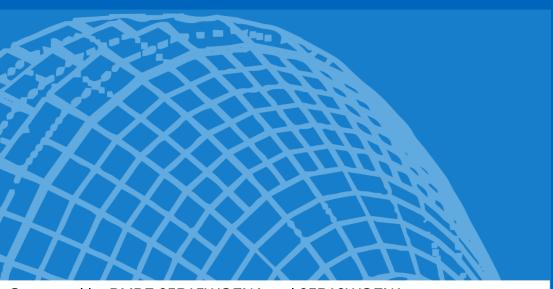


ПΠ

Time Stitching and Eventbuilding for Experiment S455 (and S515)



Supported by BMBF 05P15WOFNA and 05P19WOFN1.

The results presented here are based on the experiment s444/s473, which was performed at the beam line/infrastructure Cave C at the GSI Helmholtzzentrum für Schwerionenforschung, Darmstadt (Germany) in the frame of FAIR Phase-0.

GEFÖRDERT VO







Tobias Jenegger

R3B Collaboration Meeting 2021

First Unpacking Issues

Event Building before/after 2020

Recipe for Time Sorting and Stitching

First Results from S455 (p,2pf) reaction

TUM Members:

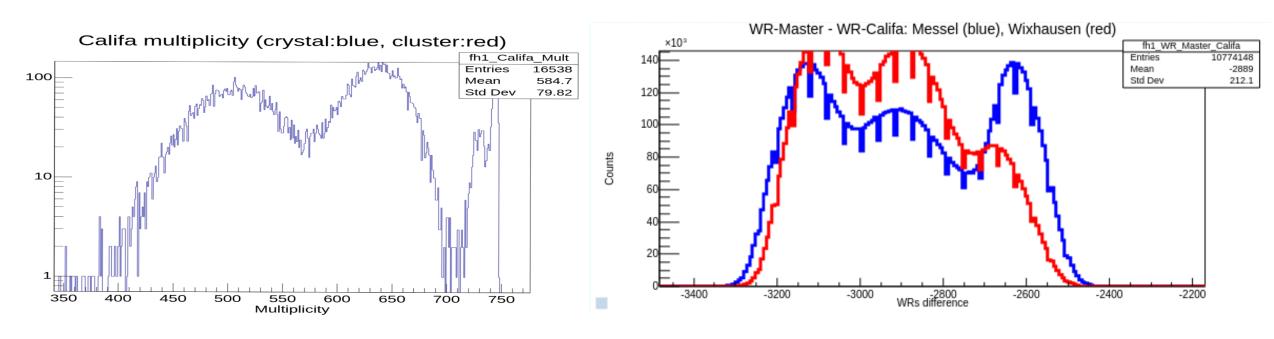
Roman Gernhäuser, Lukas Ponnath, Philipp Klenze, Tobias Jenegger



Issues in online/offline Data Analysis starting from S455



When using the known unpacking tools from previous experiment we got:



Events with hit-multiplicities > 500!?

Sharp peak for strong correlation WRM- WRCalifa expected!



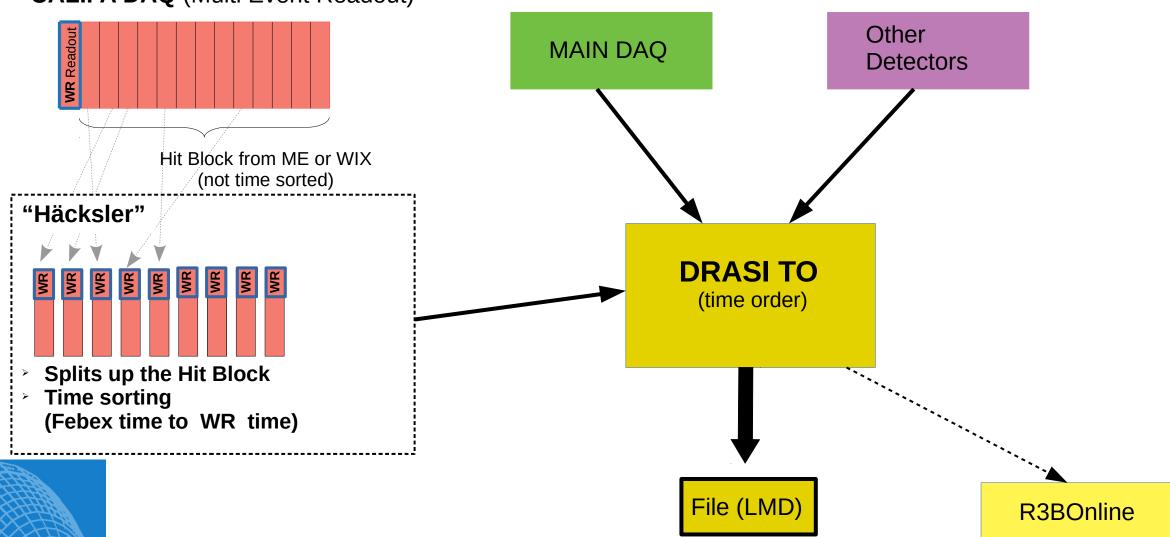
- → Is the data corrupted? Has something gone wrong during data acquisition?
- → How is data processed and stored to .lmd file?



Data Acquisition/Event Building until 2020



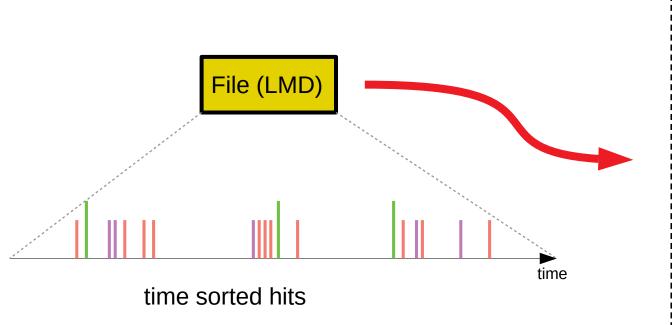
CALIFA DAQ (Multi Event Readout)





Offline Time Stitching until 2020

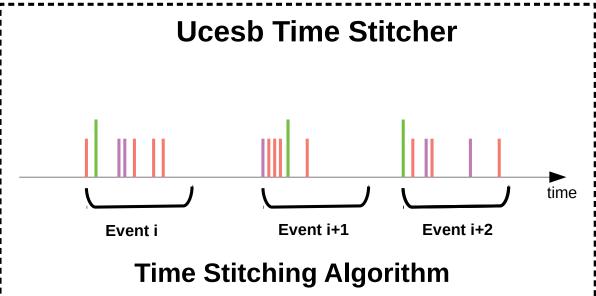




: main DAQ hit

: CALIFA hit

: other detector hit (e.g. MWPC1, TWIM,...)

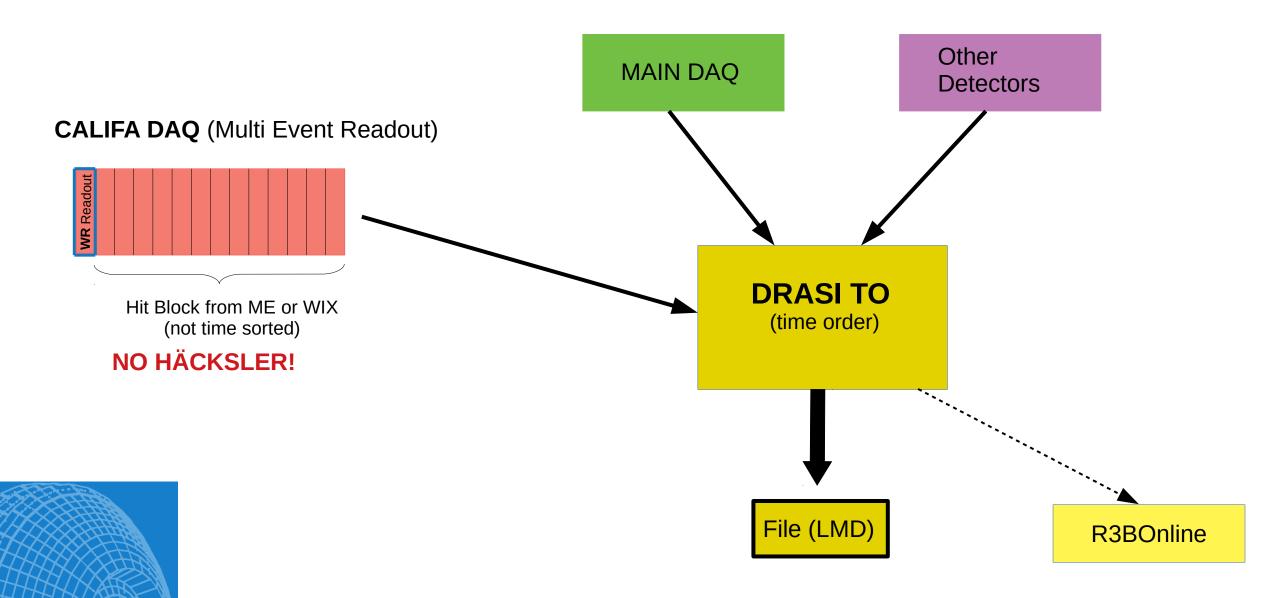


- First detected hit opens event window
- Time Stitcher does not distinguish between CALIFA, Main DAQ,.. hits
- Event window is closed after predefined time (eg. 8 µs)



Data Acquisition/Event Building since 2021







Output LMD File since 2021







- CALIFA Hits are collected to Hit Blocks (Messel or Wixhausen)
- Hit Blocks are sorted according to the WR Readout timestamp ()
- Hits inside Block are not time sorted!

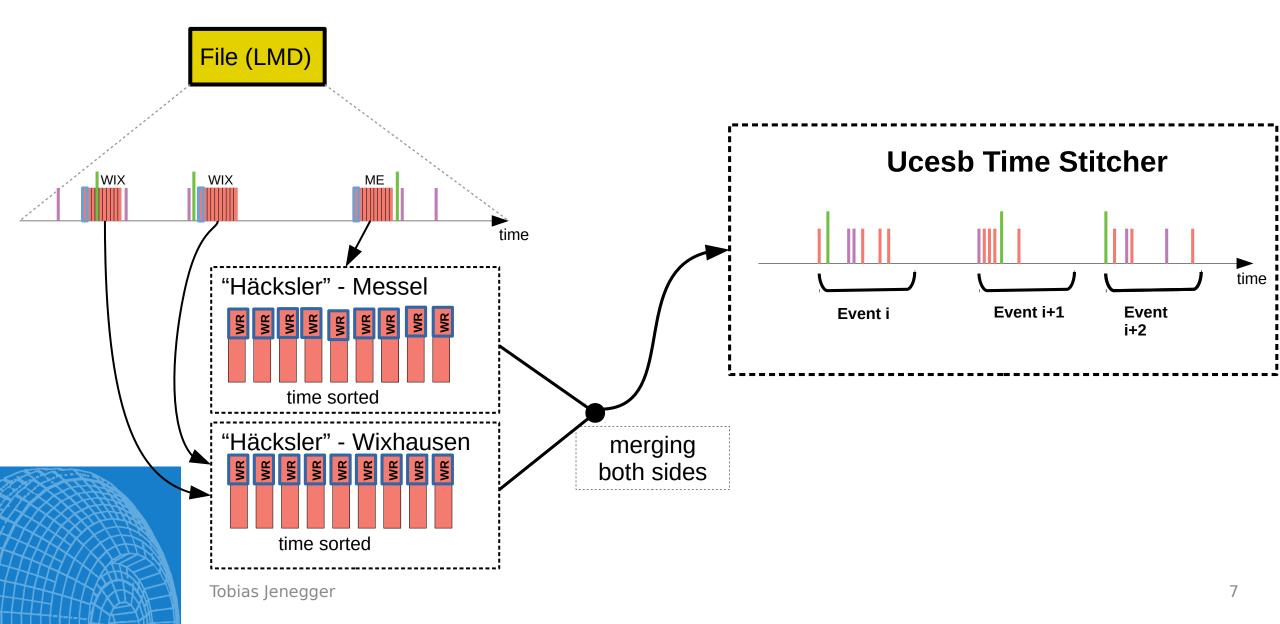


CALIFA single hit time sorting needed!



Offline Time Sorting and Stitching since 2021



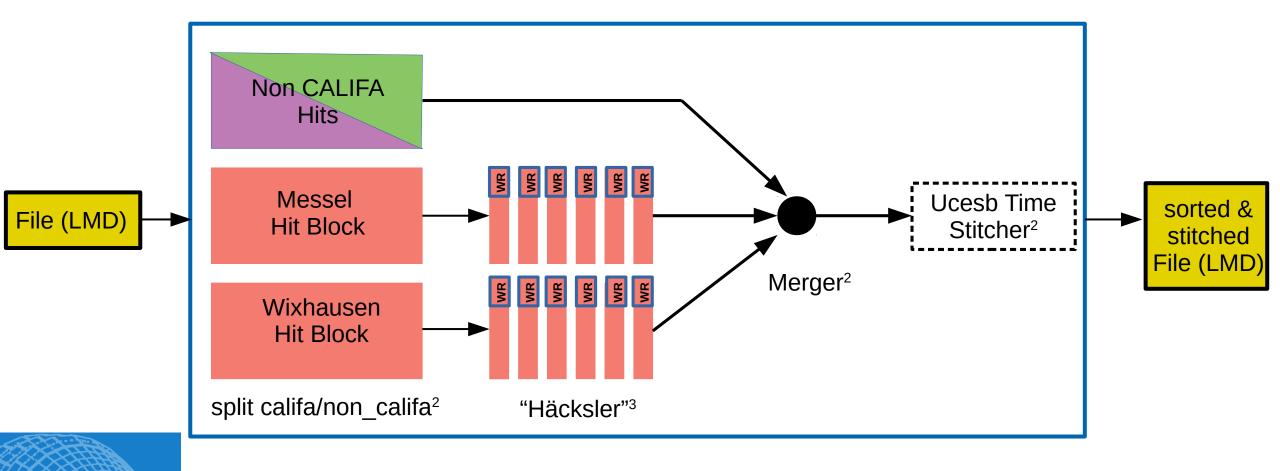


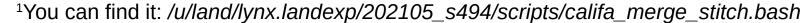


Time Sorting and Stitching with Philipp's script



califa_merge_stitch.bash1





²Unpacker: /u/land/klenze/ucesb_gcc6/empty/empty

³Unpacker: /u/land/landexp/202105_s494/x86l-76/daq_bootstrap/ucesb_wip/empty/empty



How to use the script



Location:

Always use the newest version you find in: /u/land/lynx.landexp/202105_s494/scripts/califa_merge_stitch.bash

Call Parameters:

- > an input, either a file path, stream of trans server
- > an output for the merged data (*.lmd file; " " for "no output; ucesb -server option)
- > an output for the stitched data (*.lmd file; " " for "no output; ucesb -server option)

Example:

Califa_merge_stitch.bash run10.lmd " " stitched_run10.lmd

Output .lmd stitched data

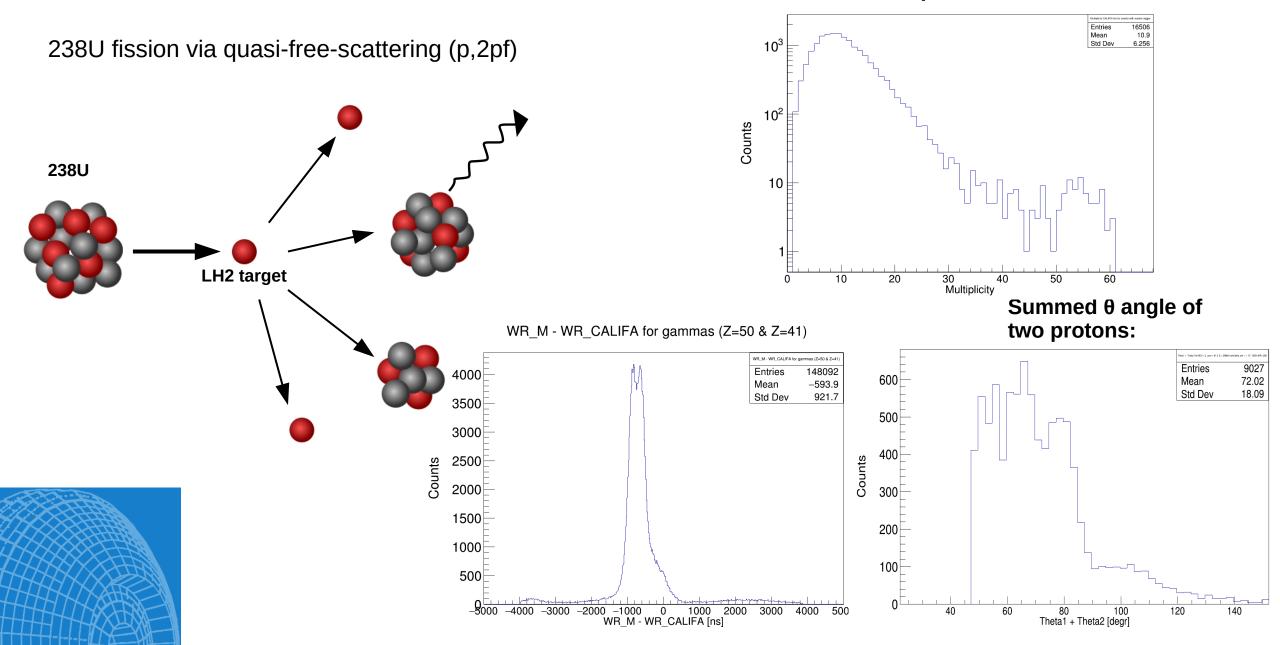
No output merged data



Looking again at S455 Data



Multiplicities:

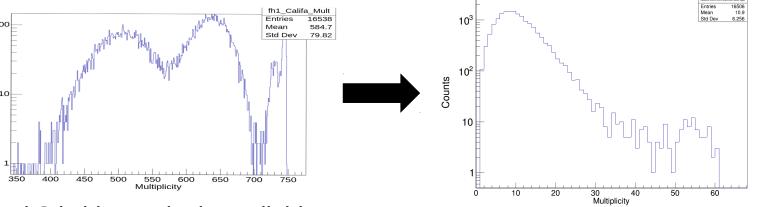




Summary & Outlook



We successfully unpacked the S455 data



- Well tested and user friendly Time Sorting and Stitching script is available: /u/land/lynx.landexp/202105_s494/scripts/califa_merge_stitch.bash
- > Same procedure for experiments after S455 (e.g. S515) → this has to be accurately tested
- > CALIFA upexps code was recently unified, experiment independent with one mapping file for all experiments











Thank you!

CALIFA @ Technical University of Munich (TUM)

Roman Gernhäuser, Lukas Ponnath, Philipp Klenze, Tobias Jenegger













Backup

