

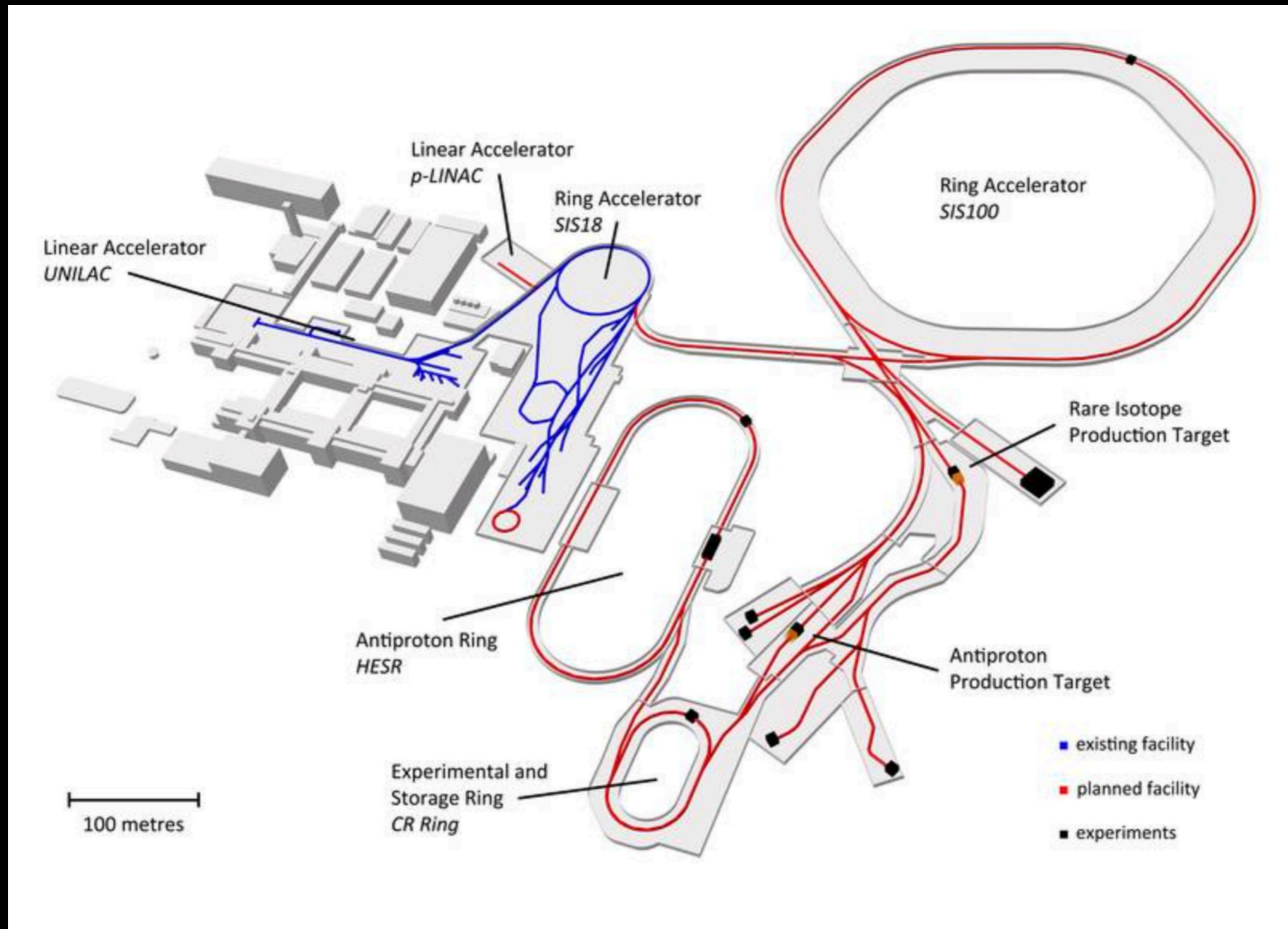


High-Precision Tracking Algorithm for Mass Reconstruction of Heavy Fragments in R3B Experiment at FAIR

D. Kresan, M. Al-Turany, M. Heil
GSI, Darmstadt, Germany

23rd International Conference on Computing in High Energy and Nuclear Physics
9 - 13 July 2018
National Palace of Culture
Sofia, Bulgaria

Facility for Antiproton and Ion Research in Darmstadt, Germany



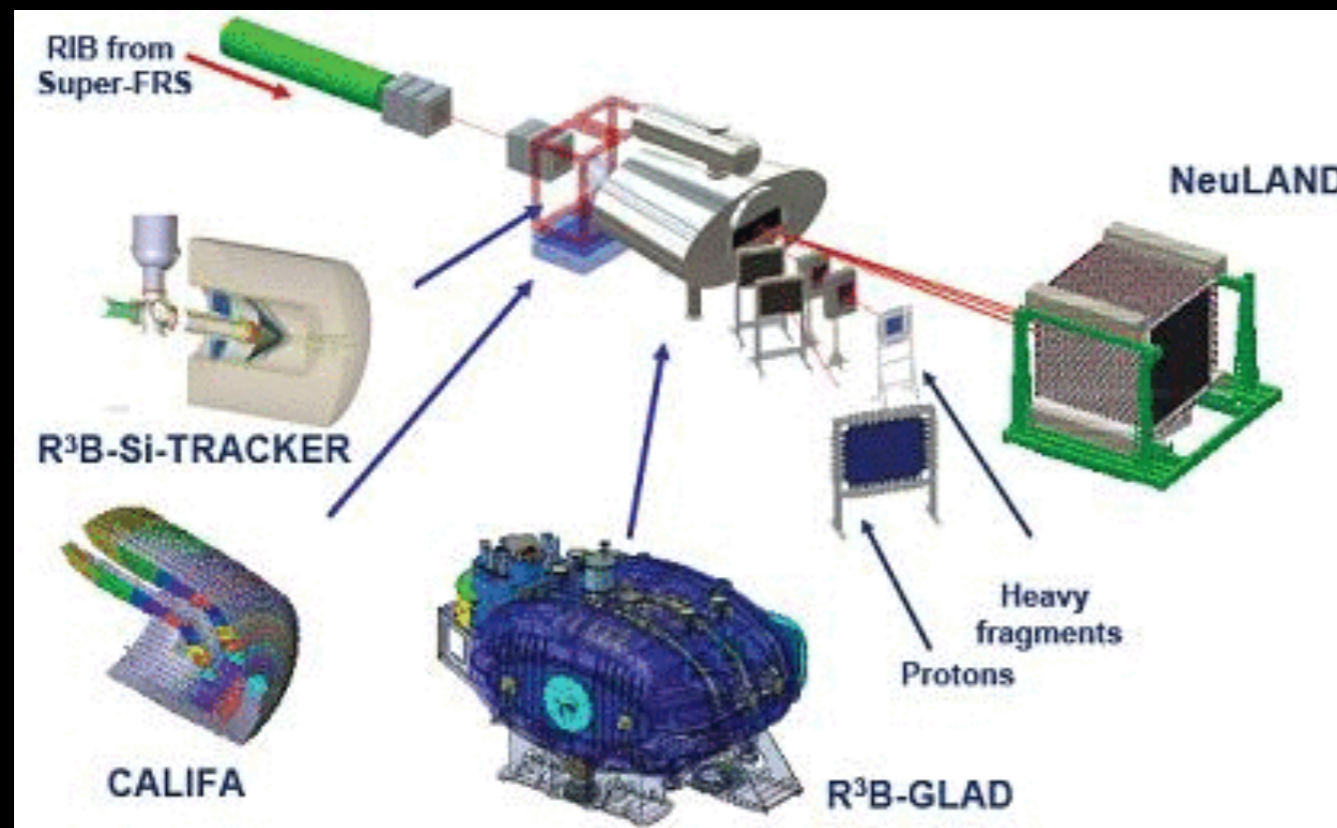
Construction site

April 2018



Reactions with Relativistic Radioactive Beams - R³B

<http://www.gsi.de/r3b>



- Properties of exotic nuclei far off stability
- Study nuclear structure and dynamics
- Astrophysical aspects
- Technical applications

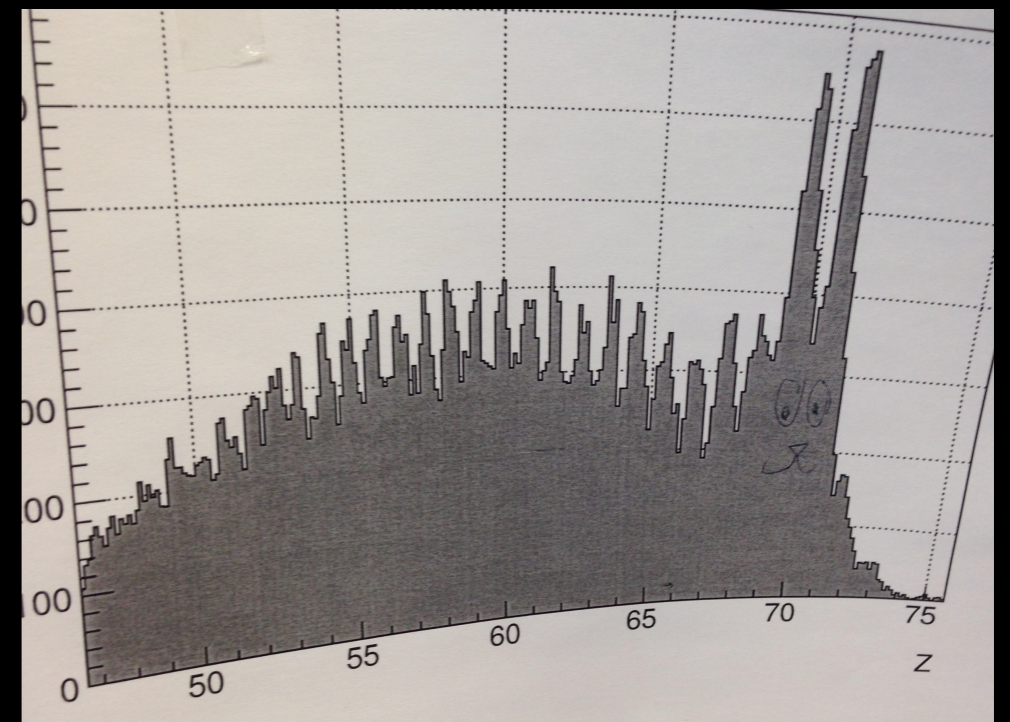
Fragment tracking arm

Measurement of the whole final state is needed to reconstruct the properties of a reaction: neutrons, protons, reaction fragment

Small effects in transversal relative energy on top of Lorentz boost

Momentum resolution in the order of 10^{-3} required

Clean separation of heavy isotopes



Challenges compared to HEP experiments

- Variety of different setups — changes in configuration and detector types
- Due to moderate beam energies and high charge, energy loss in detector layers impacts velocity (and thus trajectory) significantly. Has to be corrected for in the reconstruction
- Highly non-homogeneous magnetic dipole field
- No position measurement inside of field

Tracking algorithm



Depending on the setup:

- I. 2 detectors before the magnet, 1 after — forward fit
- II. 1 detector before the magnet, 2 after — backward fit
- III. 2 before, 2 after — both fits sequentially

No improvement if both fits are applied in cases I. and II.

Components

- Simplified geometry description

No sophisticated stepping navigation
Current approximation model: 1 layer per detector

Components

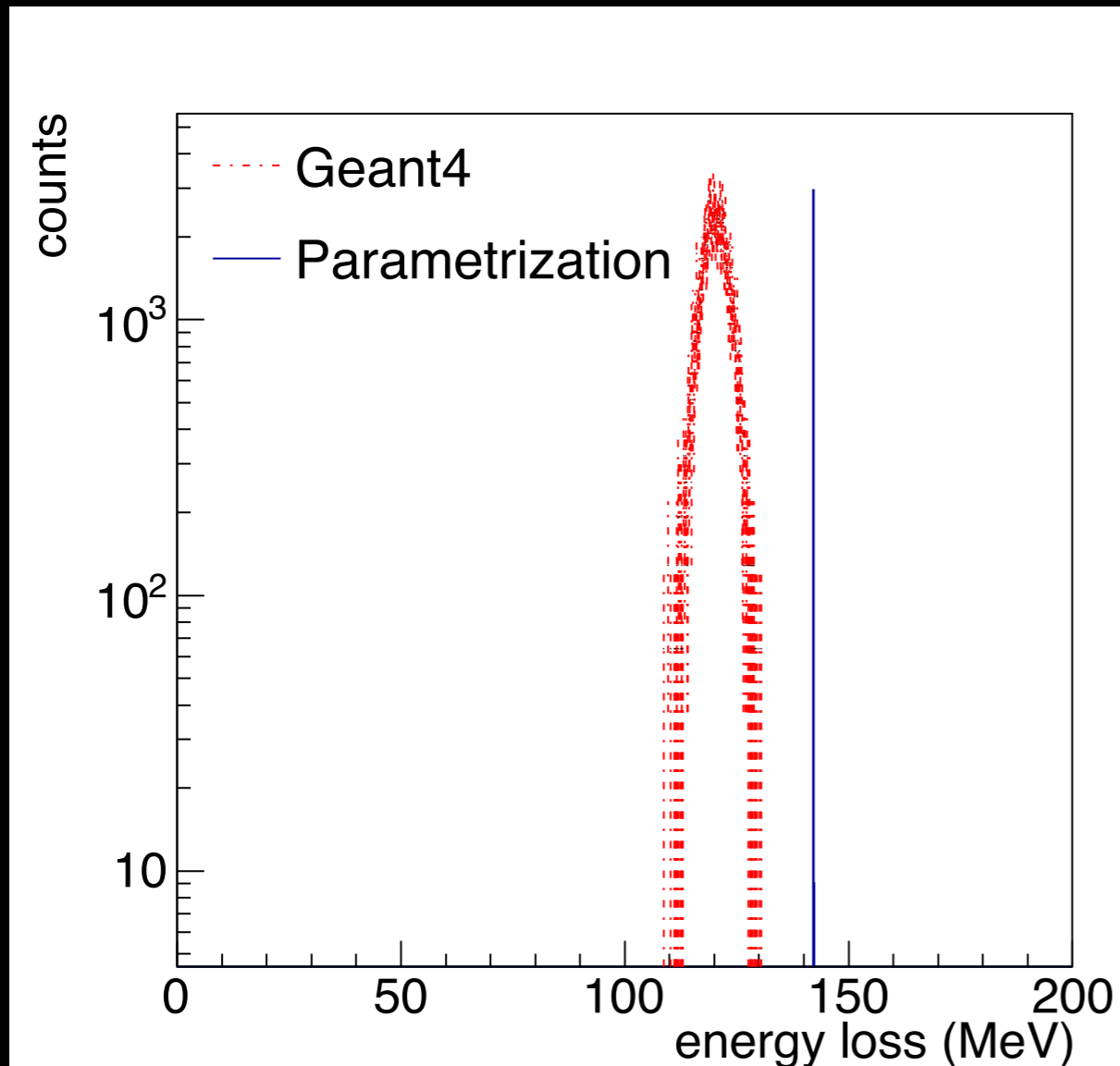
- Simplified geometry description
- Propagation in magnetic field

Highly non-homogeneous dipole field
Step-like propagation based on
Runge Kutta 4th order method
(implemented in the FairRoot framework <https://fairroot.gsi.de>)

Components

- Simplified geometry description
- Propagation in magnetic field
- Energy loss calculation

Energy loss calculation



Bethe Bloch formula with density effect

Weak point: mean excitation energy

Test if can be fitted during alignment with known beam

Components

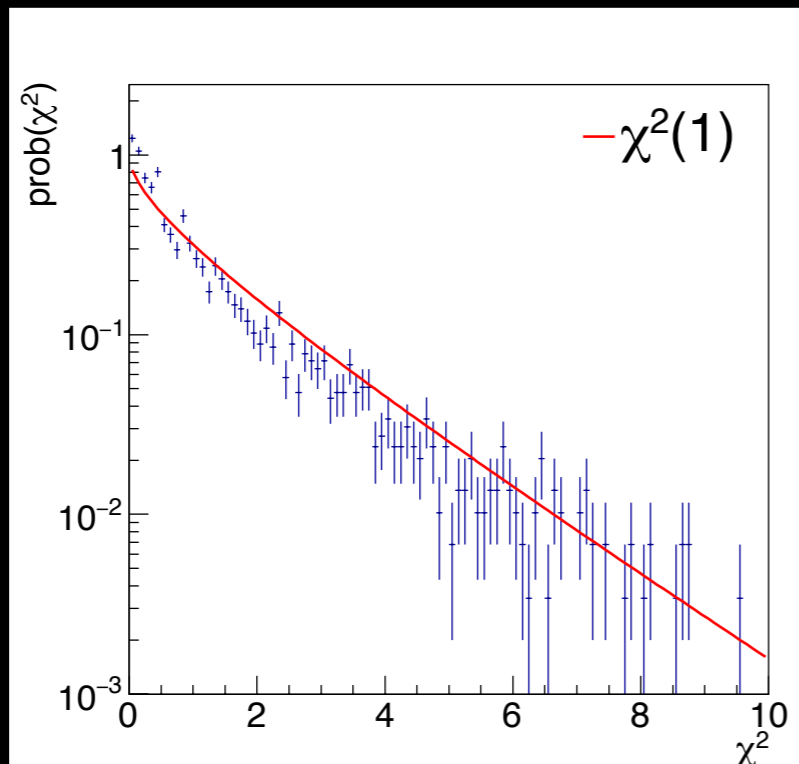
- Simplified geometry description
- Propagation in magnetic field
- Energy loss calculation
- Forward and backward propagation

Components

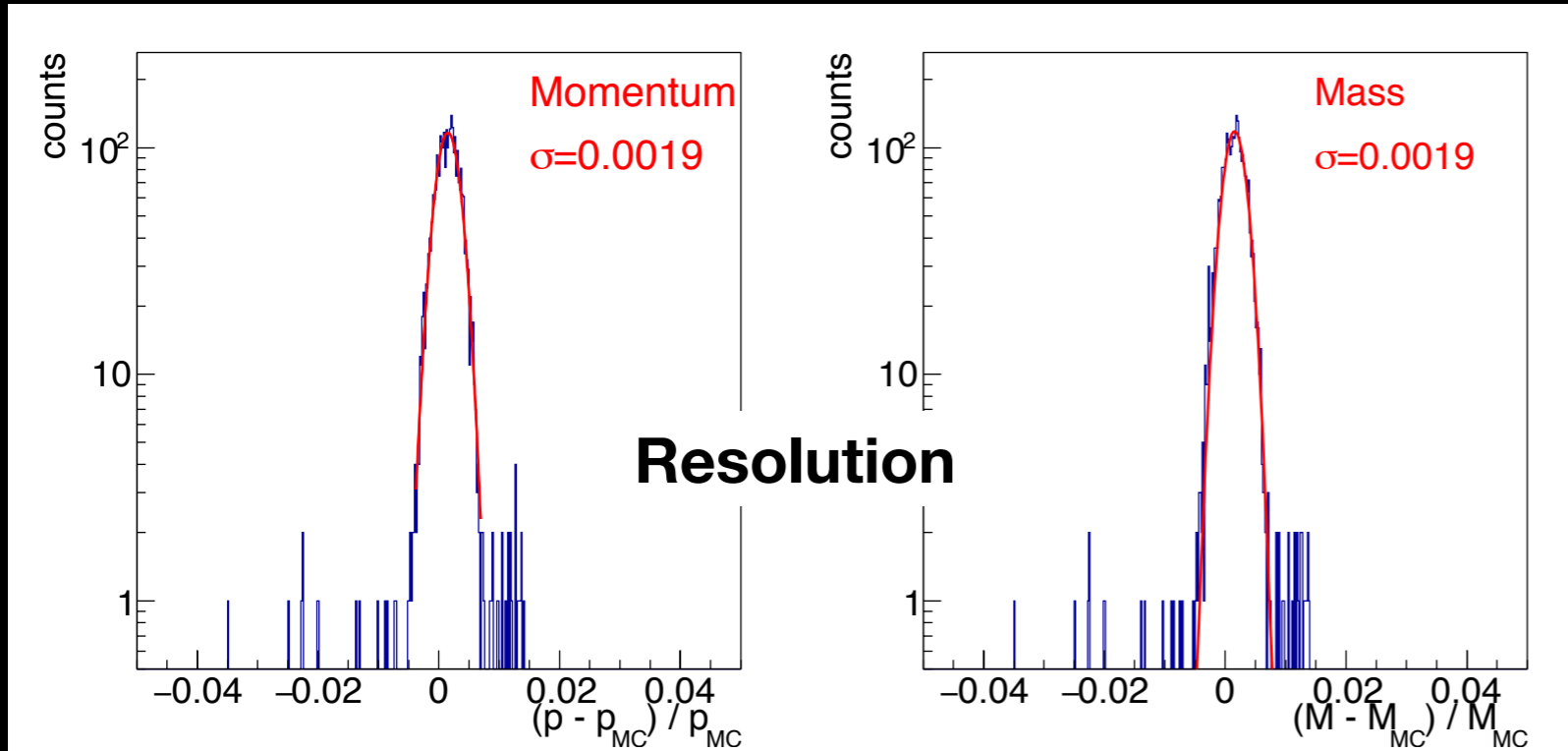
- Simplified geometry description
- Propagation in magnetic field
- Energy loss calculation
- Forward and backward propagation
- Mass fit with Minuit2 from ROOT

Simulation results

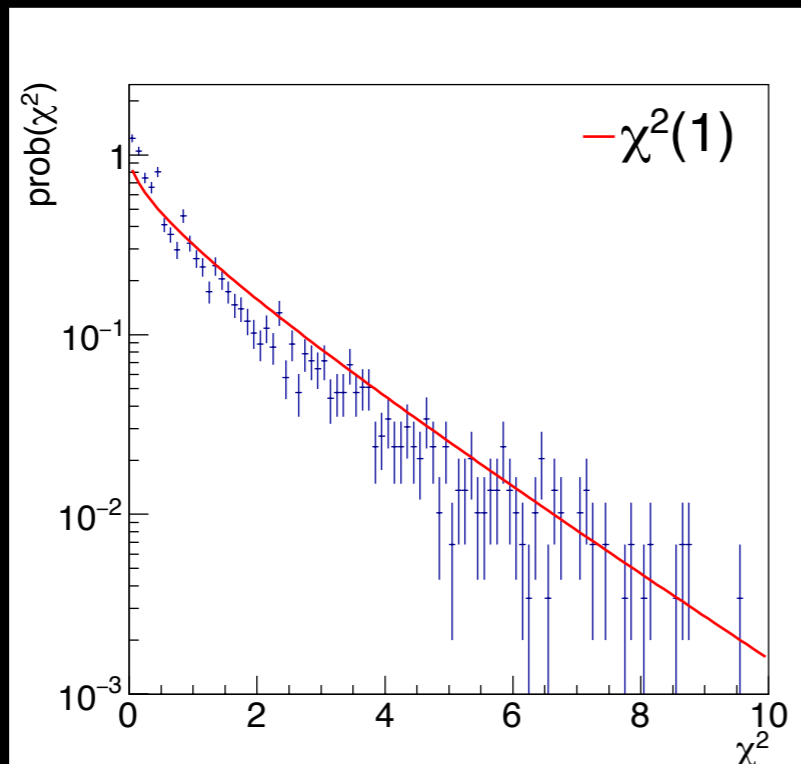
3 Sn isotopes, 1 ion per event
3.000 events
Geant4
 $p_{\text{LAB}} = 1,4 \text{ AGeV}$
backward tracking



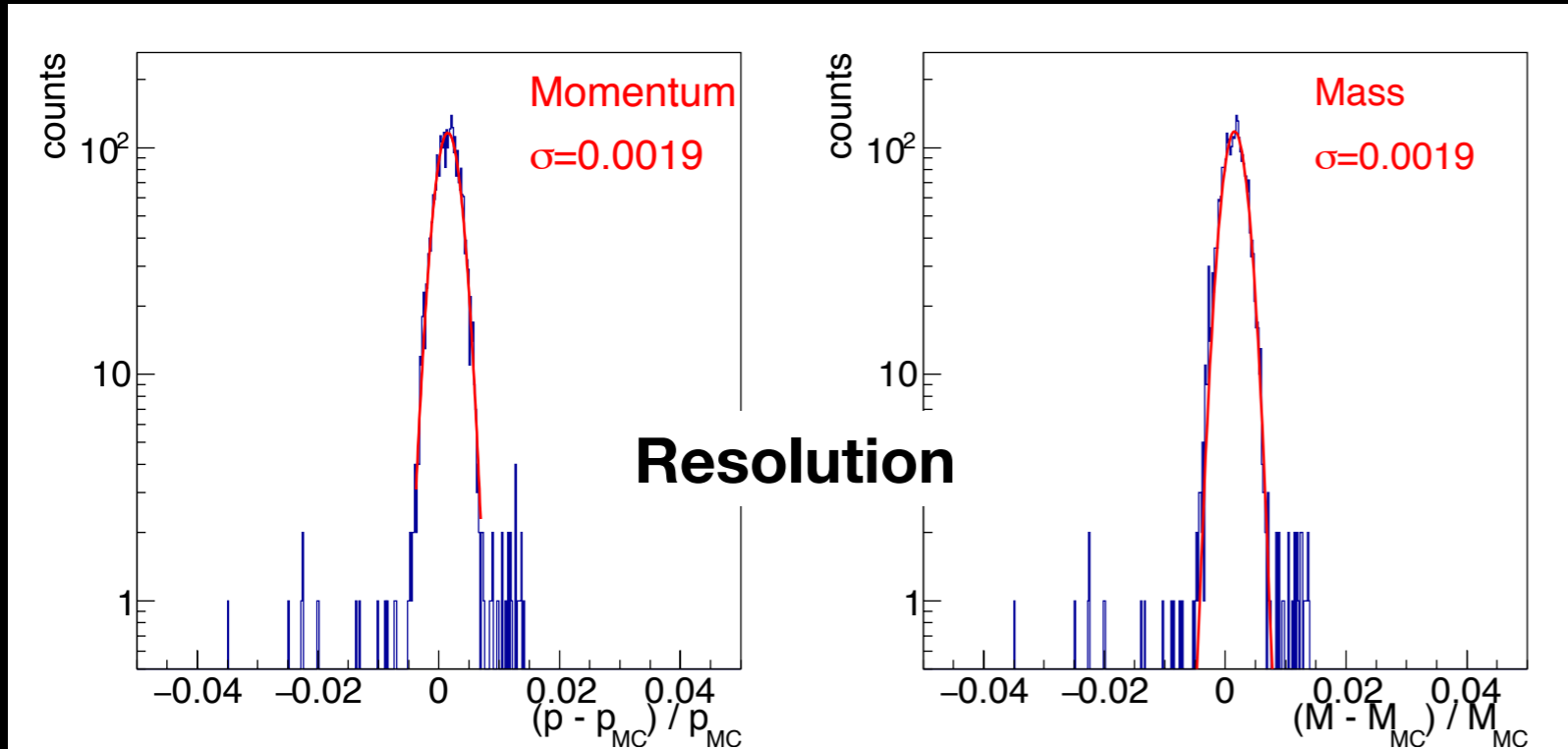
Simulation results



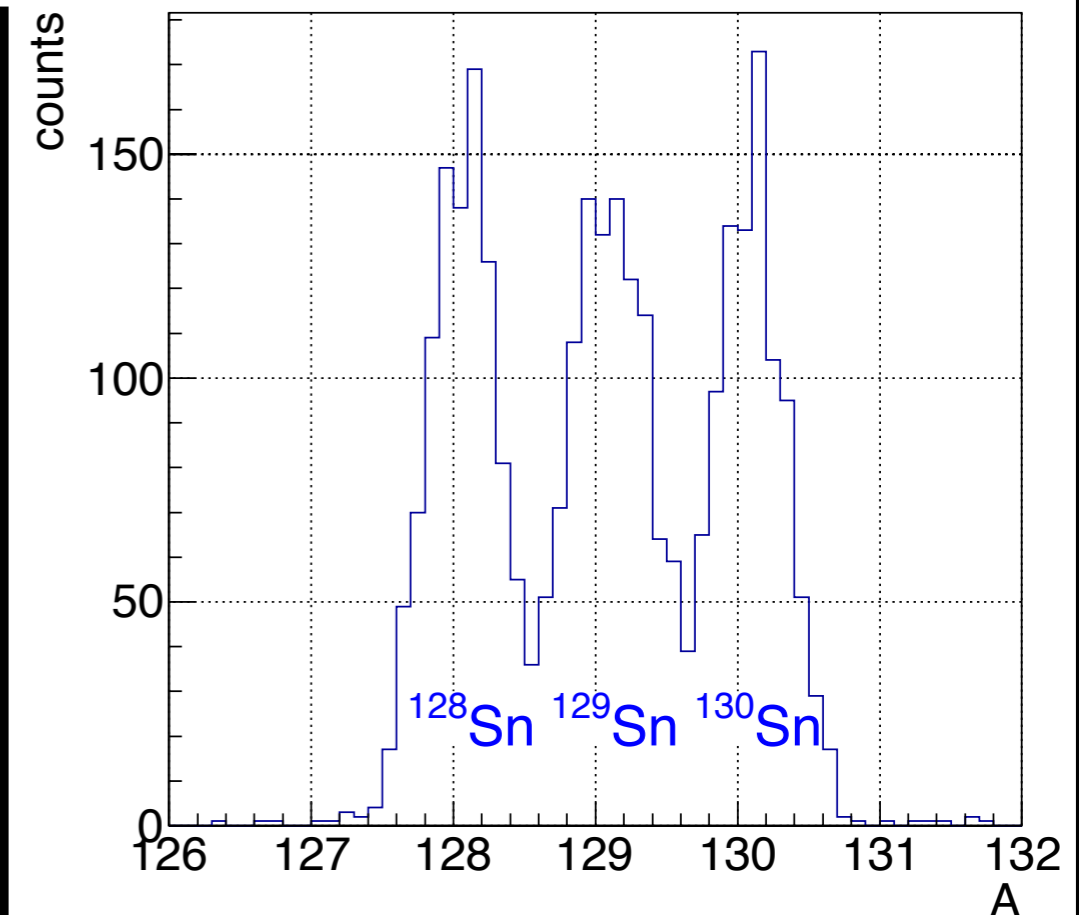
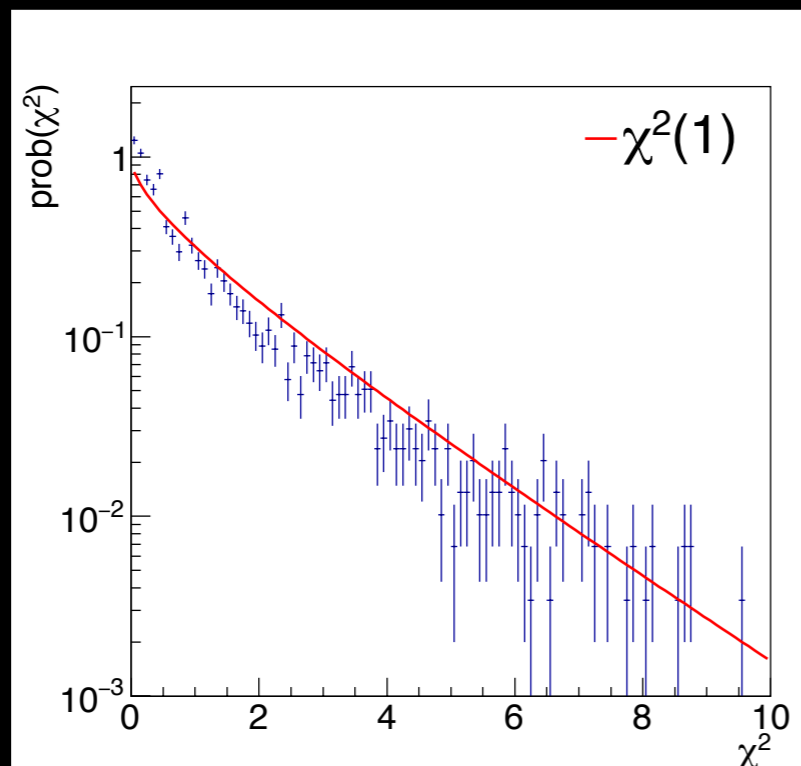
3 Sn isotopes, 1 ion per event
3.000 events
Geant4
 $p_{LAB} = 1,4$ AGeV
backward tracking





Simulation results



3 Sn isotopes, 1 ion per event
3.000 events
Geant4
 $p_{\text{LAB}} = 1,4 \text{ AGeV}$
backward tracking



Performance

-  • Total reconstruction time of 150 ms / event
- Combinatorics penalty: 12.8 candidates / event
- Single fit performance: 11.6 ms / track
-  • 2000 lines of code

Status

- Tracker code is integrated into R3BRoot framework
<https://www.r3broot.gsi.de>
- Single implementation is used to analyze experimental and simulated data
- Ready for data taking this autumn

Summary

- The R3B experiment at FAIR is equipped with the software algorithm for the mass reconstruction using heavy-ion tracking arm
- Required accuracy of 2×10^{-3} in mass reconstruction achieved
- The implementation is modular, compact, straightforward to validate and can be exported
- Further effort needed in timing optimization and in development of automated alignment procedure

Backup

Geometry example - 1 detector plane

```
[fi4GeoPar]
PositionX: Double_t -141.165491
PositionY: Double_t 0.
PositionZ: Double_t 614.
RotationX: Double_t 0.
RotationY: Double_t -16.7
RotationZ: Double_t 0.
DimensionX: Double_t 20.
DimensionY: Double_t 20.
DimensionZ: Double_t 0.02
Z: Double_t 5.57354
A: Double_t 11.0723
Density: Double_t 1.032
IonisationEnergy: Double_t 57.4e-6
```

No recompilation needed when changing geometry