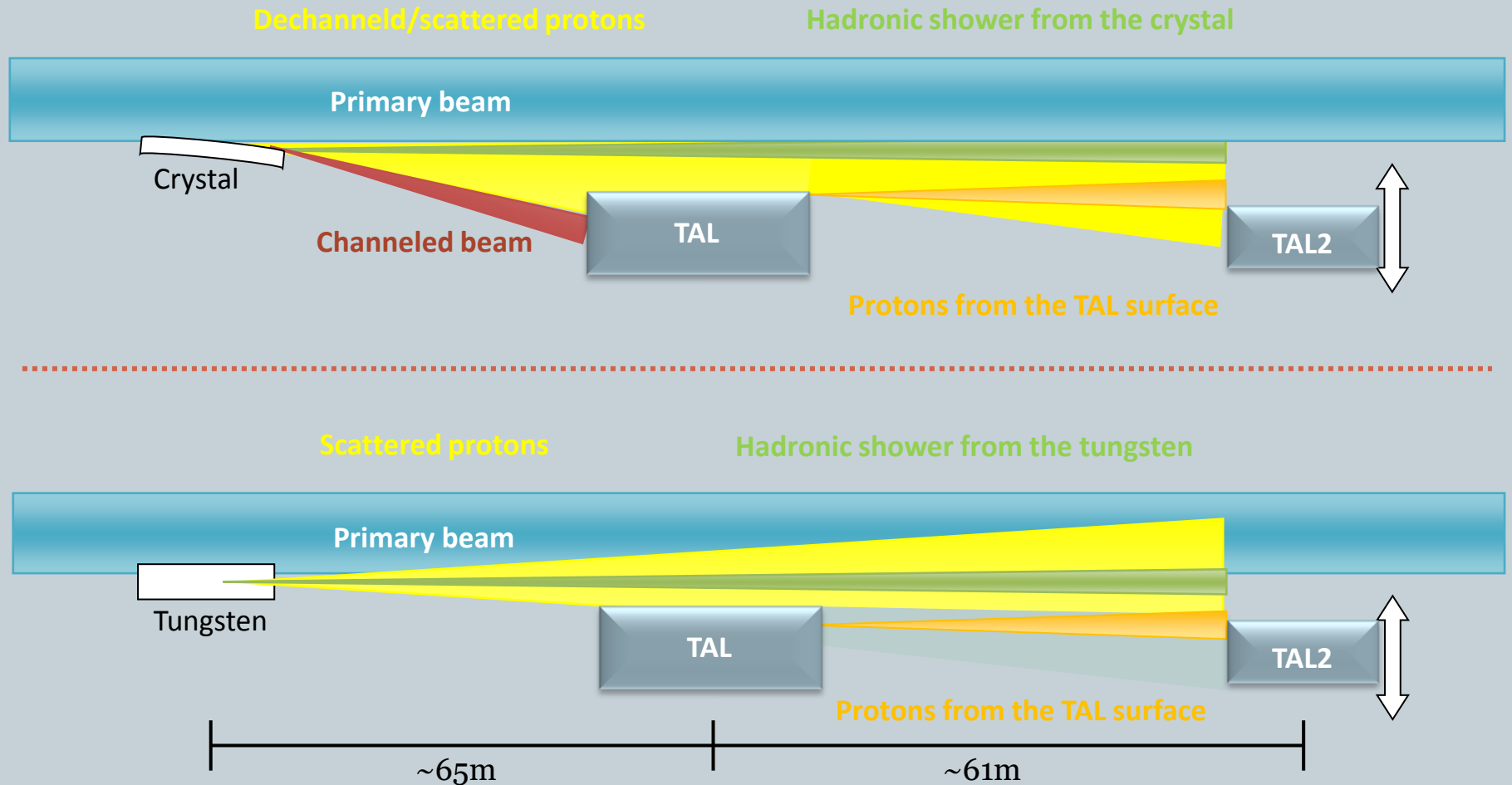
What is the effect of the collimation process, on the shape of beam?



Beam tails scan

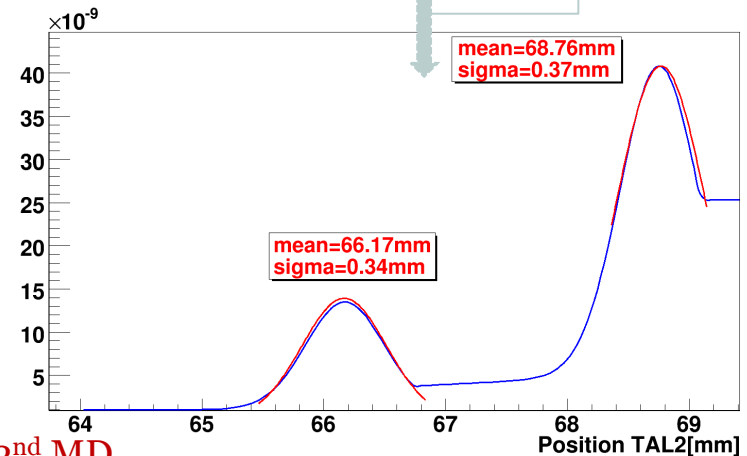
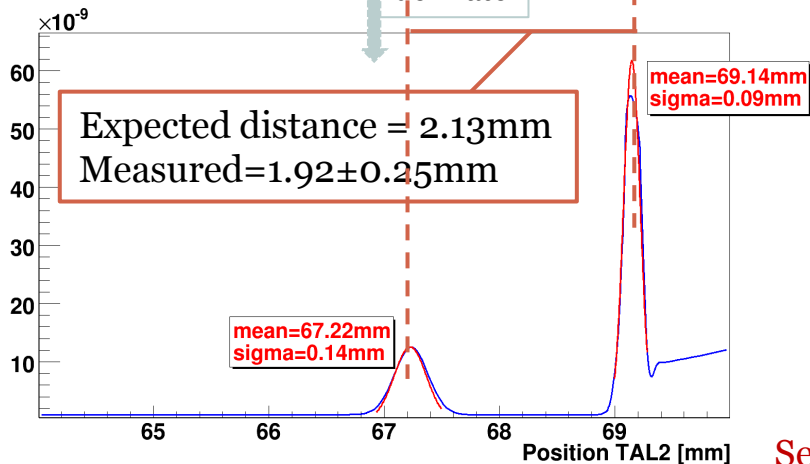
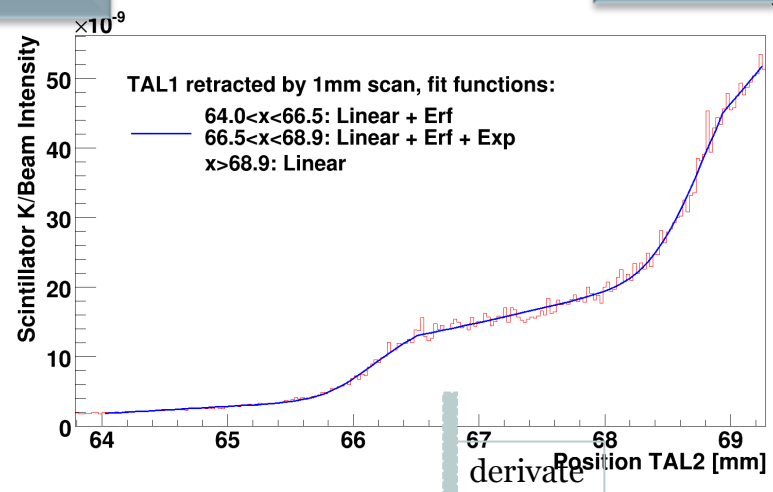
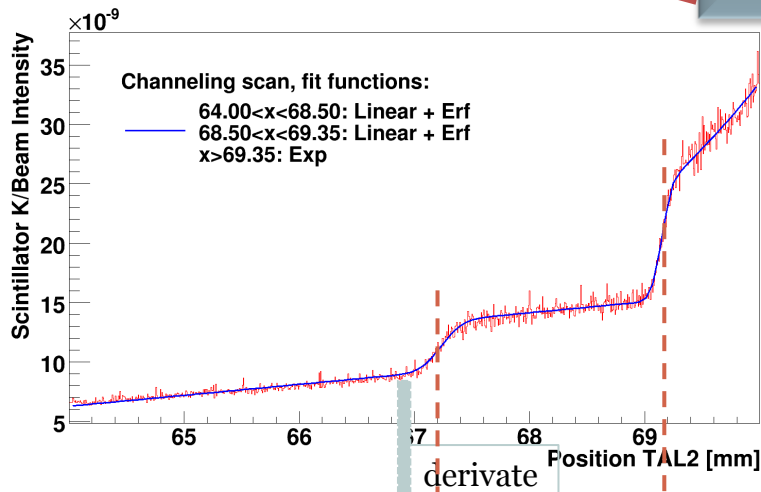
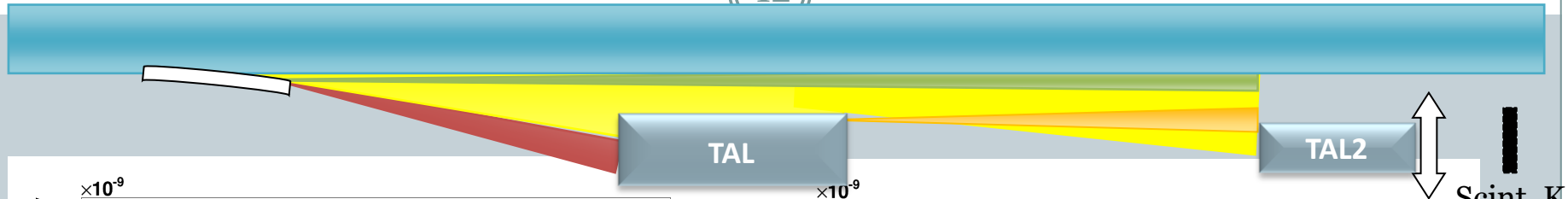
DISPERSIVE AREA SCANS

11



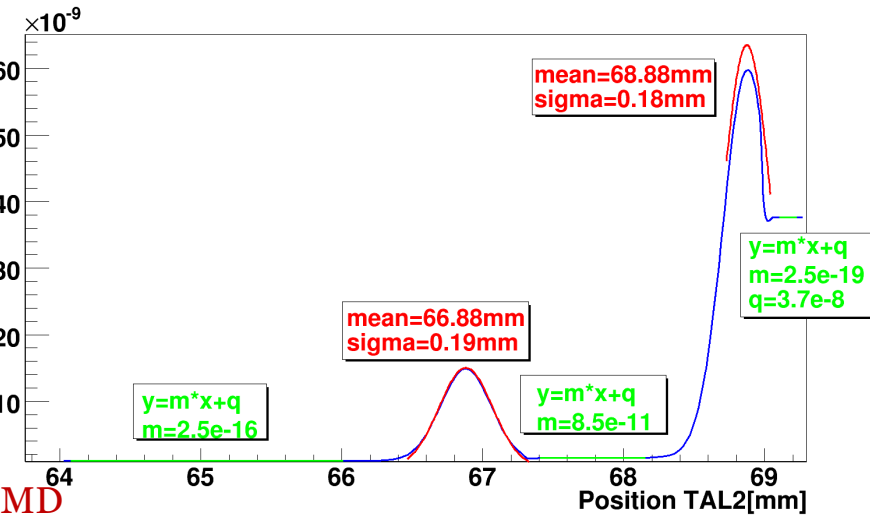
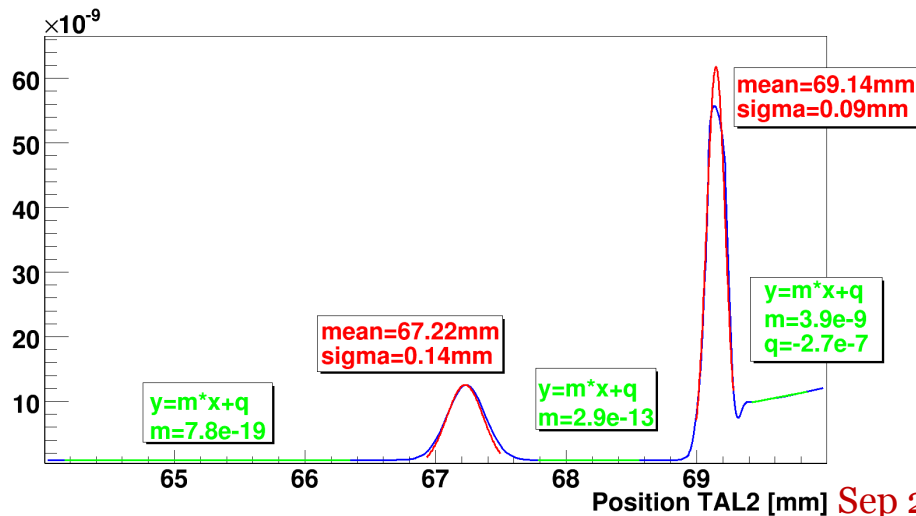
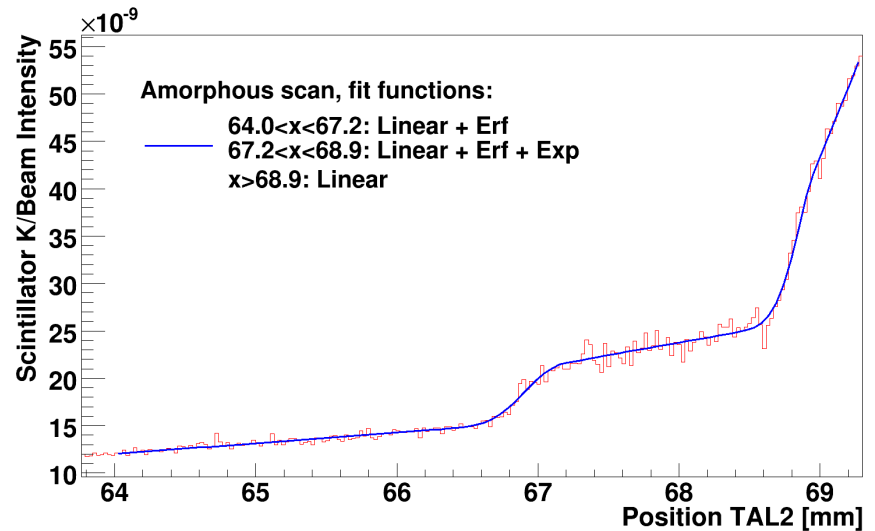
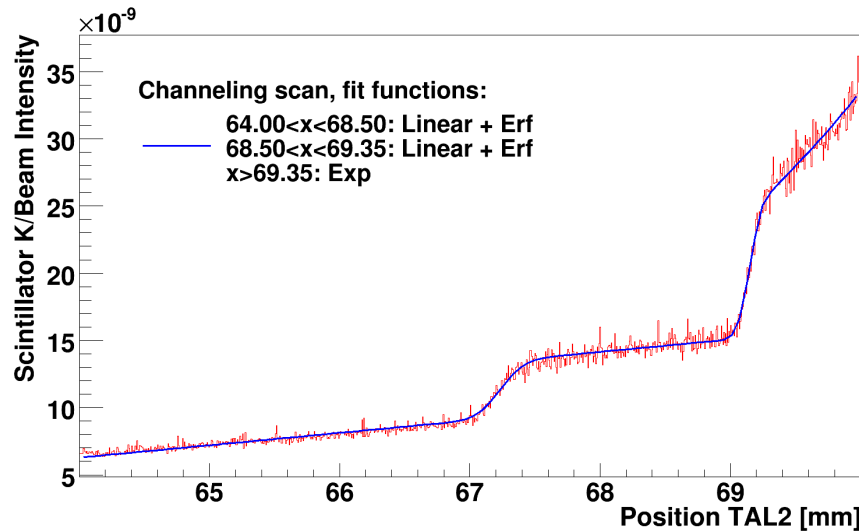
DISPERSIVE AREA SCANS

12



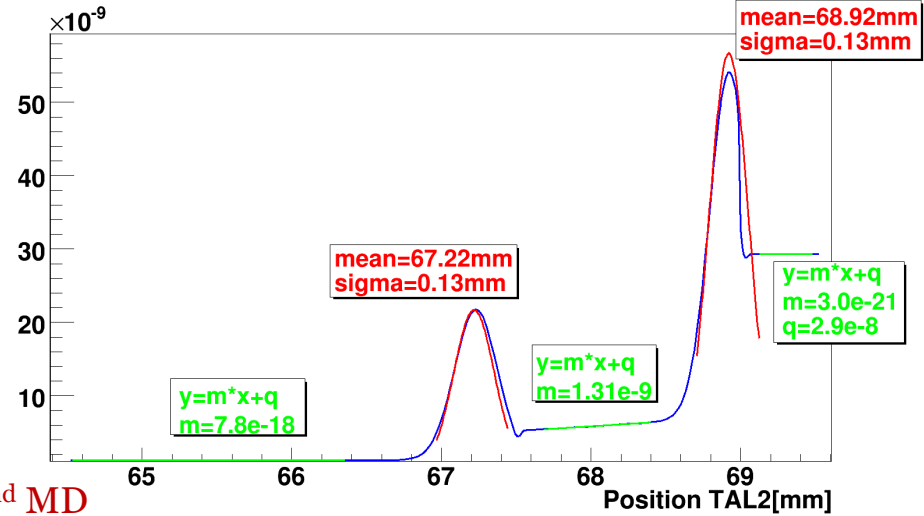
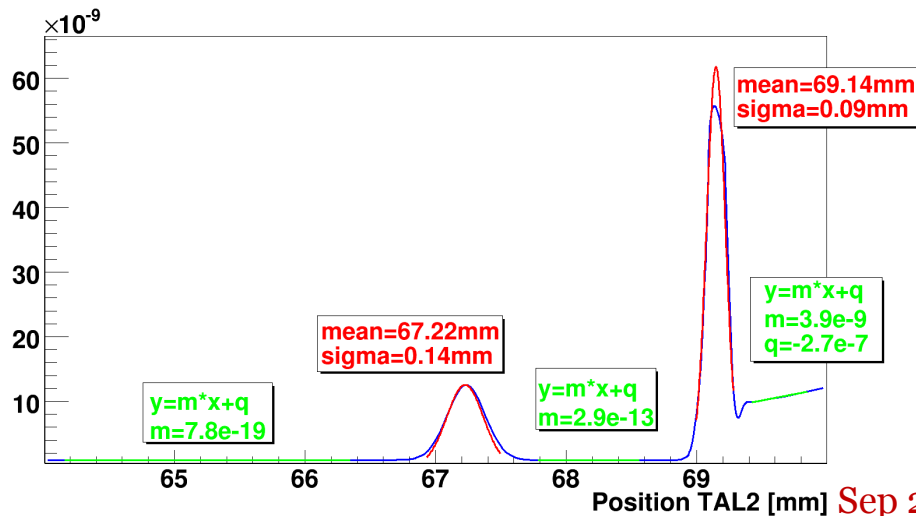
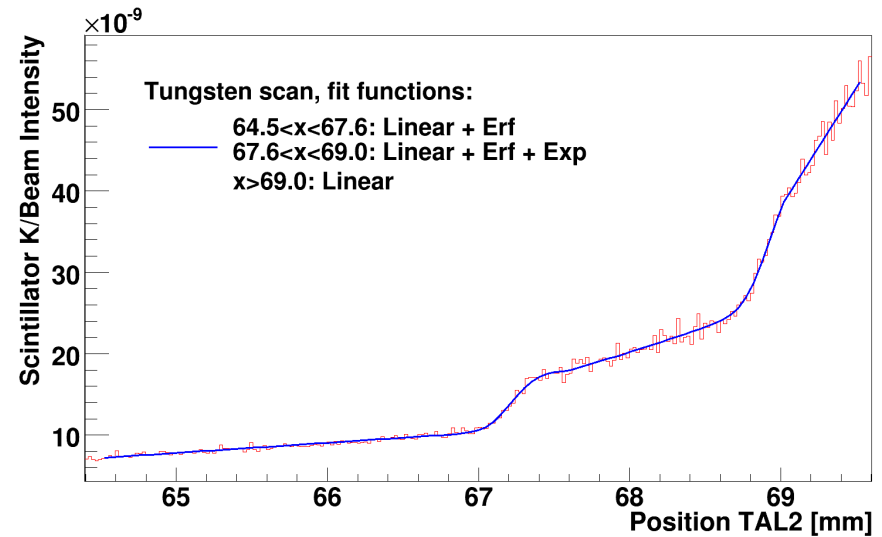
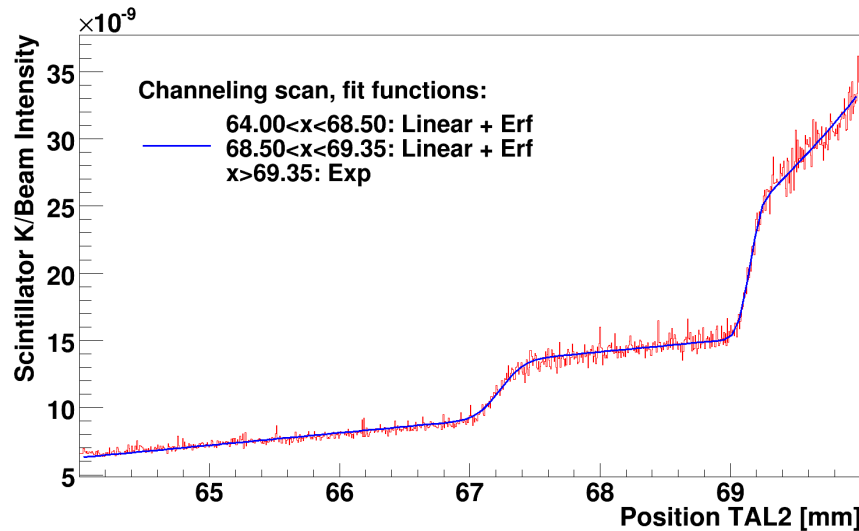
DISPERSIVE AREA SCANS

13



DISPERSIVE AREA SCANS

14



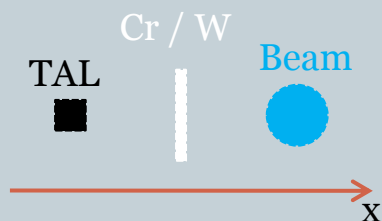
Sep 2nd MD

DISPERSIVE AREA SCANS

15

Derivative slope %	Before TAL	Between TAL & Crystal	Between Crystal & Beam
Channeling	$< 10^{-9}$	$3 \cdot 10^{-4}$	3.9
Amorphous	$< 10^{-6}$	0.1	$< 10^{-9}$
Tungsten	$< 10^{-8}$	1.3	$< 10^{-11}$

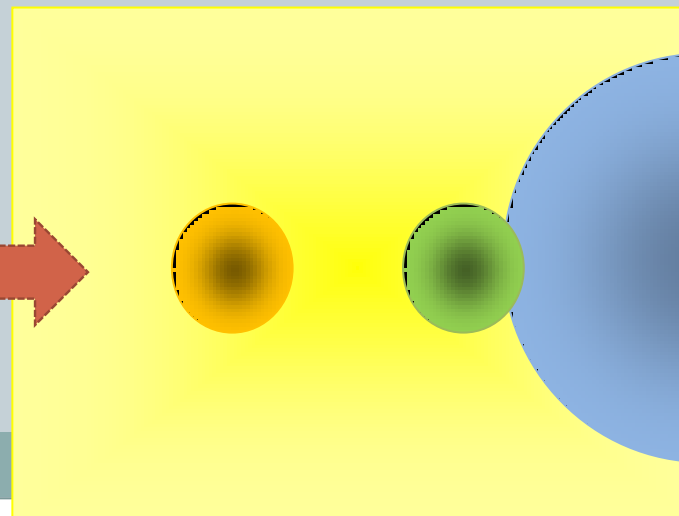
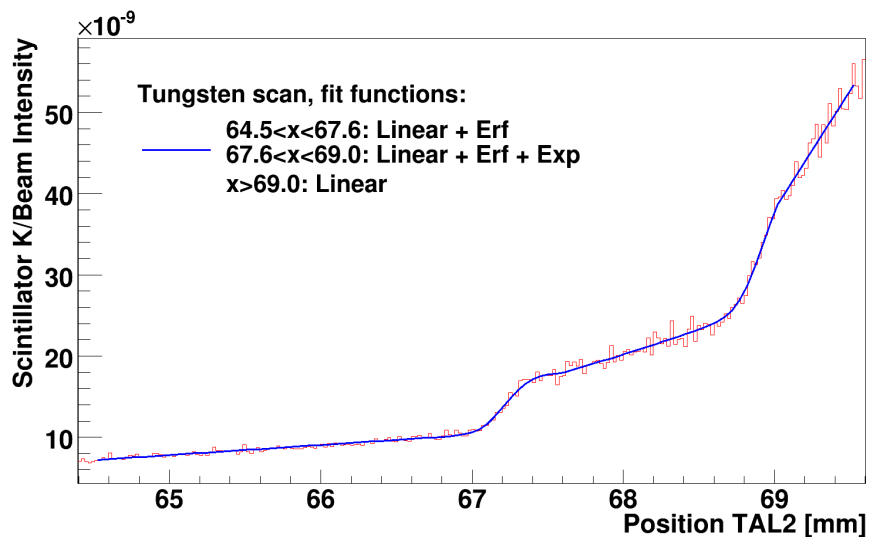
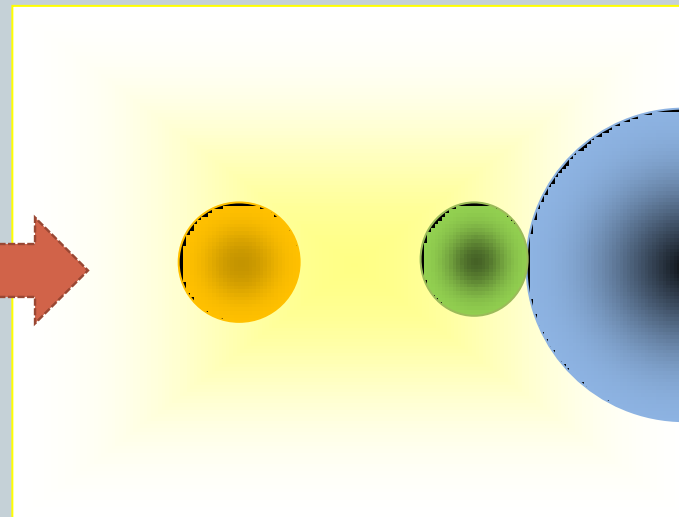
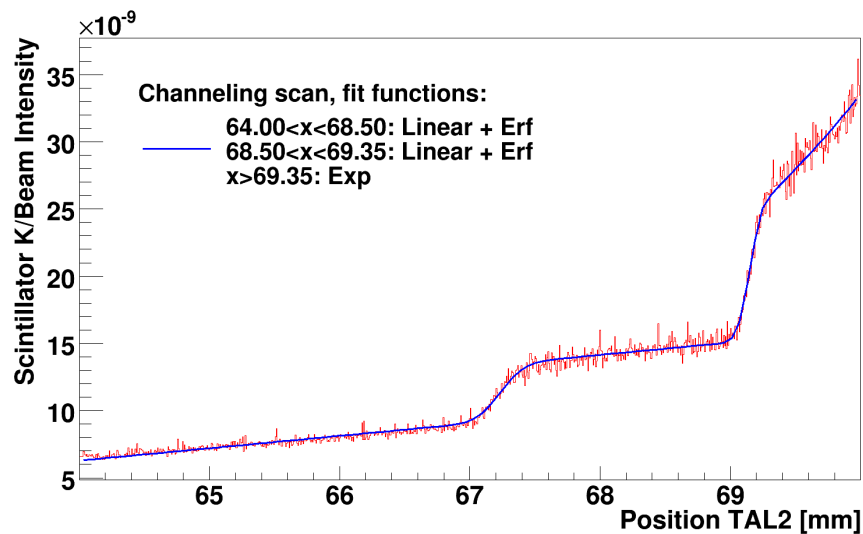
Normalized particles density (10^{-9})	Before TAL	After TAL	After Crystal
Channeling	0.9	0.9	10
Amorphous	1	1.5	37
Tungsten	1.2	5.4	29



Normalized counts after Crystal (10^{-9})	
Channeling	25
Amorphous	40
Tungsten	40

DISPERSIVE AREA SCANS

16



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17

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CONCLUSIONS

18

- ✓ With this synchronization program, we have the complete picture of what happen during the run, second by second.
- ✓ The alignment procedure is crucial and challenging: we developed two methods (with or without LHC-Collimator) to align the devices with a good precision in every condition.
- ✓ First qualitative analysis of the dispersive area scans, shows that with crystal collimation we seems have a more clean and definite beam, with respect to amorphous collimation.

For the future:

- Do an online synchronization during data taking.
- Collect more dispersive area scans, also with a larger range.
- Do a FLUKA simulation for dispersive area scans.

ALIGNMENT

19

Two different procedures due to the presence or not of the LHC-Collimator.

Without LHC-Collimator:

1. Close both sides (one at time) of the Roman Pot.
2. Close the absorber of channeled beam.
3. Open both Roman Pot sides. Absorber stays in the same position.
4. Approach the beam with one mobile device.
5. Retract the device.
6. Repeat points 4 & 5 for each mobile device.

Same picture on
Medipix



Same distance
from closed orbit

See the shadow
on Medipix

Cross the absorber shadow



Local losses increase

Basic configuration after the alignment:

- Insert crystal 1mm inside respect the alignment position.
- Retract absorber of channeled beam of 2mm respect the alignment position.

ALIGNMENT

20

With LHC-Collimator:

1. Close both jaws (one at time) until they touch the beam.] →
2. LHC-Collimator stays closed.
3. Approach the beam with one mobile device.] →
4. Retract the device.
5. Repeat points 3 & 4 for each mobile device.
6. Open completely both jaws of LHC-Collimator.

Same losses



Same distance
from closed orbit

Cross the LHC-Collimator shadow



Local losses increase

For a better estimation of the alignment position during the offline analysis we need few little steps ($\sim 100\mu\text{m}$) after touching the beam.