Nice to meet you! Federica Borgato

INFN - SESAME International School on EFFICIENT SCIENTIFIC COMPUTING ESC@SESAME







My past and current activities:

- I studied in Padova and got graduated with a thesis on TimeSPOT 3D silicon sensors. I performed the static characterization and dynamic characterization with 2MeV protons from National Legnaro Laboratories AN2000 accelerator.
- Now I am a PhD at the university of Padova and my activity is focused on R&D for the LHCb Upgrade:
 - I joined TimeSPOT test-beams at CERN SPS to establish the time resolution of such sensors, performing then the data analysis to extract the detection efficiency.
 - I joined RICH test-beams at CERN SPS to test the new electronic chain for the Upgrade 1b. I am currently performing the data analysis to extract the single photon time resolution of the MaPMTs and SiPM read out with the FASTIC asic.
 - I am performing the measurement of BR(Λ (dominant background of $R(\Lambda_c^*)$).

$$\Lambda_b^0 \to \Lambda_c^* D_s^{(*)} / \Lambda_b^0 \to \Lambda_c D_s^{(*)}$$



The LHCb RICH subsystem

LHCb relies on the Ring Imaging Cherenkov (RICH) detector system for the charged hadrons identification in a wide momentum range (2 - 100 GeV/c).

The Cherenkov light produced by the particles is redirected by an optical system towards the photodetector planes and outside the acceptance of the spectrometer.



Photon-Detection chain

MaPMT

CLARO chip

RICH commissioning: threshold scan analysis

Threshold scans analysis are a fundamental tool to **optimize the detector performance** since it allows to:

- choose the optimal threshold of operation
- fine tune the HV
- check the aging of the sensors
- Before the columns installation a full set of scans was taken at the CERN Comlab. This means:
- One integrated charge spectrum for each anode (threshold scan)
- One DAC scan for each anode, to correlate threshold and charge







RICH MaPMT Single Photon Time resolution

In 2022 two beam tests have been performed at CERN-SPS to test a prototype readout chain with fast-timing information.

The MaPMTs/SiPM are coupled with FastICs and read out by a TDC-in-FPGA.

The analysis is performed subdividing the data with respect to their ToT. In this way it is possible to create subsets of events sharing the same signal time length above threshold allowing to decouple the time resolution measurement from the time the time resolution is the time to the time the time the time to the time the time the time to the time the time to contribution.

Multiple CrystalBall fits are performed changing the fit range in order estimate the systematic uncertainty as well. In this way for each channel under study there will be a subset of 16 time resolution for each ToT bin.



each ToT bim.















Sicherheit in Technik und Chemie

29.05.2023

COMBINED EXPERIMENTAL AND SIMULATIONAL APPROACHES TO ACCESS RADIATION DAMAGE TO DNA-PROTEIN COMPLEXES

Marc Benjamin Hahn marc-benjamin.hahn@bam.de

Radiation damage to DNA



Ionizing radiation damage to DNA:

Cancer therapy and

medical imaging

DNA carrier of genetic

information



Hahn et al. Commun Chem 4, 50 (2021)

DNA-Protein interaction



DNA binding proteins

Tasks during DNA replication and repair

Interact directly with DNA



Hamon et al. Nucl. Ac. Res. 35, e58 (2007)





Experiments often give only access to microscopic dose.

But we are interested in fundamental processes.







- 1. Microscopic processes on molecular level?
- 2. Involved scattering processes?
- 3. Direct (scattering) & indirect (radicals) effects?



Access microscopic events by simulations



Particle scattering simulations with Geant4/TOPAS to access events on differents scales



Hahn J Phys Comm (2023)

Example I: Electron irradiated DNA in water



Determine microscopic dose-damage relation



Hahn et al. Phys. Chem. Chem. Phys. 2017a





Radiation damage and Nanoparticles in Cells: effects of nanoparticles and their location on dose in cell organelles



Zutta-Villate et al. Sci Rep (2020)

Dorothea Hallier

- Tihomir Solomun
- Heinz Sturm

Harald Seitz

DFG Deutsche
Forschungsgemeinschaft
DFG (Grant 442240902)

Contact: marc-benjamin.hahn@bam.de

9

Acknowledgements

Thank you for your attention!





Philipp Hans

philipp.hans@sesame.org.jo



I am a crystallographer. - I am interested in e.g. machine learning, approximation theory, automation, philosophy.

My work is about X-ray diffraction data processing and correction strategies any quality assessment, investigations of phase change materials for data storage applications, pharmaceuticals. This all involves scientific computing, where high performance computing can be an important aspect.

Hobbies: programming, sitting in the sun, climbing, kettlebell, socializing

I will be happy if you contact me to discuss.





Phase Change Materials (PCM) for PCRAMs

Ge-Te-Sb allov PCMs: •

used in memory devices: CD/DVD RW, PC RAM ٠

Reversibility between 2 states

+25 %

 Δ Optical reflectivity

crystalline

heating

250 300 35

T(°C)

cooling

200

Δ Resistivity -10⁴









technical realization



S. Raoux, Chem Rev. (2010)



Ge-rich composition shifts T and data retention to higher temperatures (important for automotive applications)



N-doping increases endurance



Next aim: time resolved + high throughput

measured with 2D detector -> reduction here: 1D patterns before and after in situ annealing



- patterns with a range of peaks.
- find out which peaks belong together ("which phase"; can be difficult)
- analyze the peaks to estimate ensemble effects and dynamics + quantification

Mixture of Ge and metastable cubic $Ge_2Sb_2Te_5$ (c-GST 225)

😭 pyFAI

naster

Search docs

General introduction

Example of usage

Cookbook recipes

Calibration of a diffraction setup using the Graphical User Interface (GUI) Calibration of a diffraction setup using the Command Line Interface (CLI)

Calibration of a diffraction setup using pupyter notebooks

Conclusion

Azimuthal integration using the graphical user interface

Performing the azimutal integration from shell scripts

Integration with Python

Tutorials

Application manuals

Design of the library
Python programming API
Installation

Ecosystem

Project

Change-log of versions

Publications about pyFA

Bibliography

Glossary

Calibration of a diffraction setup using Jupyter notebooks

This notebook presents a very simple GUI for doing the calibration of diffraction setup within the Jupyter lab or notebook environment with Matplotlib and Ipywidgets. It has been tested with widget and the notebook (aka nbagg) integration of matplotlib.

Despite this is in the cookbook section, this **tutorial** requires advanced Python programming knowledge and some good understanding of PyFAT.

This tutorial is also available as a video:



The basic idea is to port directly the original pyFAI-calib tool which was done with matplotlib into the Jupyter notebooks. Most credits go Philipp Hans for the adaptation of the origin PeakPicker class to Jupyter.

The PeakPicker widget has been refactored and the Calibration tool adapted for the notebook usage. Several external tools were used with the following version:

• Contributed to ESRF pyFAI library

Ring finding + calibration + accelerated integration

v: master -



"Application of Generative Models for Event Selection and Commissioning of the Long-Lived Particle **Reconstruction Algorithm in the LHCb Trigger Systems"**



ESCOSESAME

Sabin Hashmi



OUTLINE

- 1. Small Recap.
- 2. LHCb Particle Tracking

System

- 3. Real-Time Trigger Systems
- 4. Algorithm Designs and DataProcessing
- 5. Neural Network Designs.



2



- > Developing Machine Learning Based Trigger System for the efficient track reconstruction of Downstream Tracks produced from Long-Lived Particle Decays.

ON A NUTSHELL

- ► LHCb Experiment and it's objectives.?
 - ► LHCb Experiment is a General Purpose Forward Detector, that primarily designed to study beauty quarks from Standard Model of Particles.
 - > LHCb focuses on the research in resolving the mystery of Matter-Antimatter Distribution in the Universe
- Design of LHCb Experiment.
 - ► LHCb consists of 3 Major Detectors, Vertex Locator (VELO), Upstream Tracker (UT), Scintillating Fibre Tracker (SciFi Tracker).
- ► Upgrades and Recent Developments of LHCb Experiment.
 - High Precision Read out systems.
 - ► Replacement of Readout Electronics to Overcome Hardware Trigger (L0) event rate of 1MHz to 30MHz with new software trigger systems.
 - ► And, many more.
- > Applications of the research within the experiment.





REAL-TIME TRIGGER SYSTEM

- Proton-Proton Crossing Rate is once in every 25[ns] (40MHz)
- ► Older Read-out had a bottle neck of 1MHz due to the hardware limitations.
- Upgrade-1 or Run-3 started from mid-2022, with a delay due to multiple factors.
- This is a new approach, to implement a software based trigger system to produce track samples more efficiently and with more purity.
- High Level Trigger (HLT) is split into two different types based on it's purpose.
 - HLT-1 : This produce a Partial Reconstruction of Tracks, from Upgrade-1, this is done using Graphical Processing Units (GPUs) under the "Allen" project
 - ► HLT-2 : This produce the Full-event reconstruction.
- Picking Downstream Tracks are like finding a needle in a Hay-stack due to it's nature.
- "No Storage of RAW data"







ALGORITHM DESIGNS

- > There are three projects designed to be completed this year.
- ► The first algorithm is designed to filter the track segments (Seeds) that gets extrapolated to other detectors to reconstruct the complete tracks.
 - Objective : Filter the Real Seed Tracks and the Ghost Tracks. Small improvements can significantly reduce the complexity of track reconstruction.
- > The Second Algorithm is currently developed using Artificial Neural Network with PyTorch, a Deep Learning Framework.
 - Objective : Cherry Pick the actual Downstream Tracks. that will help to have quality track samples.
- Third Algorithm is in it's very initial stage of development.
 - > Objective : This is a new approach that helps for the detector calibration of using Pre-Trained Machine Learning Models and Automation Pipelines.





5

Thank You.!

