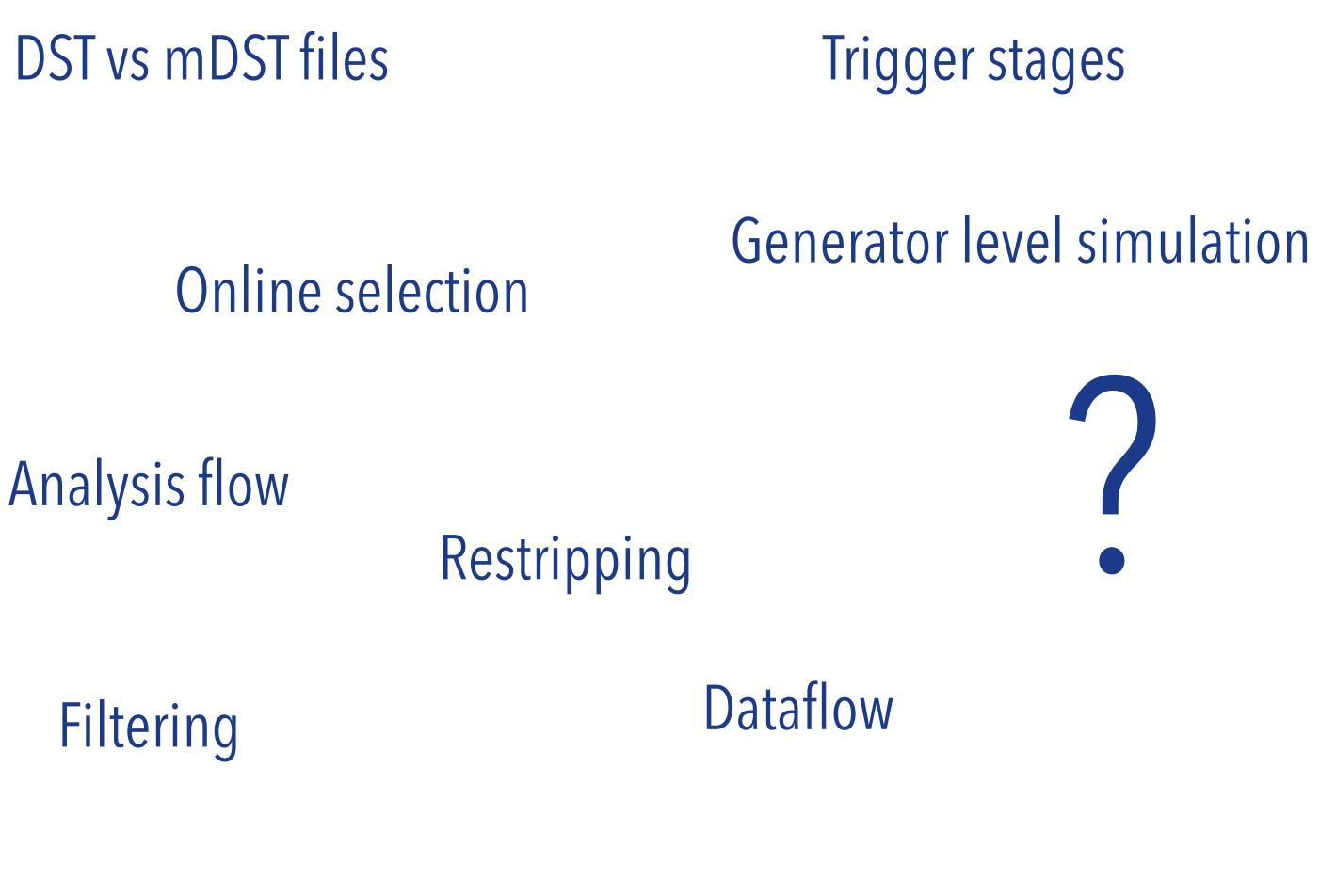


Estd. 2015

FROM COLLISIONS TO PHYSICS - And how to get to publication -

Janina Nicolini

Goal of the lesson



Fullstreams vs Streams

Prescaling

Stripping (campaign)

Offline selection

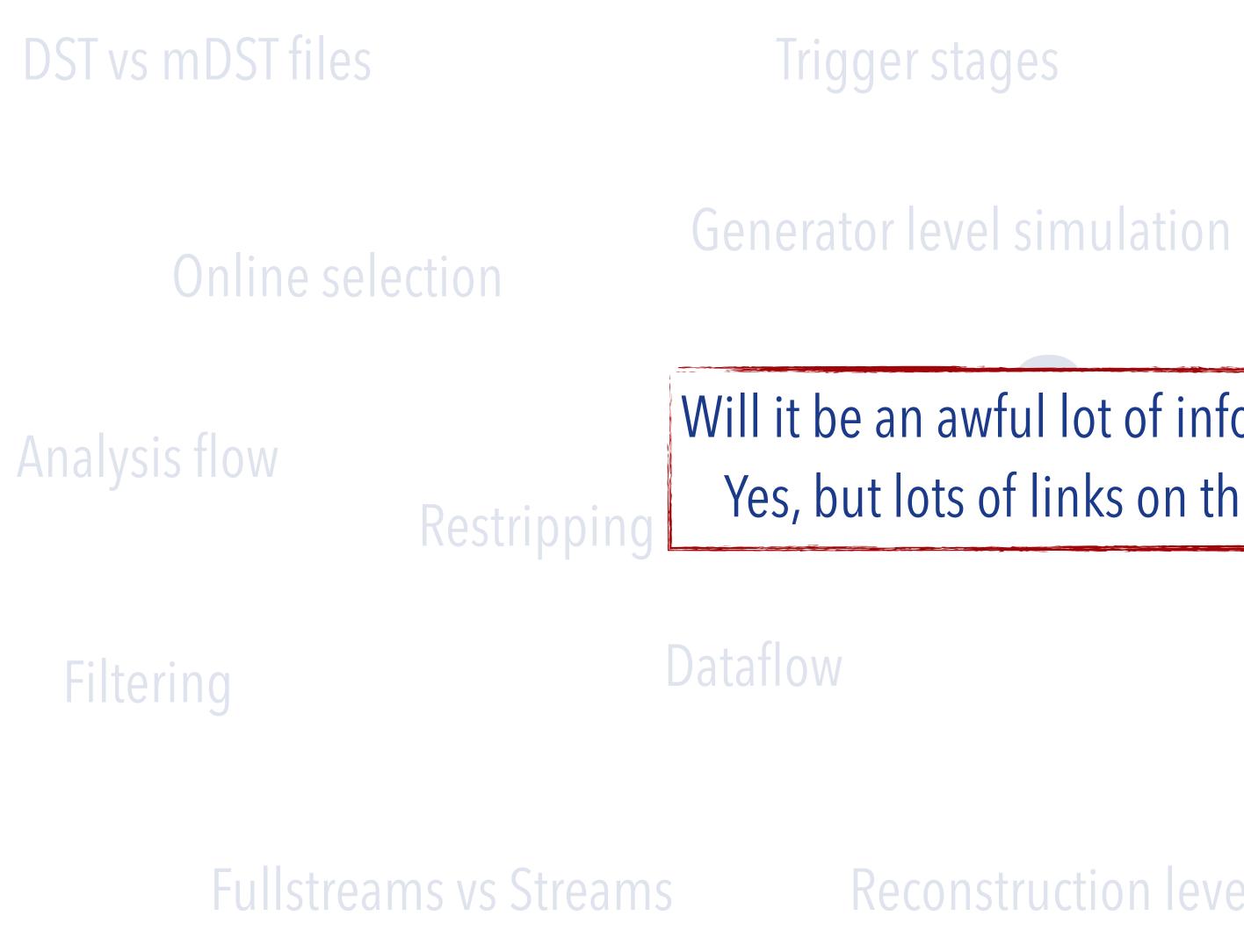
Run 1 vs Run2 dataflow

Reconstruction level simulation

Analysis preservation



Goal of the lesson



Prescaling

Stripping (campaign)

Will it be an awful lot of information? Yes, but lots of links on the slides

Offline selection

Run 1 vs Run2 dataflow

Reconstruction level simulation

Analysis preservation



Software designed to make common analyses as easy as possible. So how and what do we do in analyses?

We usually measure:

- Production
- Decay properties

Of heavy flavour hadrons

 \rightarrow short lifetime \rightarrow what can we do?

We usually measure:

- Production
- Decay properties

Of heavy flavour hadrons

 \rightarrow short lifetime \rightarrow what can we do?

Use "stable" particles

Protons p/\bar{p}

Photons γ

Electrons e^{\pm}

Deuterons

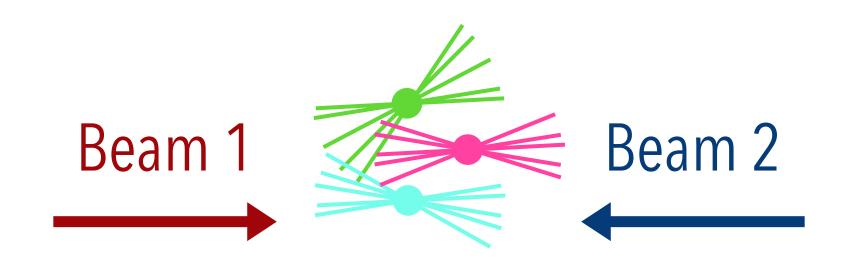
Charged pions π^{\pm}

Charged kaons K^{\pm}

Muons μ^{\pm}

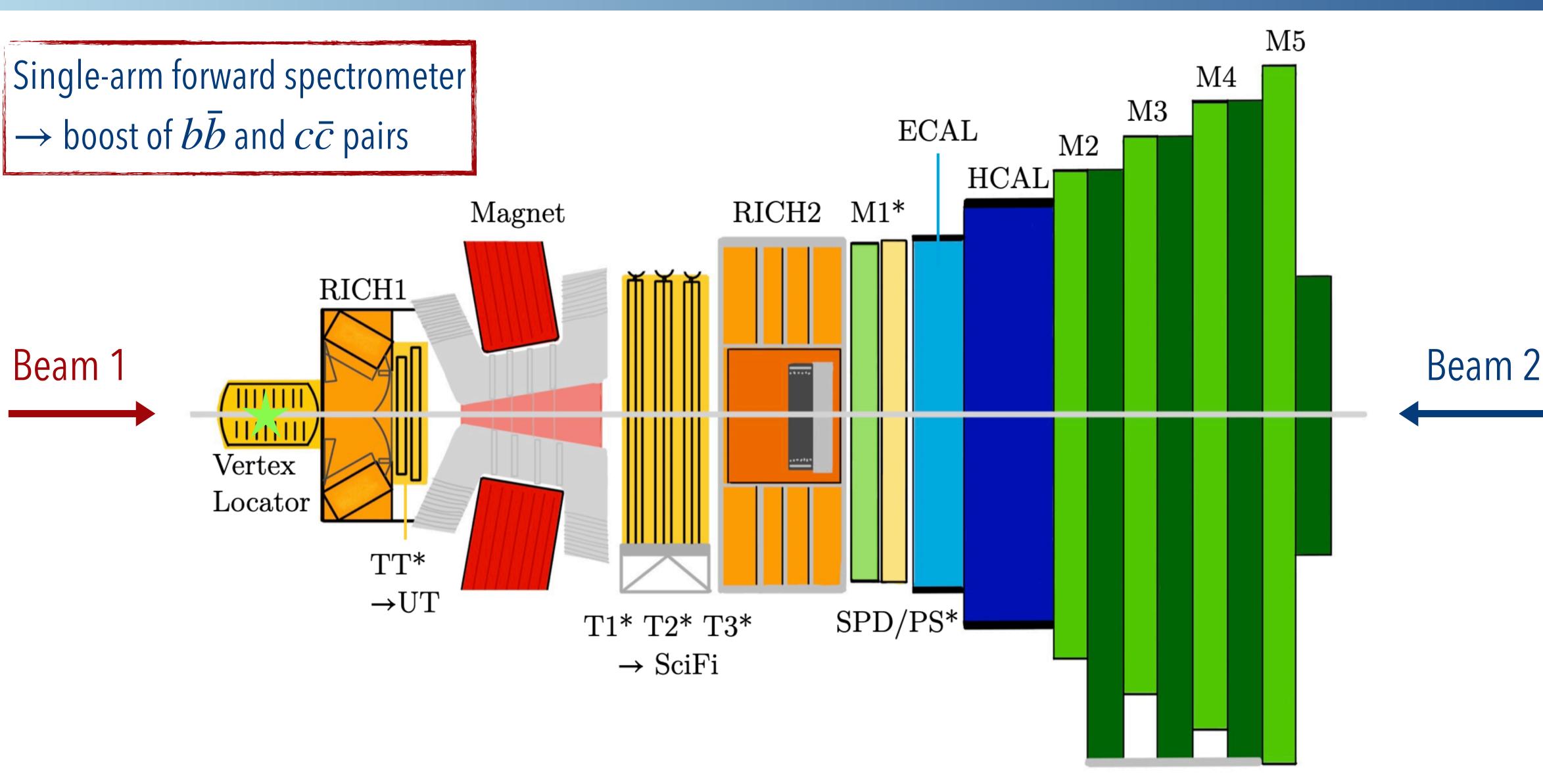


First step reconstruction of properties. Not single particle, but all charged tracks at the same time



(Several) primary vertices \rightarrow where protons collide \rightarrow large number of tracks intersecting

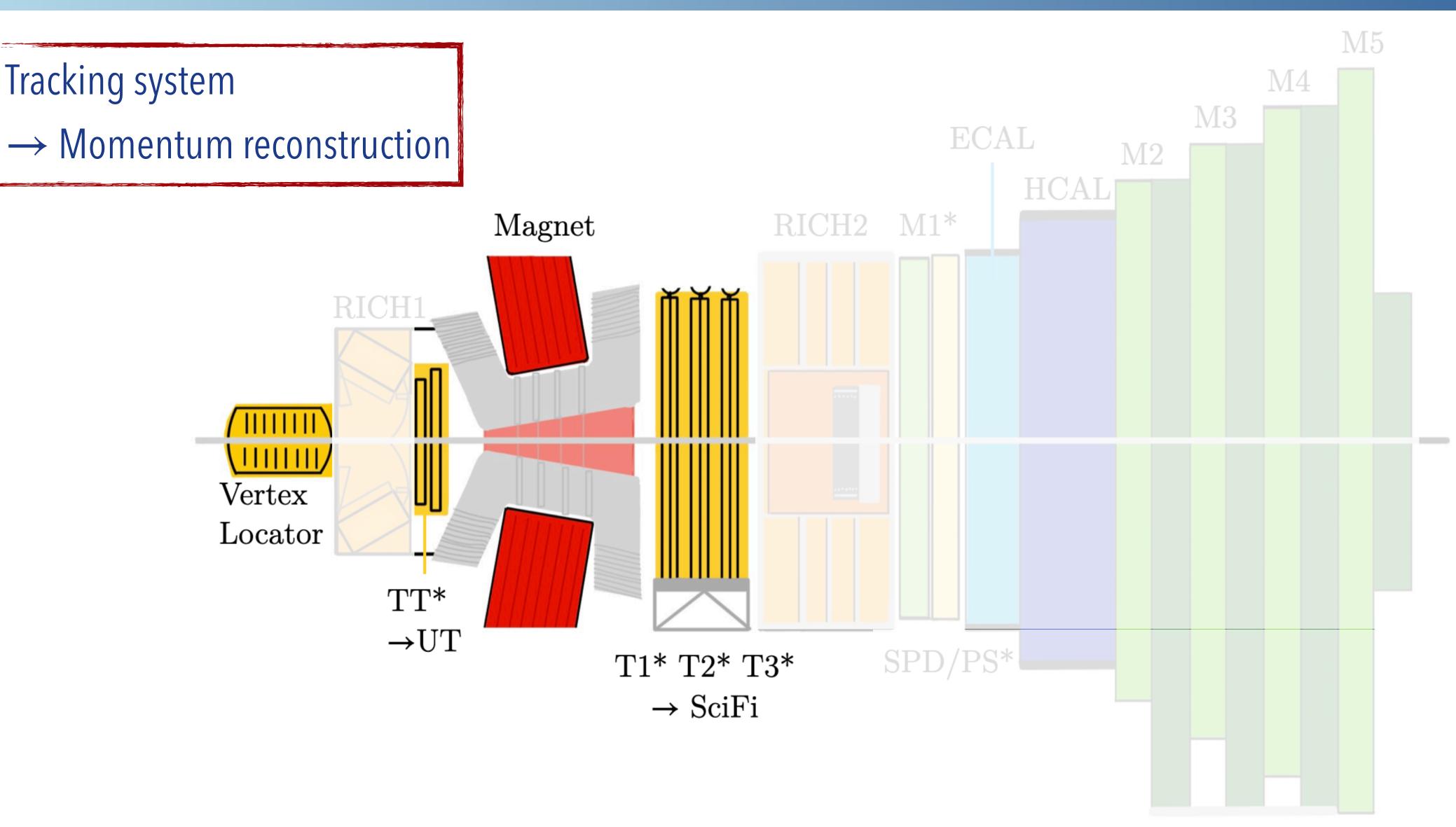
First step reconstruction of properties. Not single particle, but all charged tracks at the same time

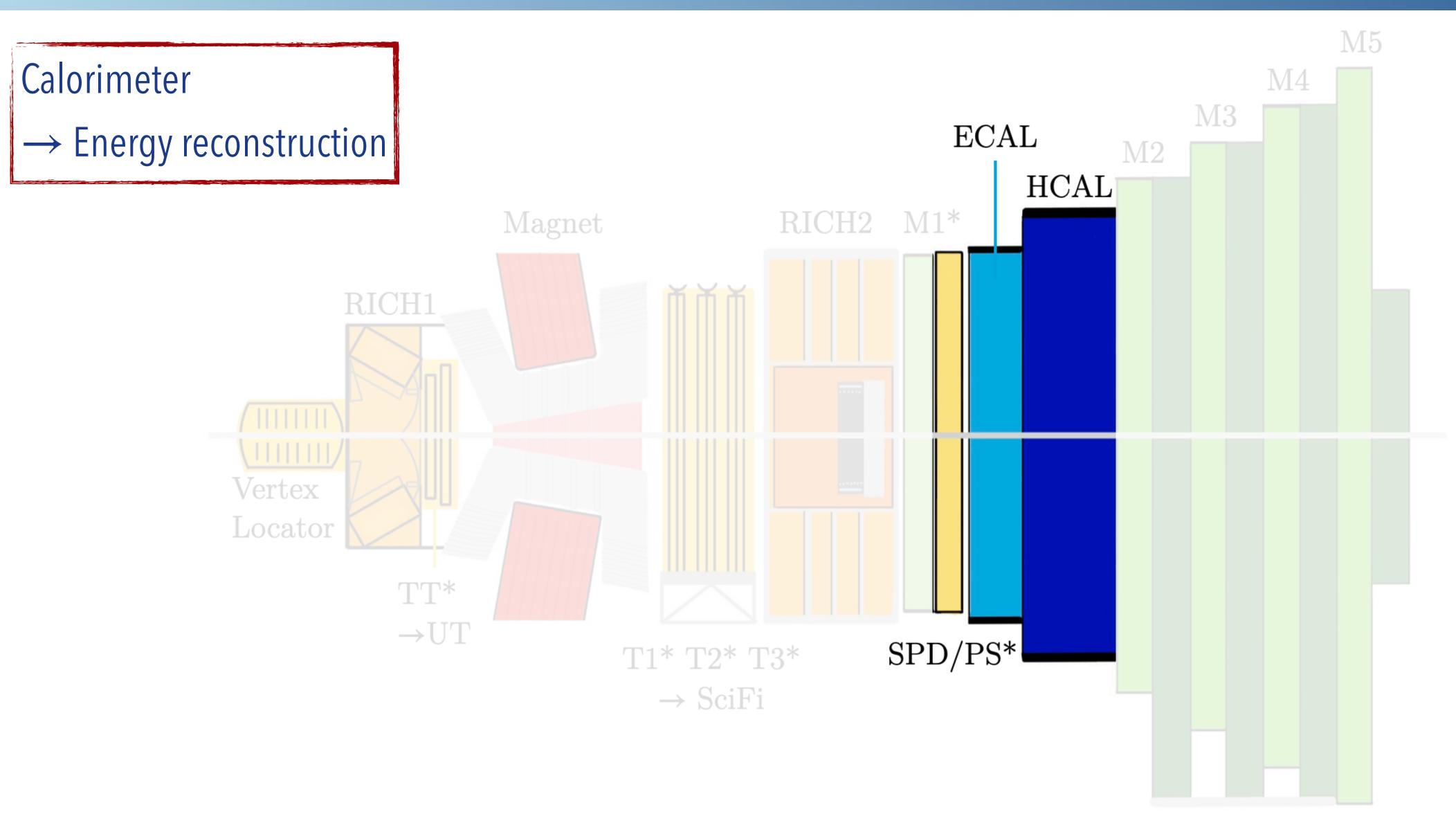


* replaced or removed during Upgrade for Run 3

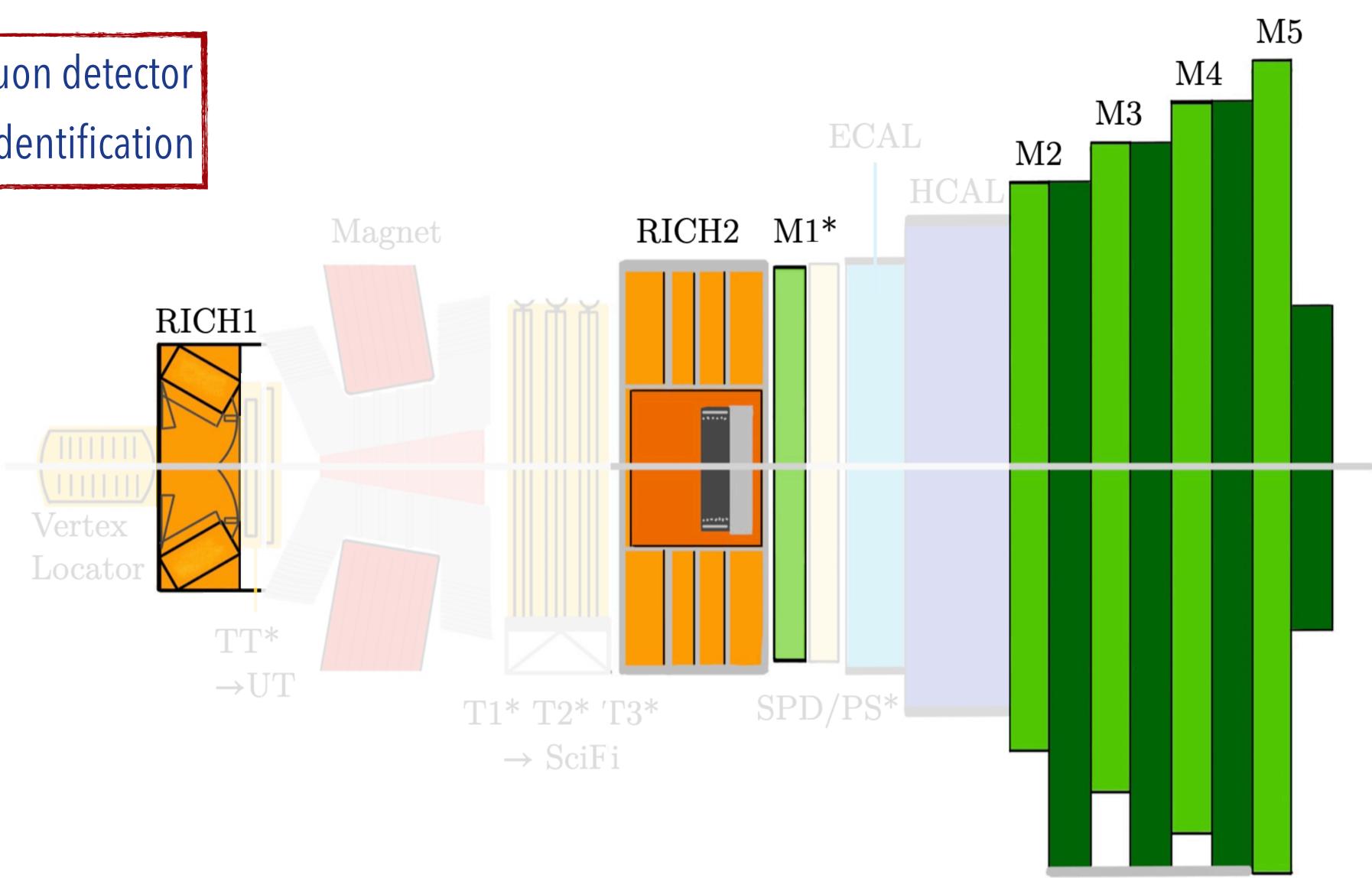








RICH and muon detector \rightarrow particle identification



Building decay candidates

Difficulties we encounter

- Contributions from detector effects
- "Ghost" tracks
 - \rightarrow combination of random hits
- Typical hundreds of tracks
- \rightarrow statistical analysis of events

Building decay candidates

Reconstructing a $J/\psi \rightarrow \mu^+ \mu^-$ decay

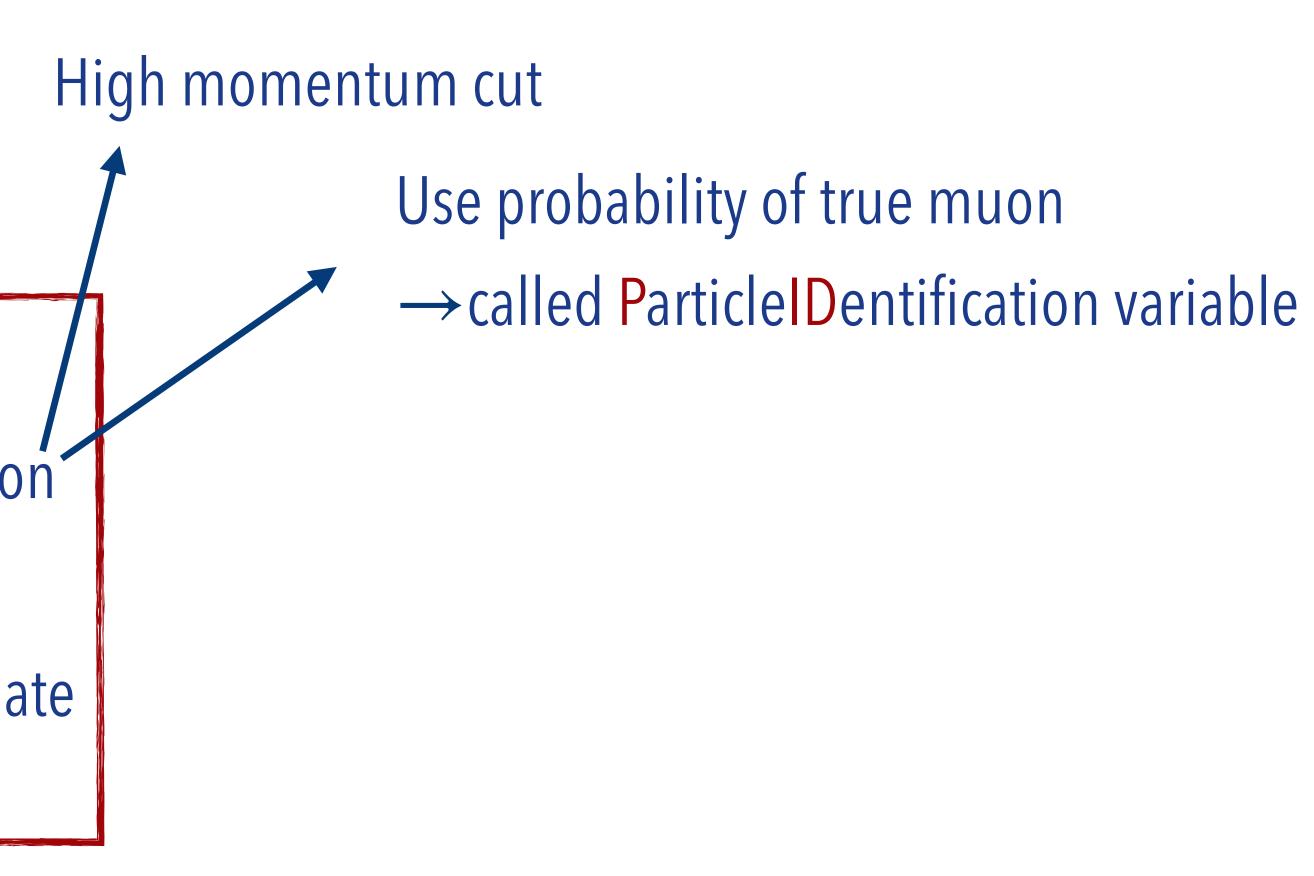
- Select suitable tracks created by reconstruction
- Create pairs of oppositely-charged tracks
- Fit each pair under the hypothesis the originate from a common point in space



Building decay candidates

Reconstructing a $J/\psi \rightarrow \mu^+ \mu^-$ decay

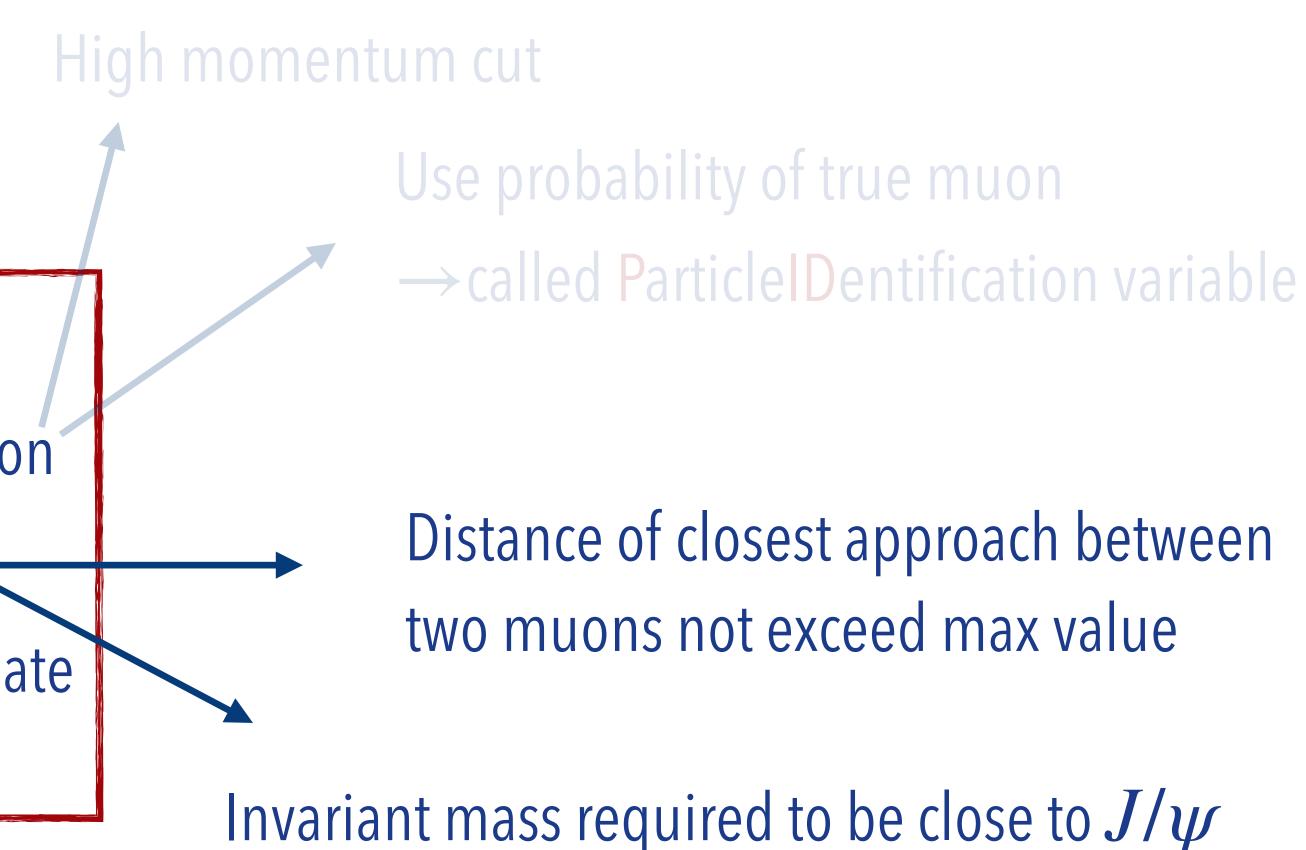
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Building decay candidates

Reconstructing a $J/\psi \rightarrow \mu^+\mu^-$ decay

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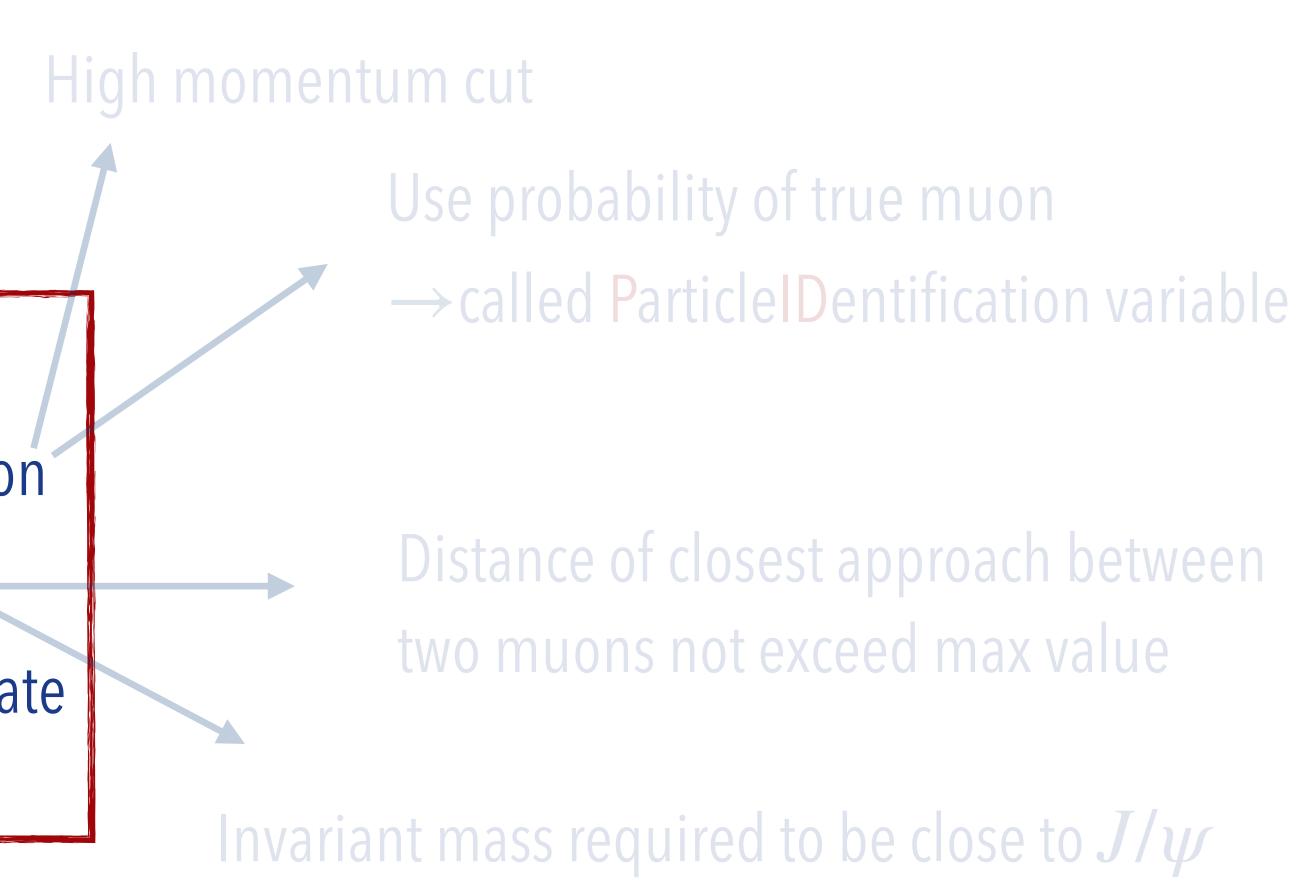


Building decay candidates

Reconstructing a $J/\psi \rightarrow \mu^+\mu^-$ decay

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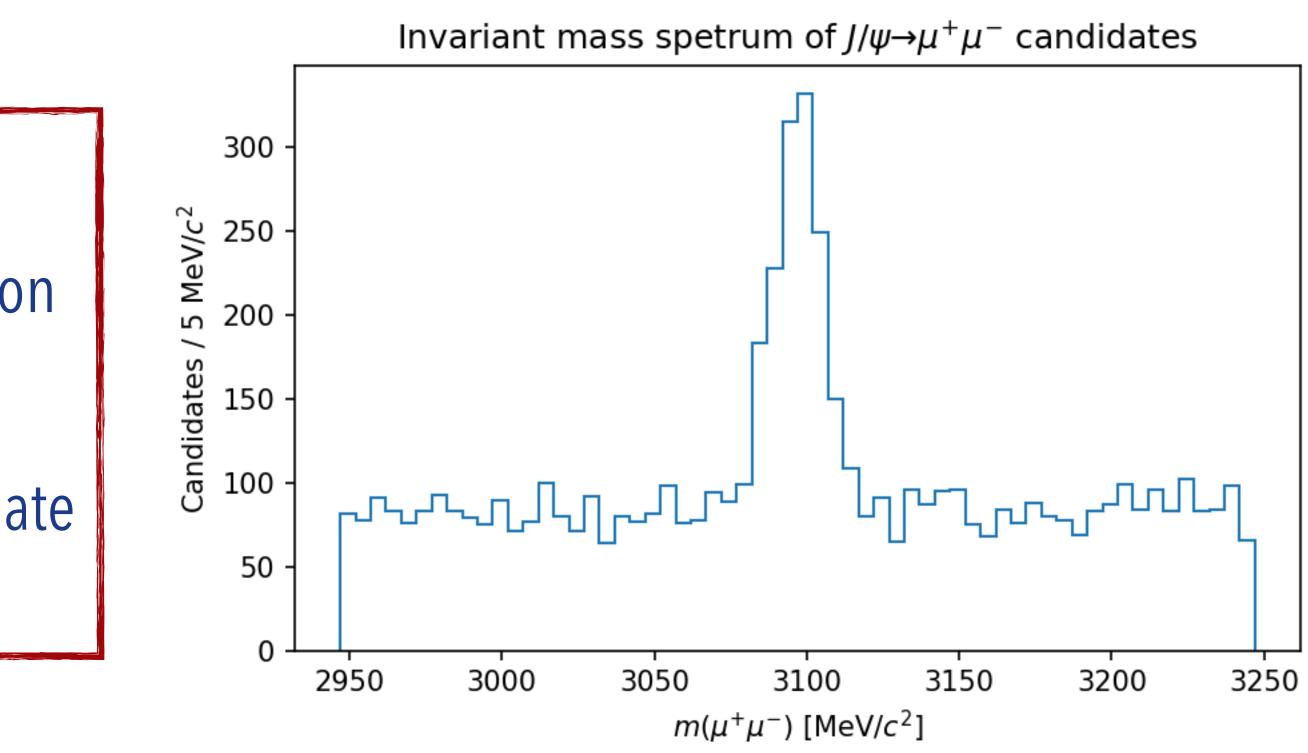
<u>DecayTreeFitter</u> tool allows to perform fit in software $\rightarrow \chi^2$ of fit can be used in selection



Creating J/ψ candidate from muon four-momenta

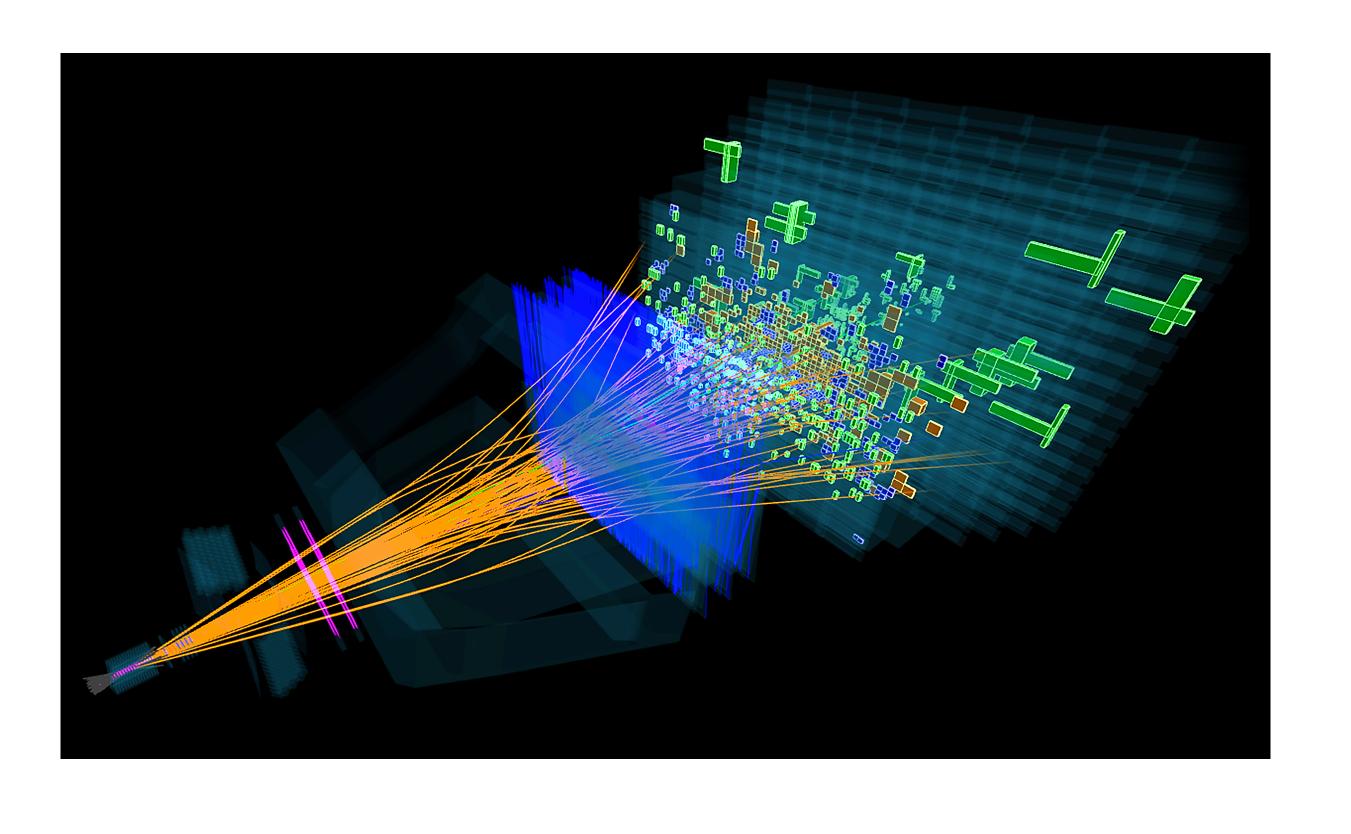
Reconstructing a $J/\psi \rightarrow \mu^+\mu^-$ decay

- Select suitable tracks created by reconstruction
- Create pairs of oppositely-charged tracks
- Fit each pair under the hypothesis the originate from a common point in space



How do we decide what to save?

The LHC can provide a bunch crossing every 25ns.



Data rate is 40MHz, but :

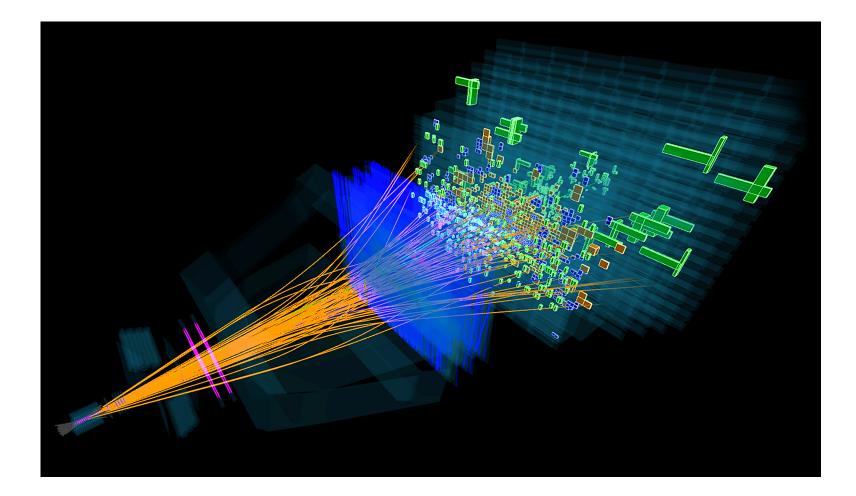
- FPGA readout max. 1MHz
- Run 1: rate to storage max. 5kHz
- Run 2: rate to storage max. 12.5kHz

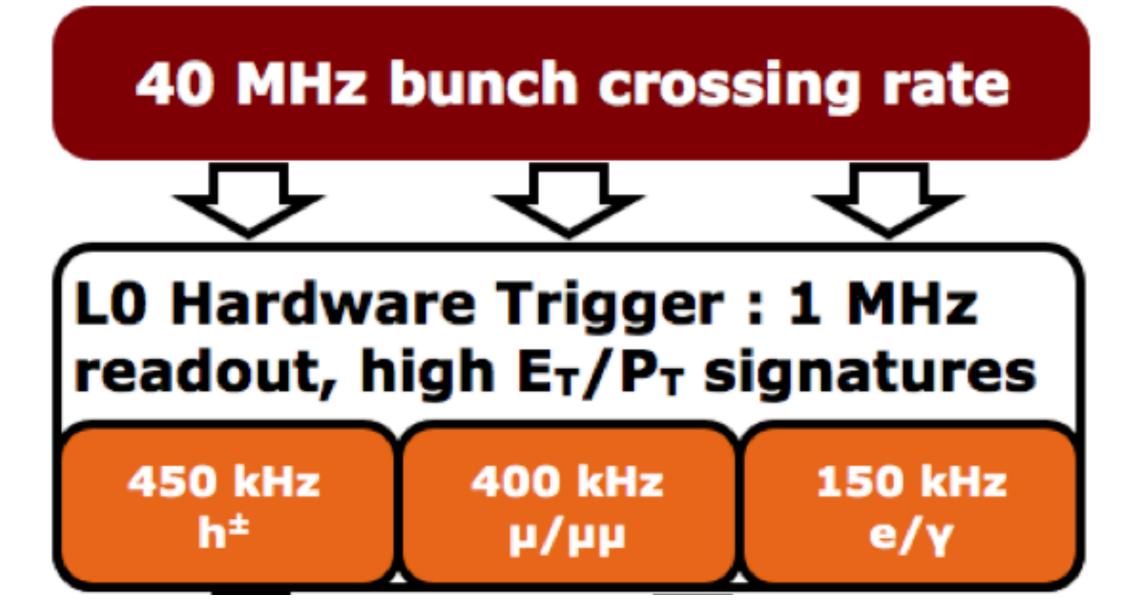
So we cannot save 1TB/s.



How do we decide what to save?

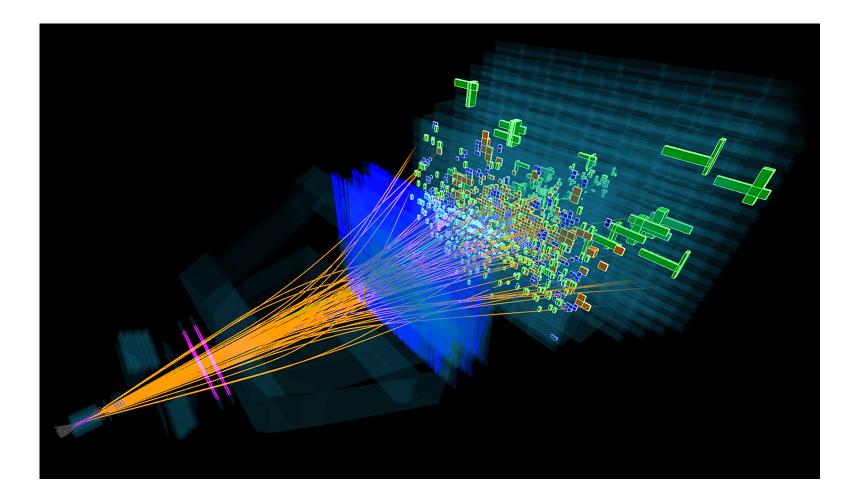
First a hardware trigger stage called LO.

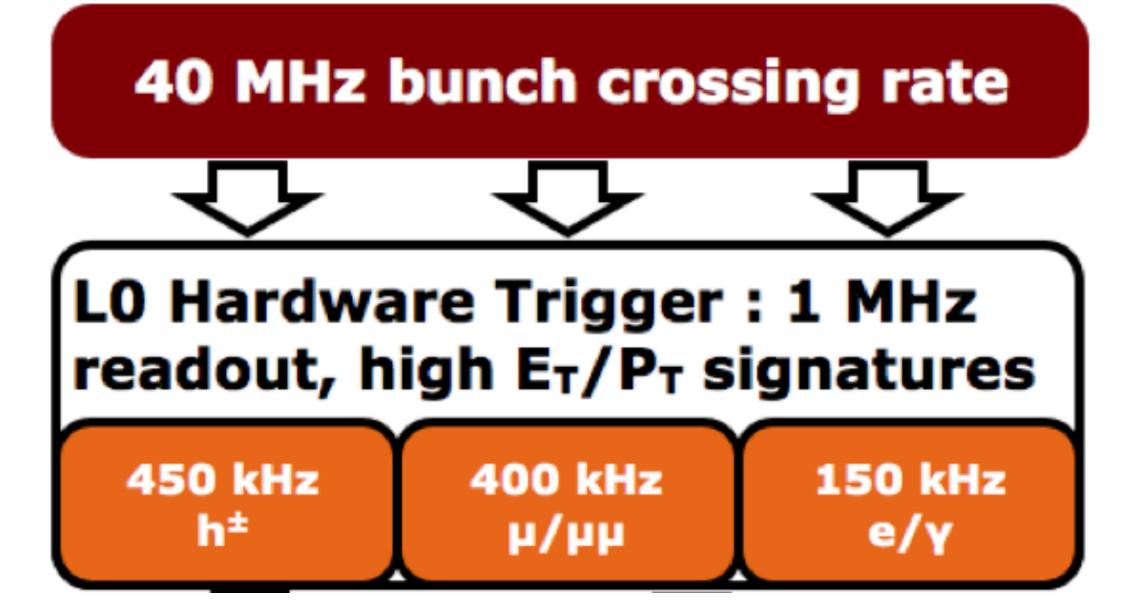


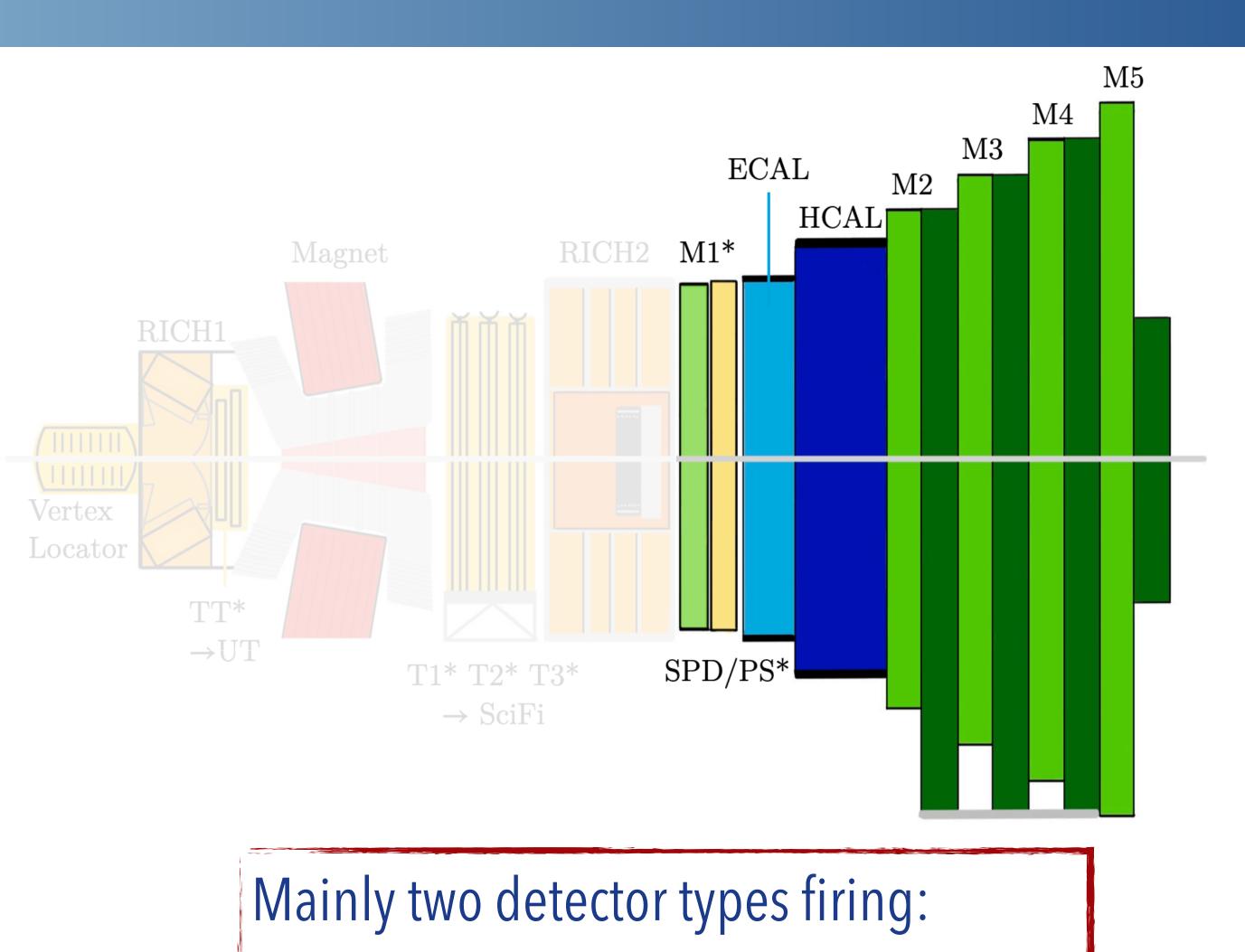


How do we decide what to save?

First a hardware trigger stage called LO.



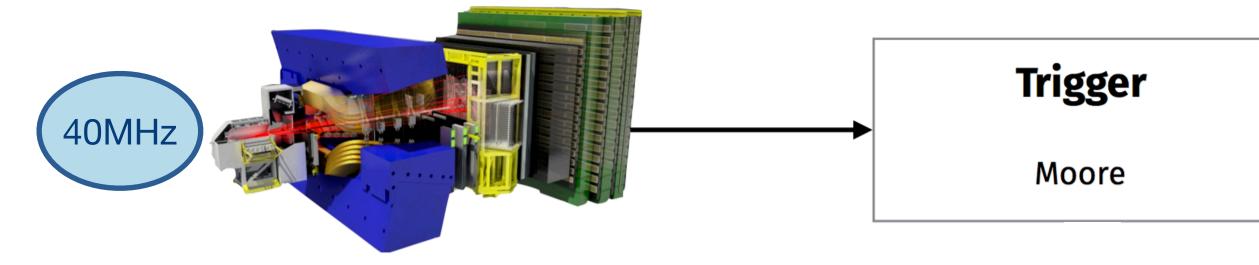




• Hits in the muon stations

• Energy deposit in the ECAL and HCAL

Second step of the online reconstruction.

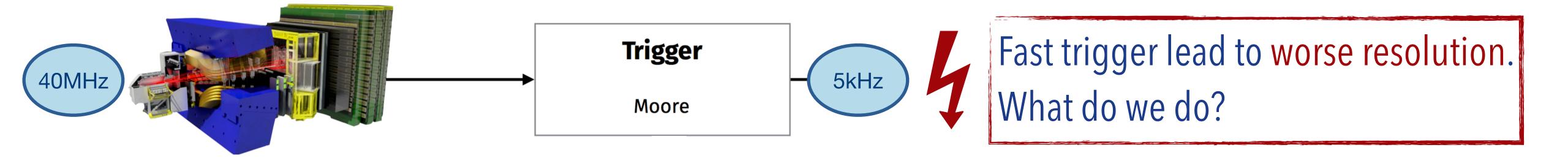


Next software stage called High Level Trigger

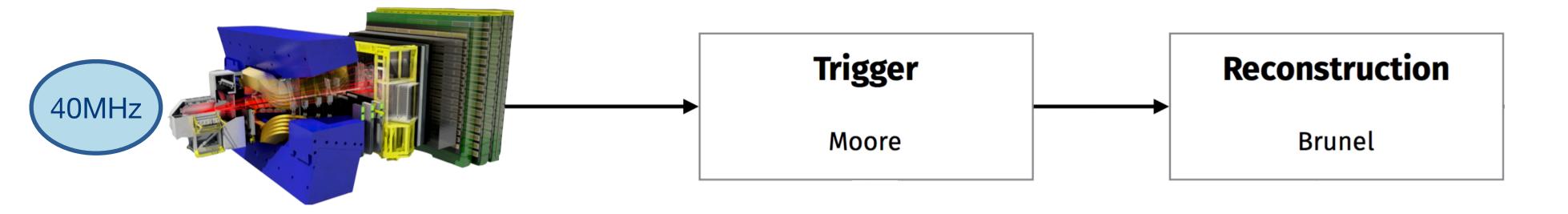
- HLT1: Adding tracking information
- HLT2: Adding RICH information
- Both run in <u>Moore framework</u>



Online reconstruction done, but..



The offline reconstruction





- Clusters,...

Improve reconstruction of:

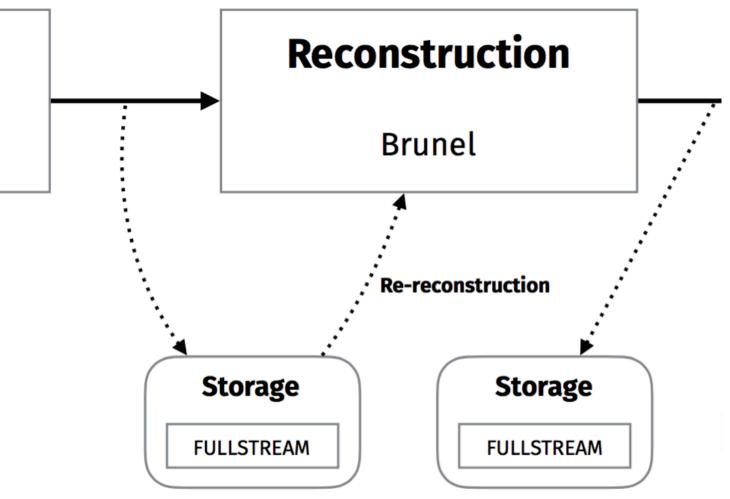
Run in <u>Brunel framework</u>

And storage of the data to tape

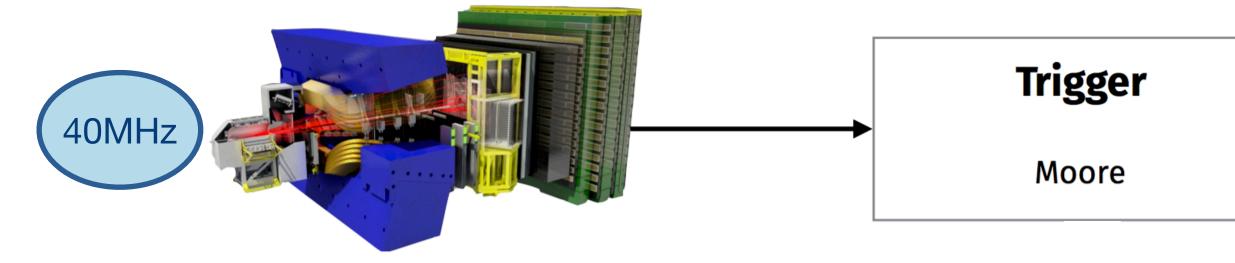


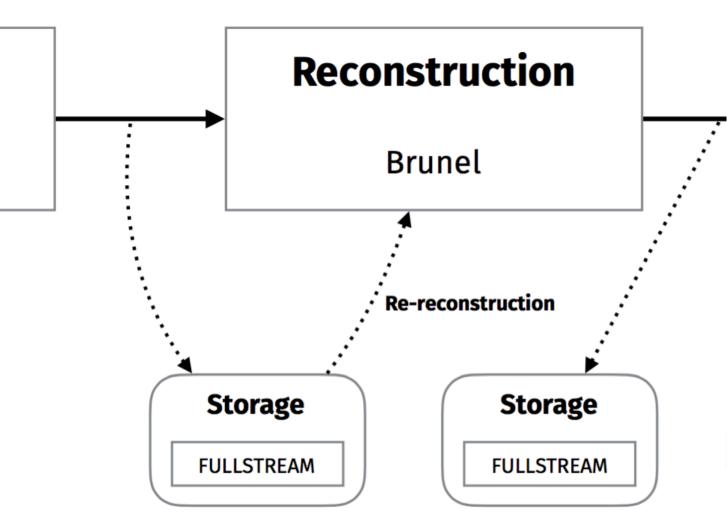
Raw banks of all the subdetectors are saved in FULLSTREAM

- \rightarrow no further selection
- → Information stored in DST Files



And storage of the data to tape

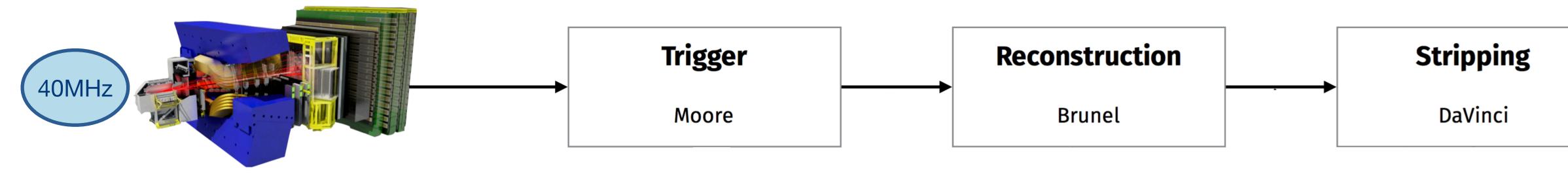




After reconstruction huge DST files. What do we do?



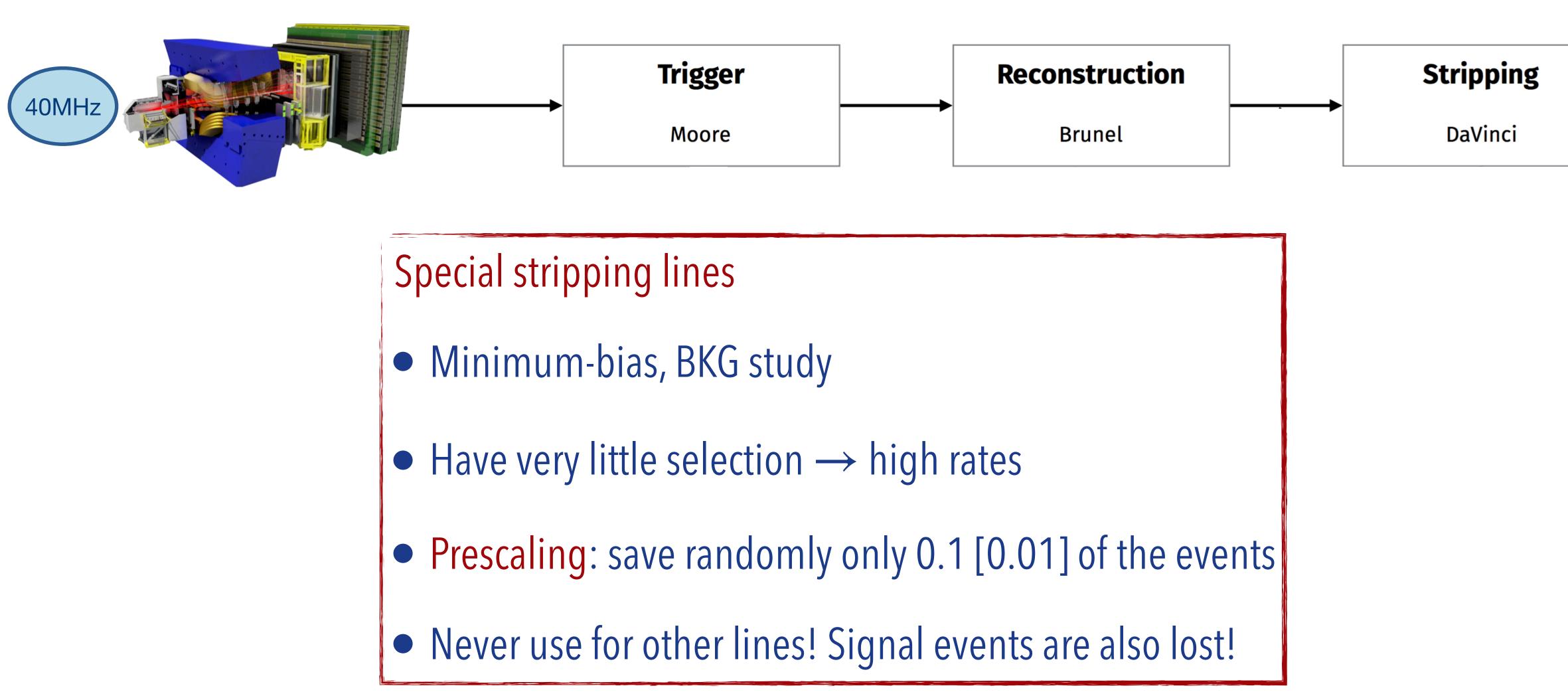
Reducing the file sizes

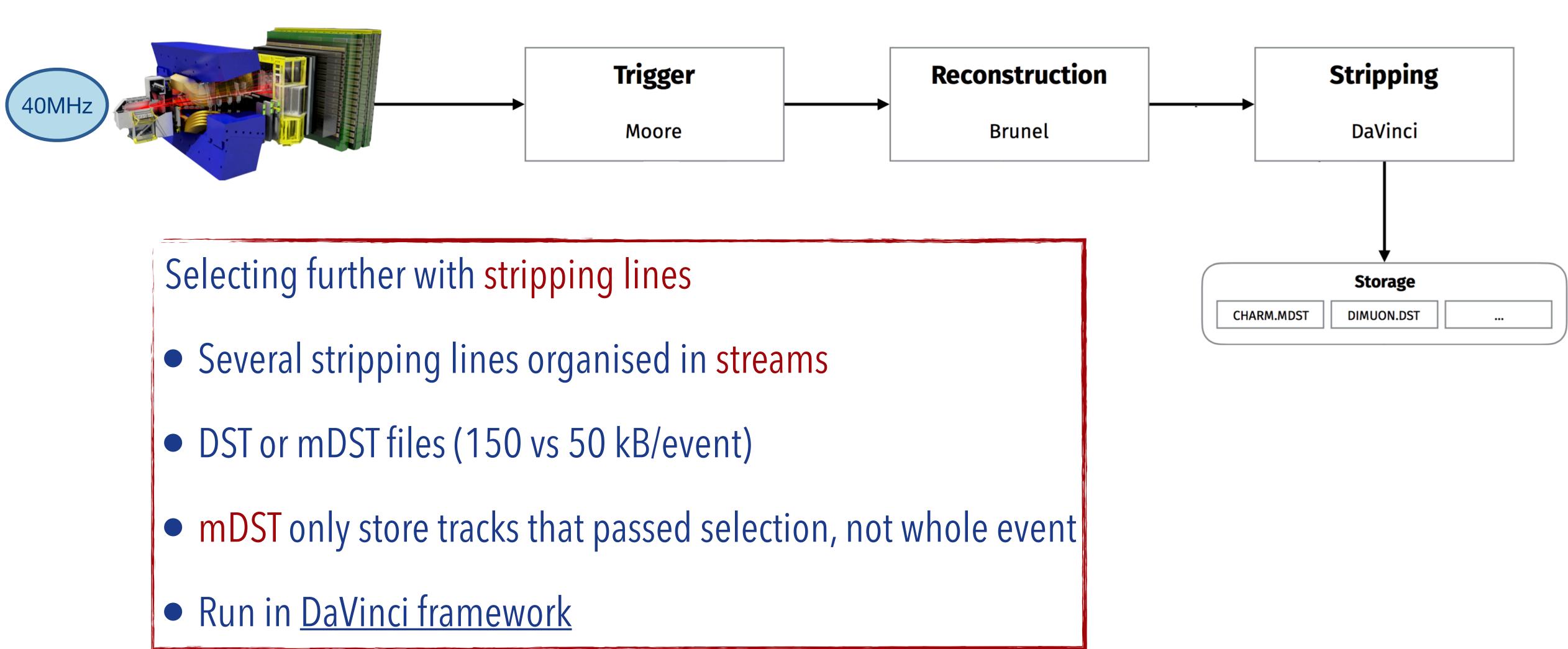


Selecting further with <u>stripping</u> lines

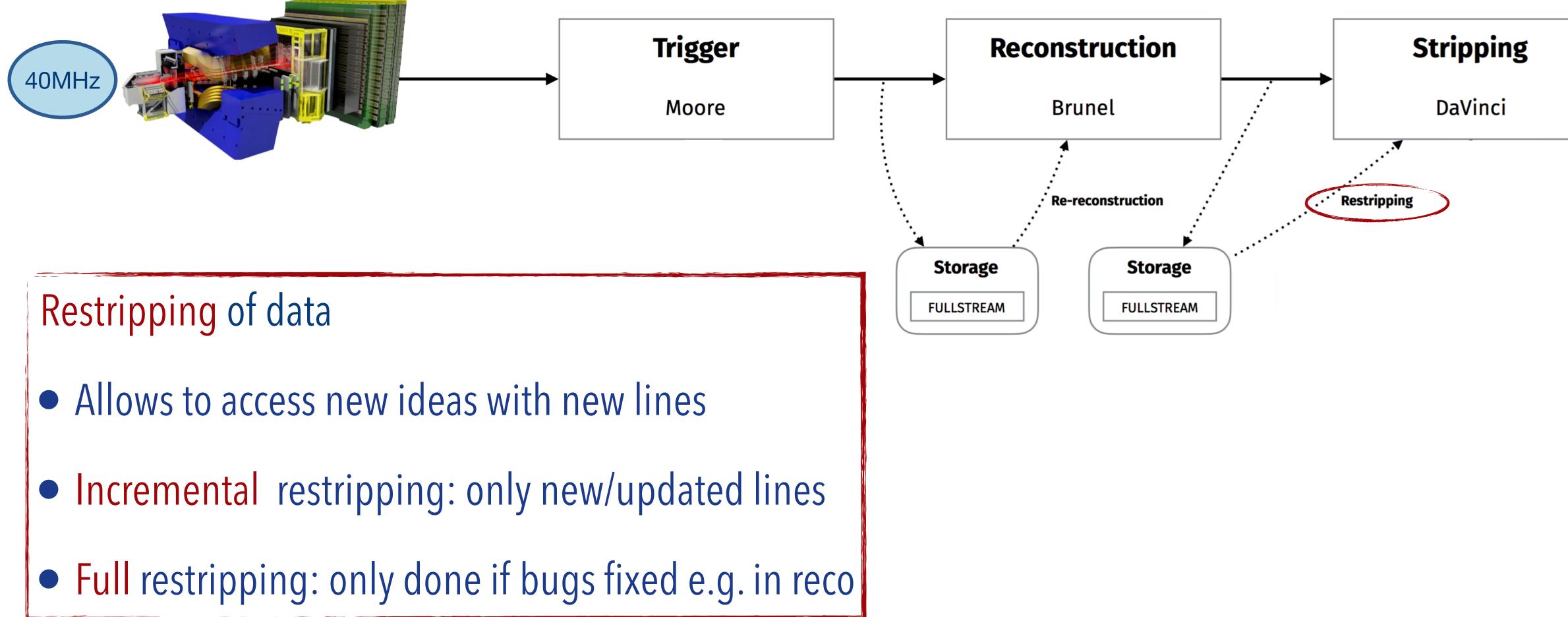
- Select certain decays based on signatures
- Exclusive: only one decay
- Inclusive: several decays combined
- Special lines: minimum-bias, BKG studies, ...

Reducing the file sizes

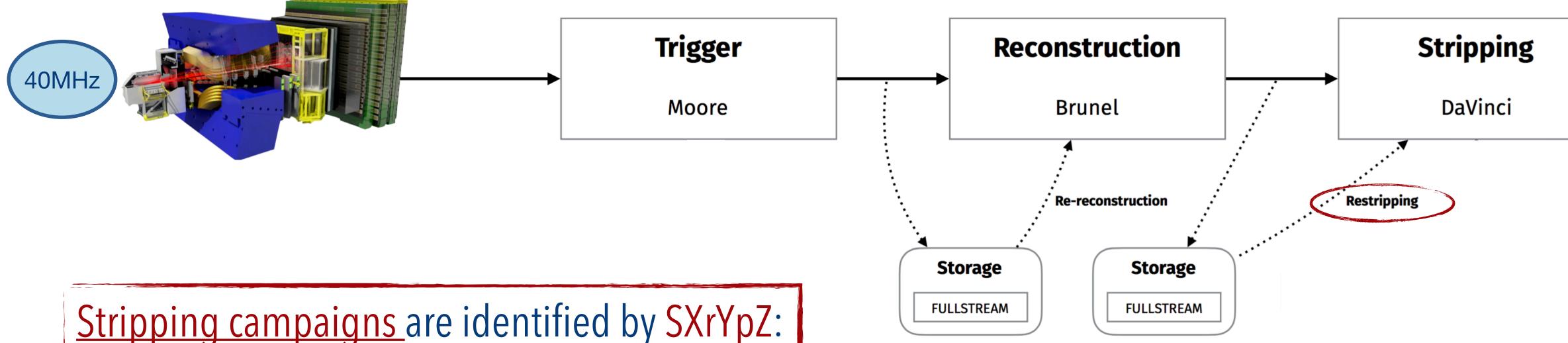




What about new ideas?



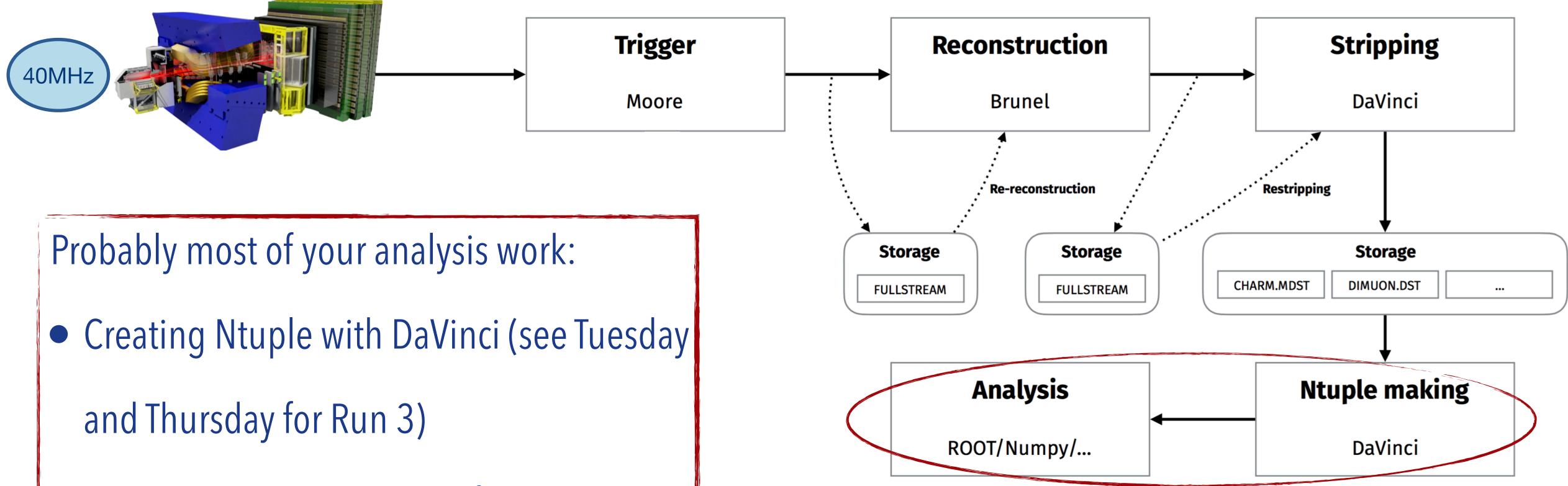
What about new ideas?



Stripping campaigns are identified by SXrYpZ:

- X: Full restripping campaign
- Y: data taking Year
- Z: incremental restripping

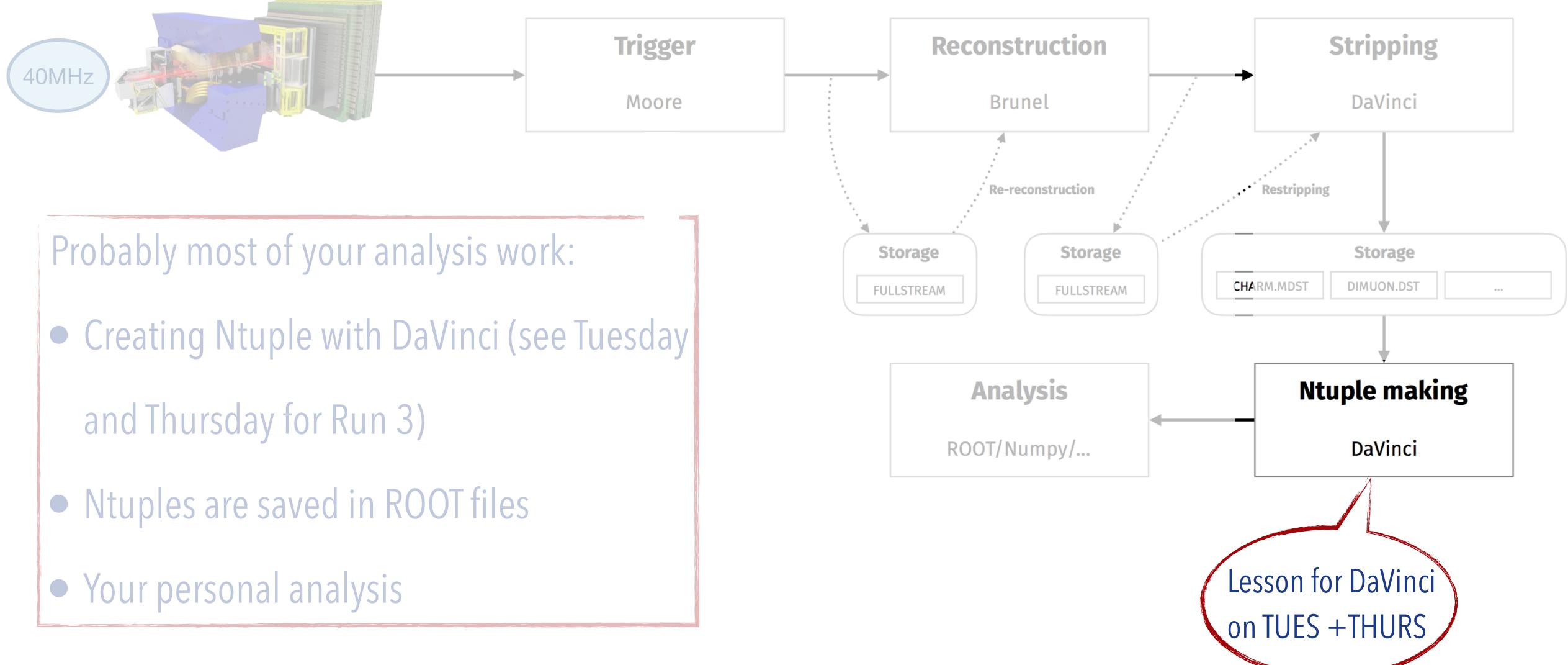
And finally:



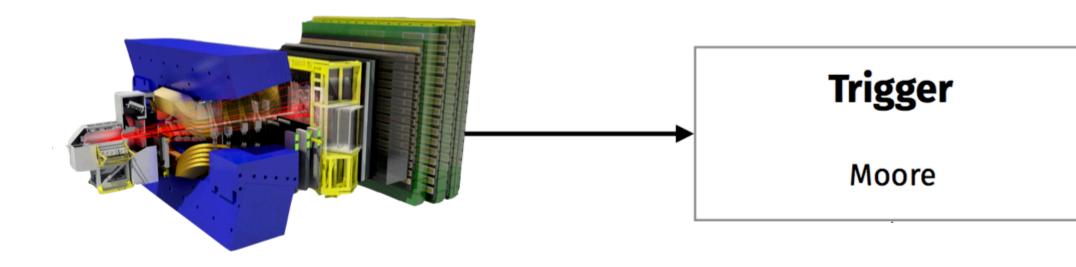
- Ntuples are saved in ROOT files
- Your personal analysis



And finally:

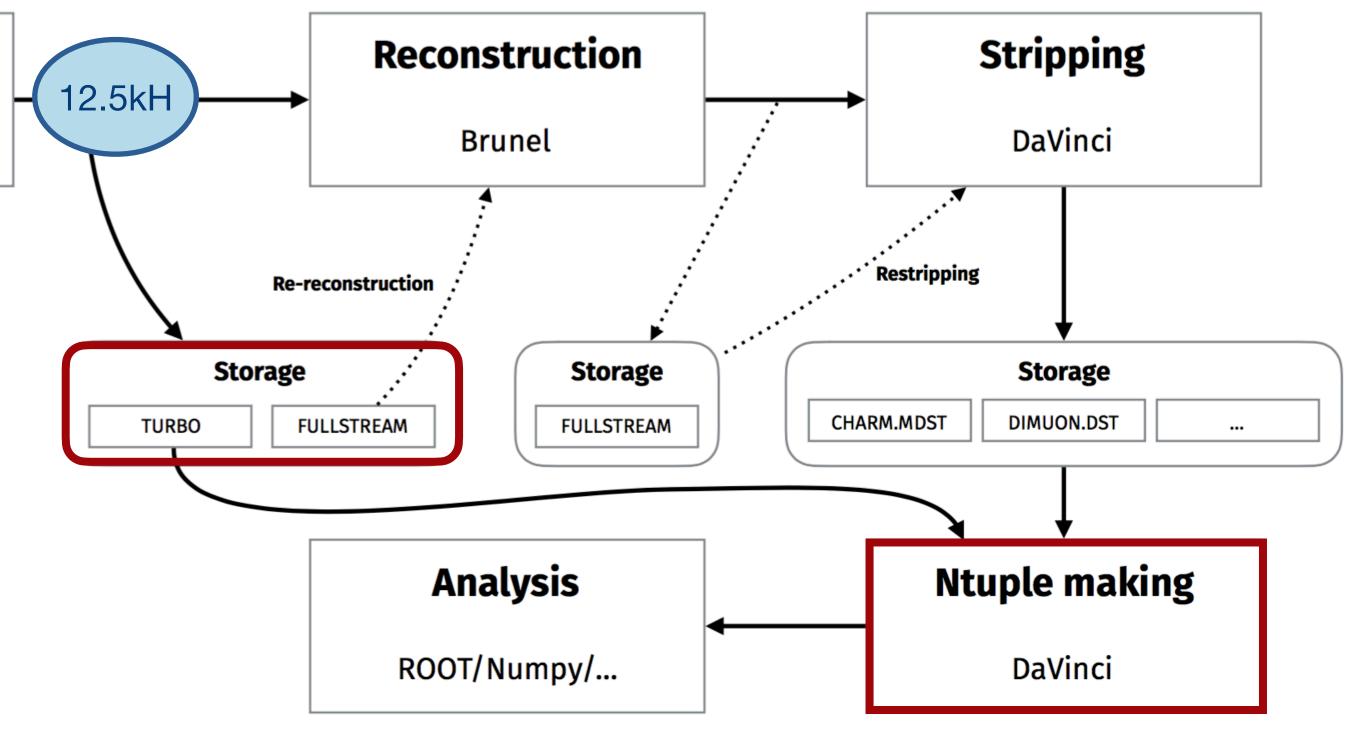


What has changed?

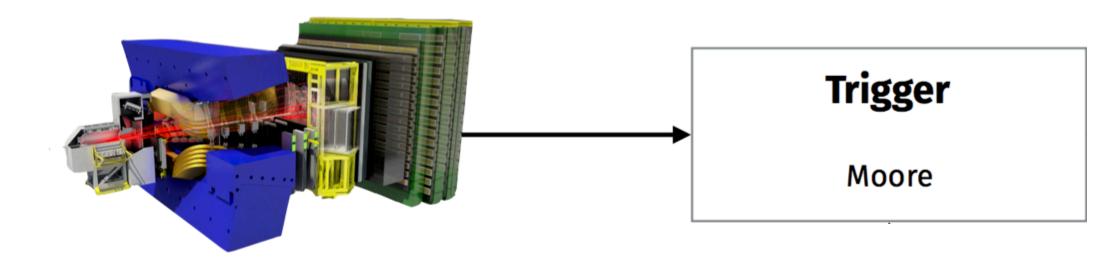


Upgrade of the event filter farm (EFF):

- HLT execution on EFF
- Improved hardware and software, automated calibration
- Full reconstruction now on HLT2!
- No offline reconstruction needed



Some details about the time for high level trigger

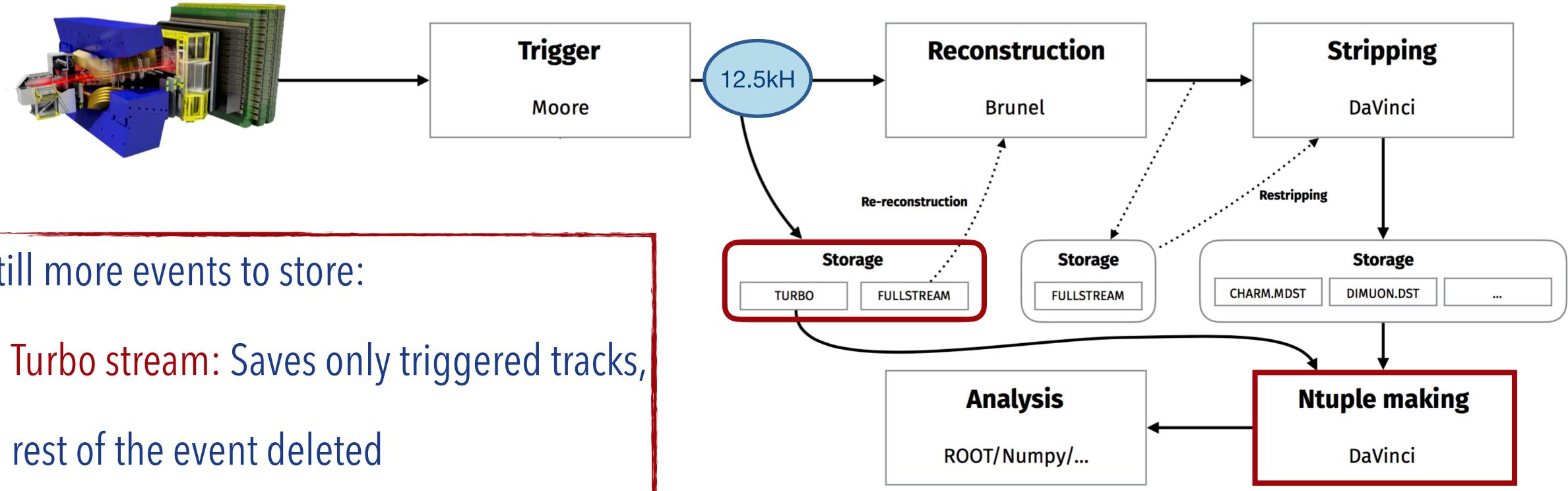


Next software stage called High Level Trigger

- HLT1: ms per event
- HLT2: full reconstruction takes hours

Alignment and calibration on 10Pb of buffer: mins/hours

Opens the option to bypass with Turbo



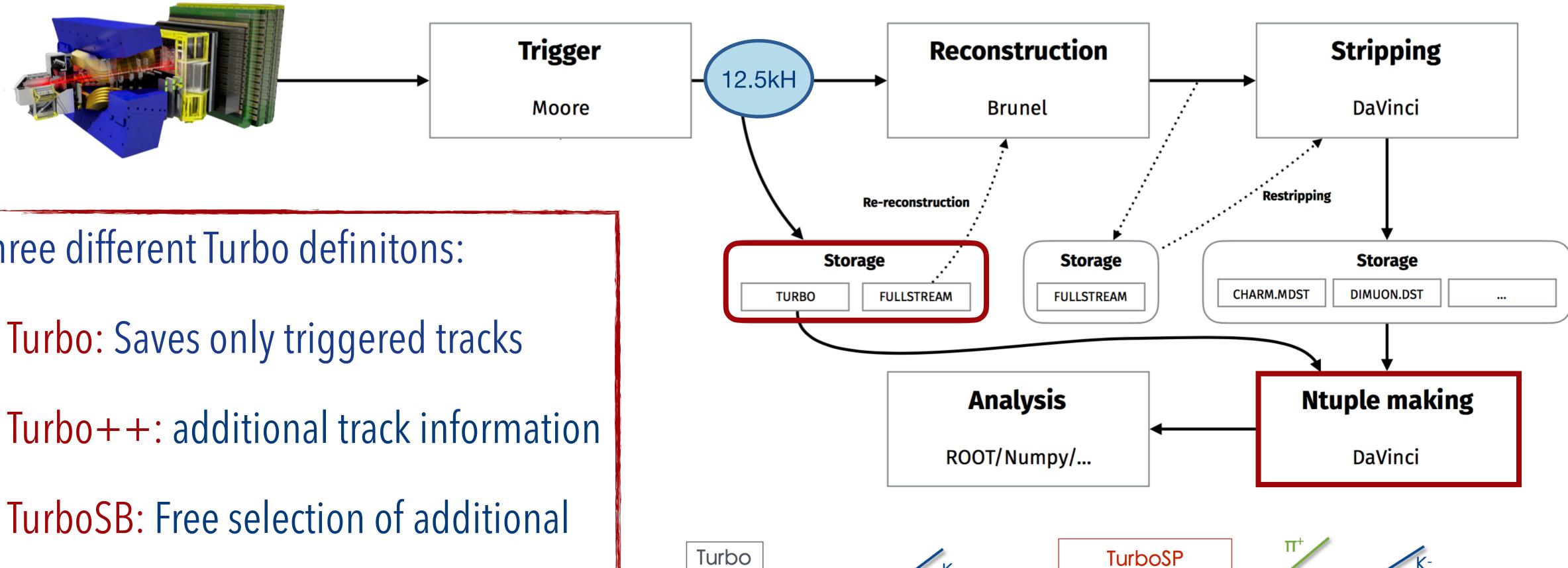
Still more events to store:

Turbo stream: Saves only triggered tracks,

- Cannot be re-reconstructed
- For available lines ask trigger liaisons

Collision data Flow during Run2

Opens the option to bypass with Turbo



PV

SV

Three different Turbo definitons:

- Turbo: Saves only triggered tracks
- Turbo++: additional track information
- TurboSB: Free selection of additional

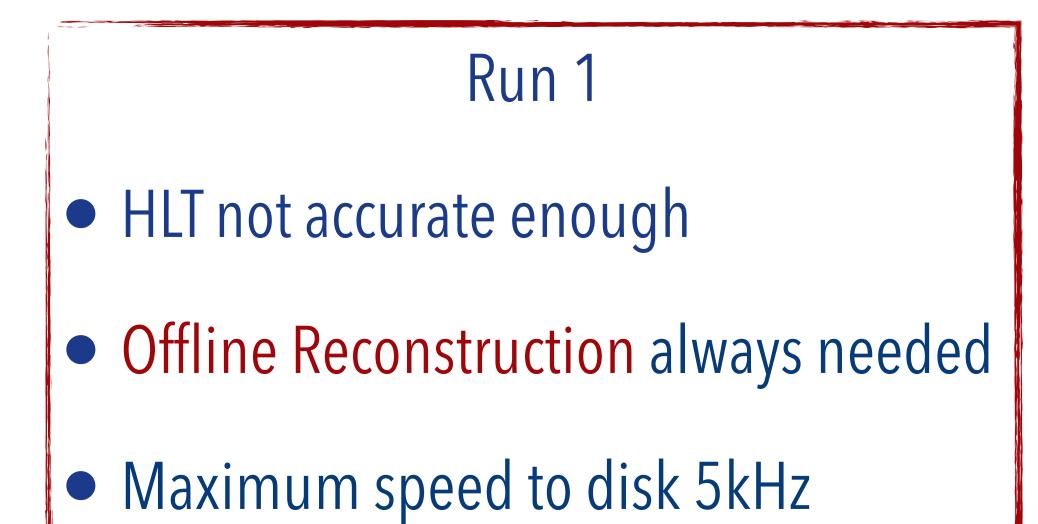
information to save



SV

PV

Summary collision dataflow



Summary collision data flow



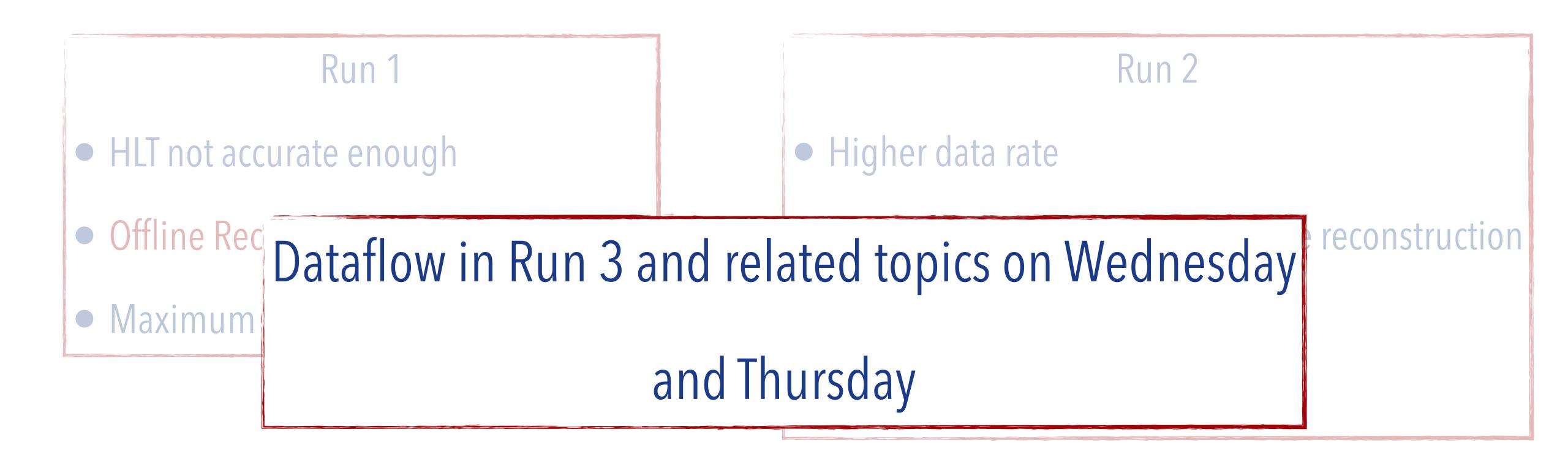
- HLT not accurate enough
- Offline Reconstruction always needed
- Maximum speed to disk 5kHz

Run 2

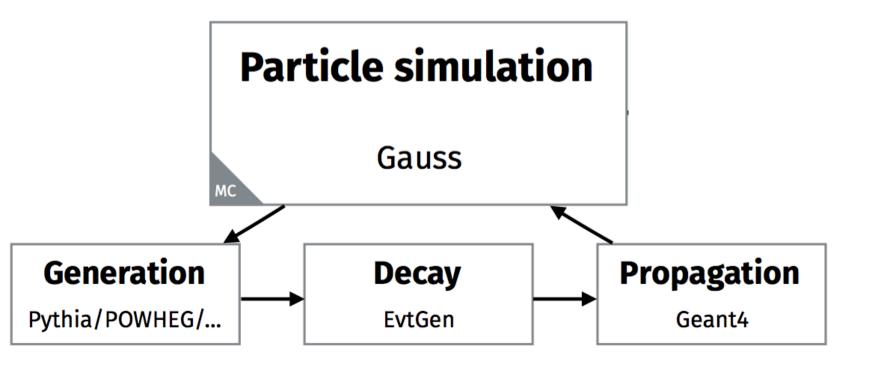
- Higher data rate
- HLT same accuracy without offline reconstruction
- Turbo Stream as bypass option
- Maximum speed to disk 12.5kHz



Summary collision dataflow



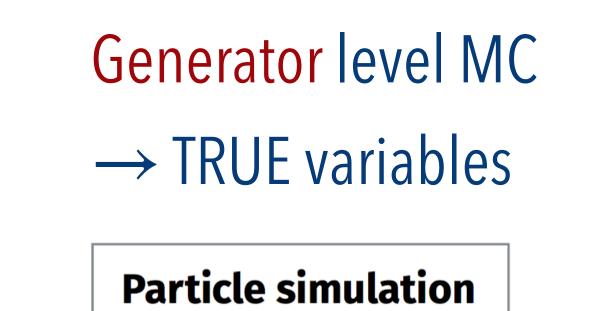
But what about simulation?

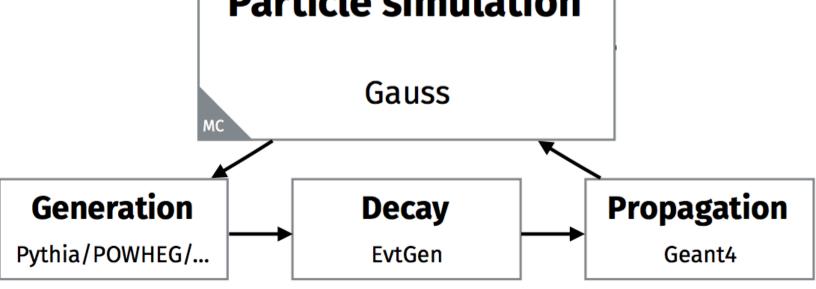


Creating particle simulation:

- Generation of the hard process e.g. Pythia
- **Decay** processed with <u>DecFiles</u> in EvtGen
- Propagation through detector: Geant4
- All executed in <u>Gauss framework</u>

But what about simulation?

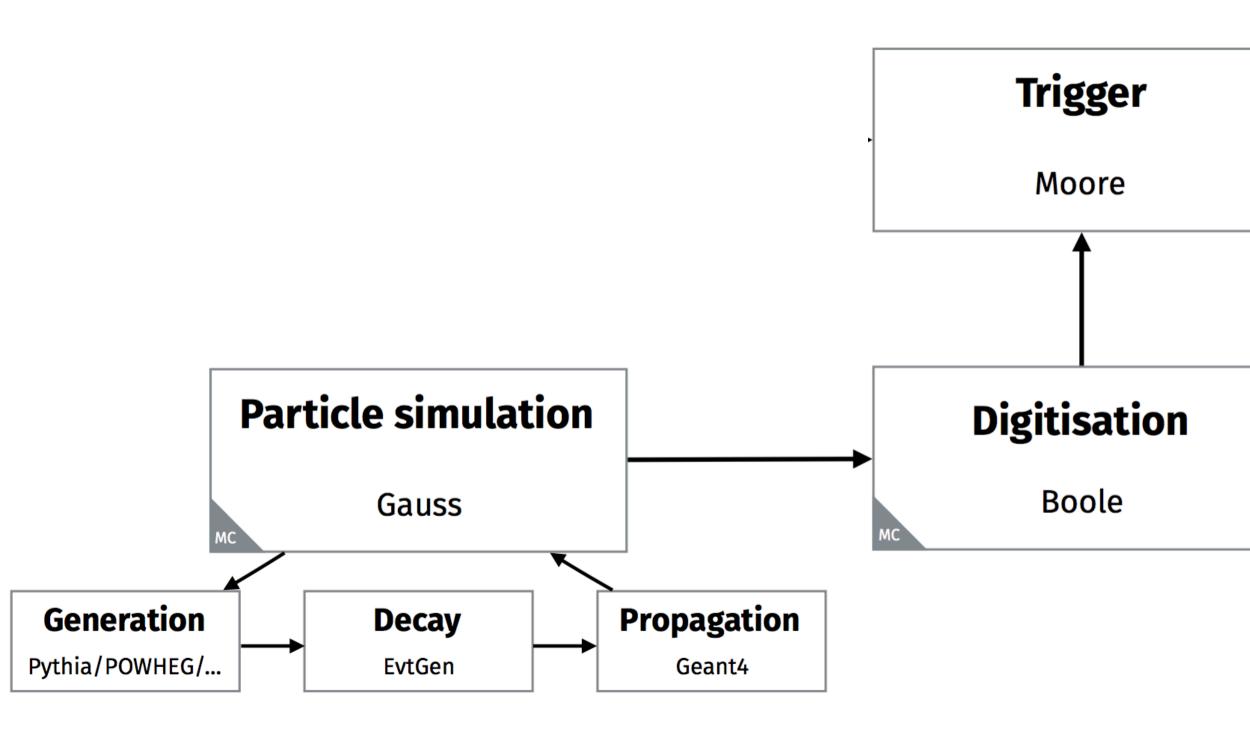


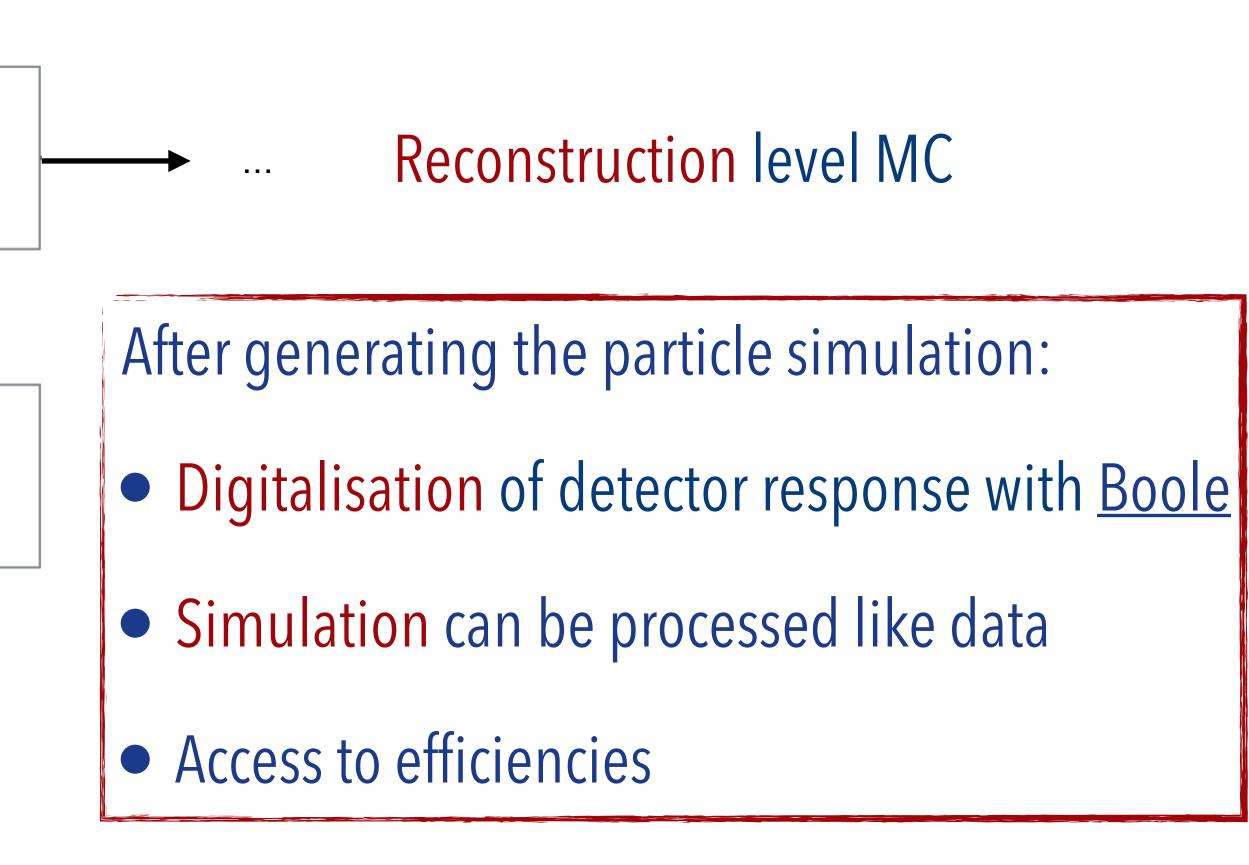


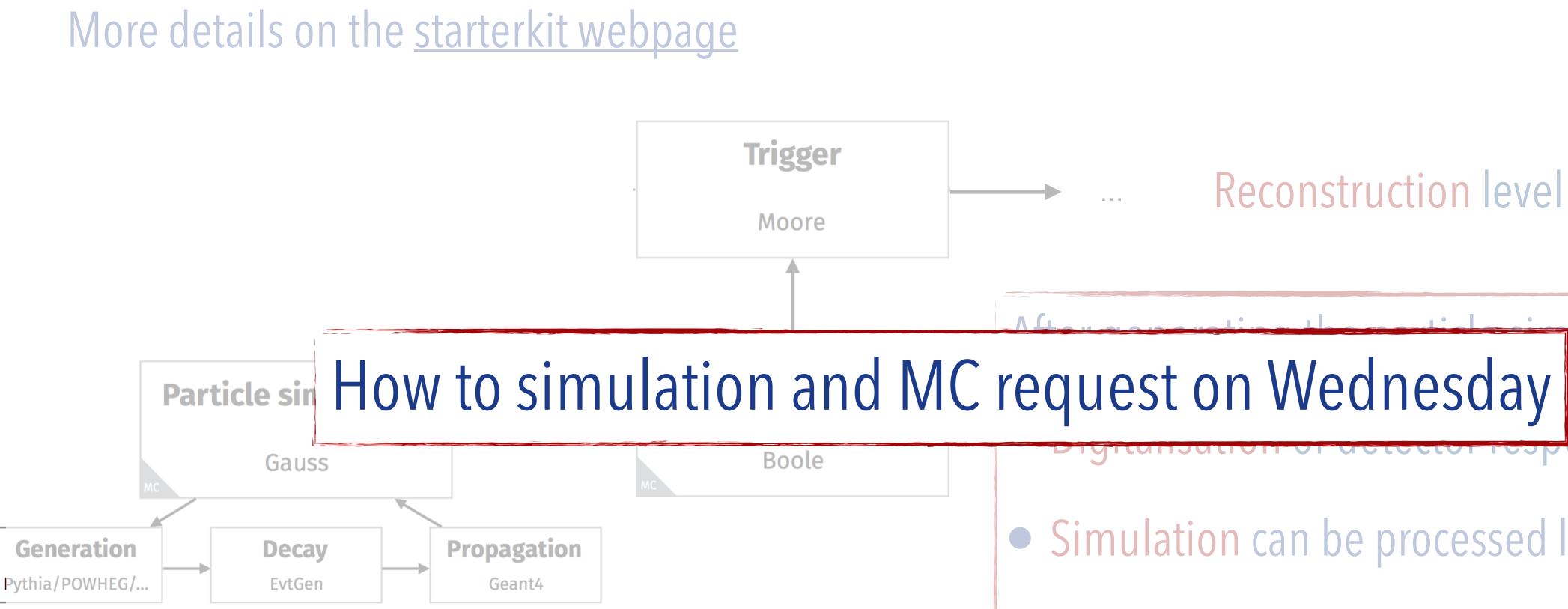
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- Propagation through detector: Geant4
- All executed in <u>Gauss framework</u>

More details on the <u>starterkit webpage</u>







Reconstruction level MC

ulation: nse with Boole

• Simulation can be processed like data

• Access to efficiencies



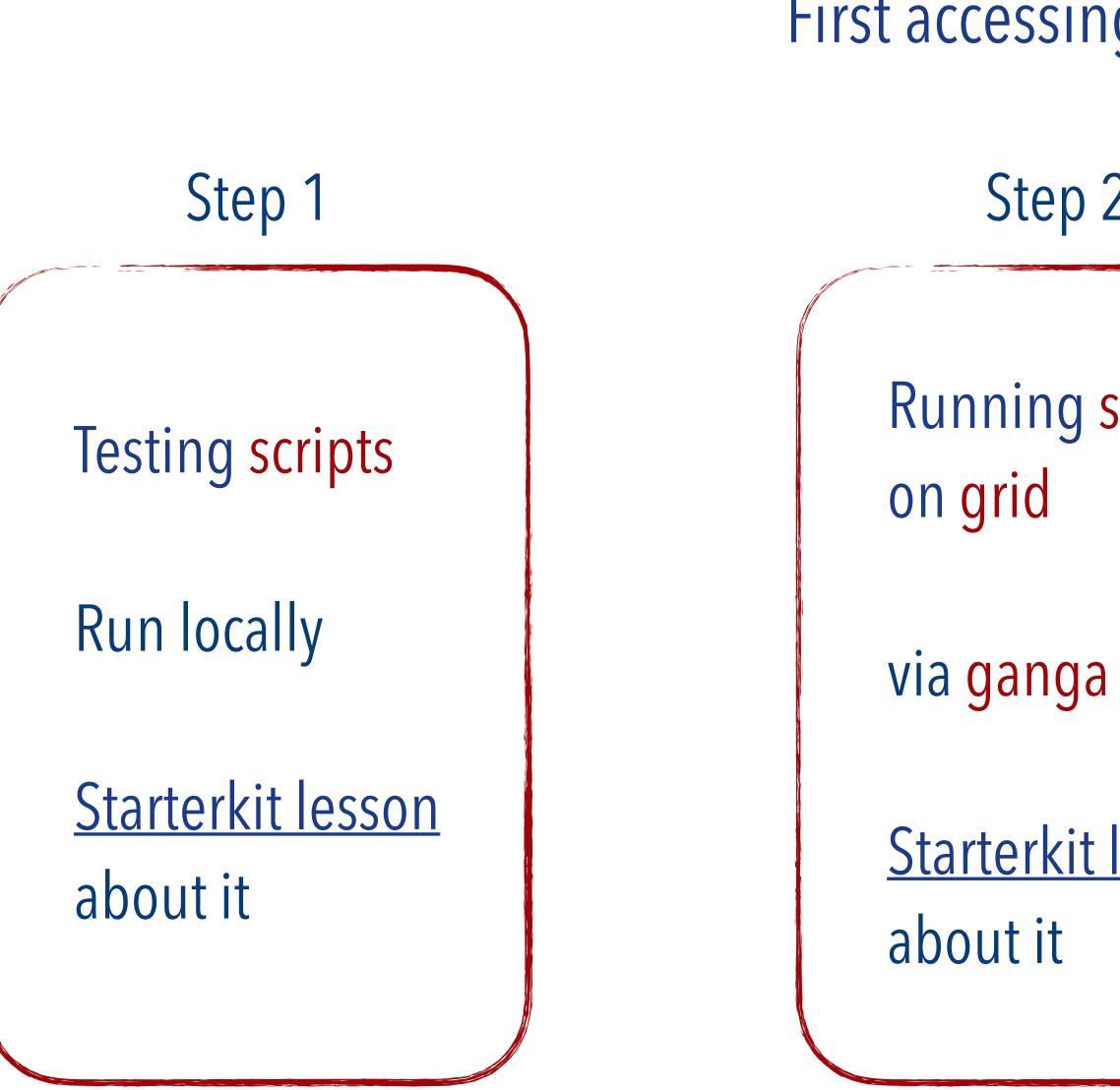
First accessing Ntuples

Step 1

Testing scripts

Run locally

<u>Starterkit lesson</u> about it



First accessing Ntuples

- Step 2a
- Running scripts
- Starterkit lesson



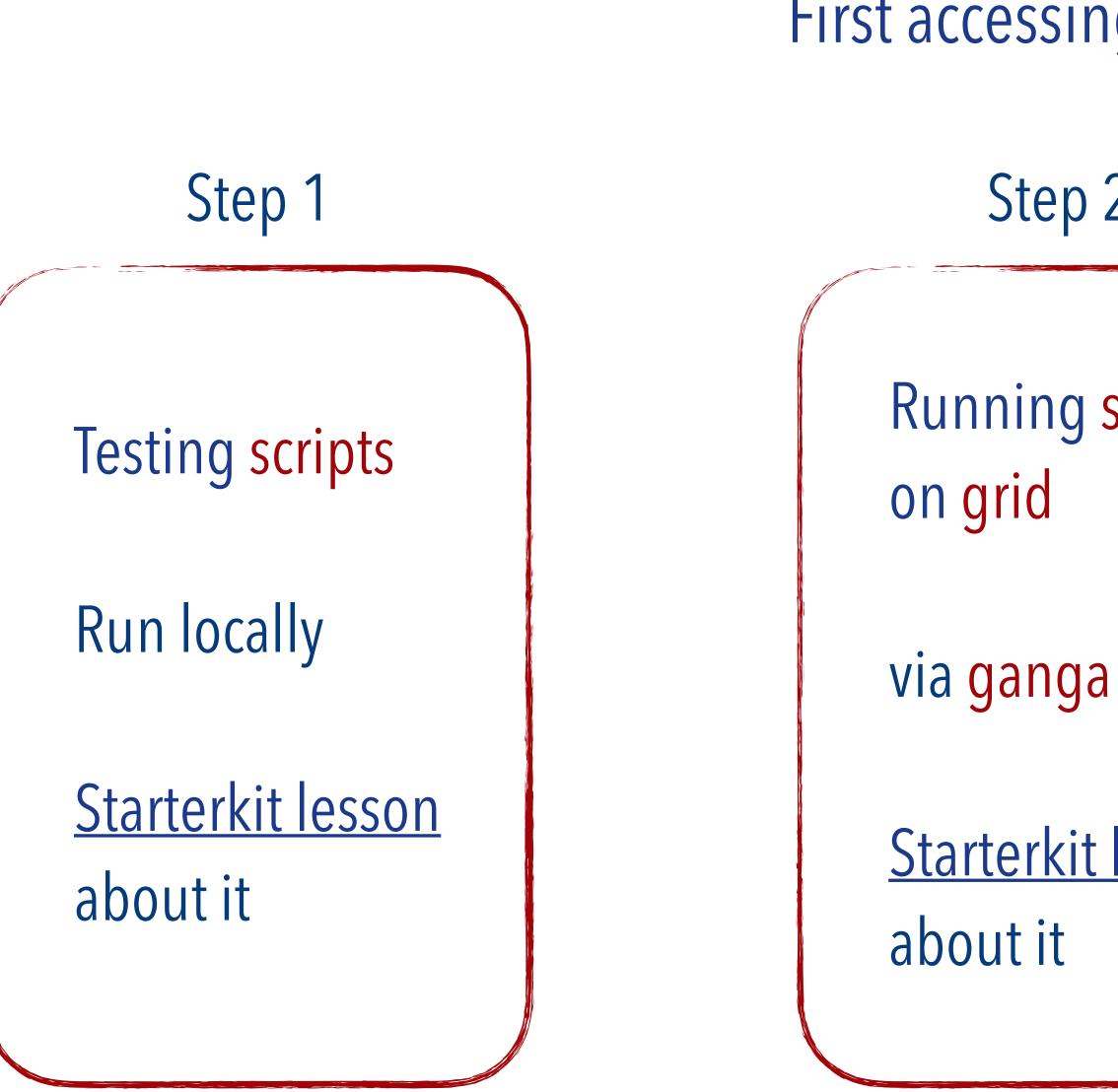
First accessing Ntuples

Step 2a

inning scripts

Can we do something more centralised to reduce human error?

Starterkit lesson



First accessing Ntuples

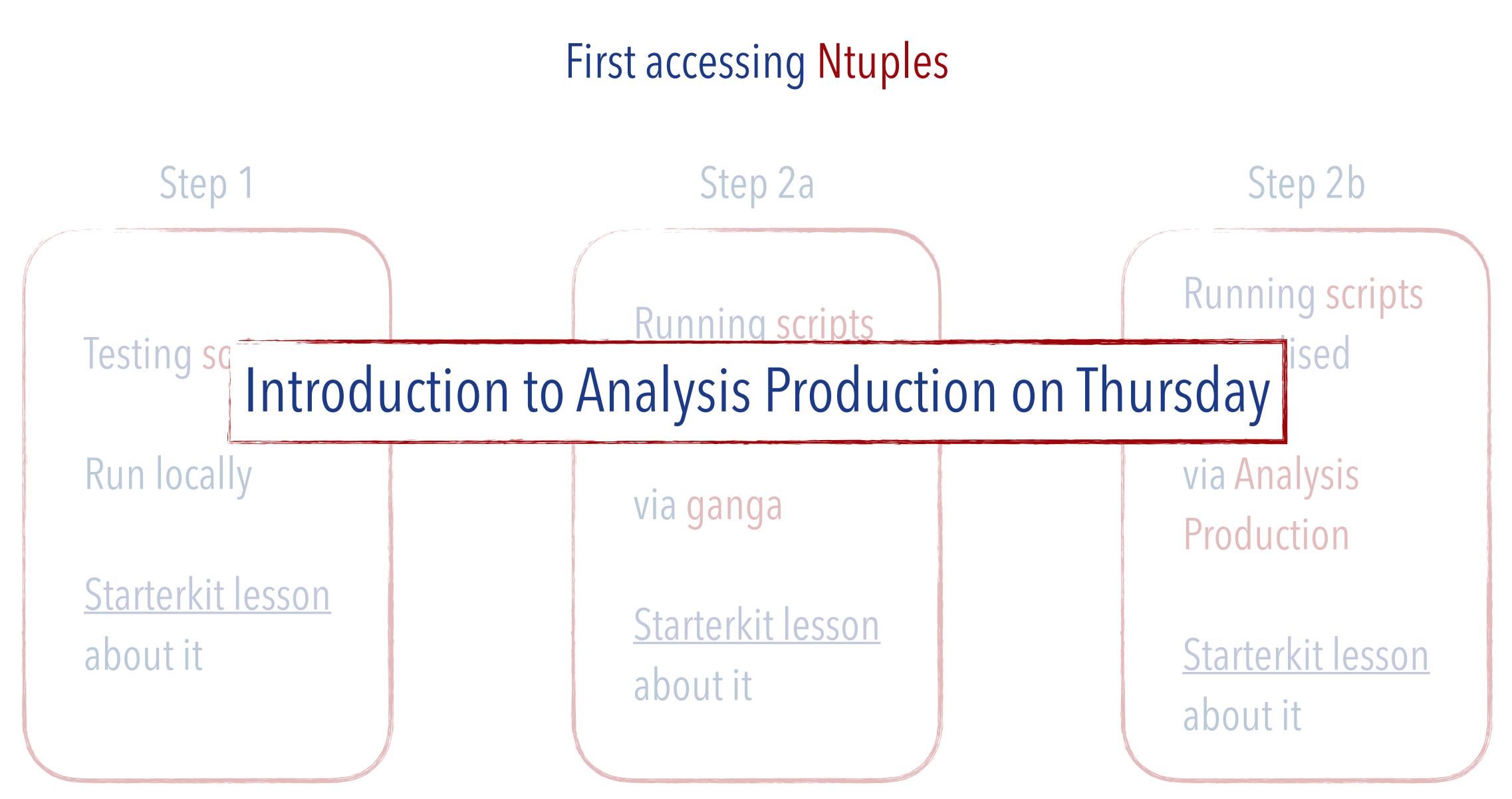
- Step 2a
- Running scripts
- Starterkit lesson

Step 2b

Running scripts centralised

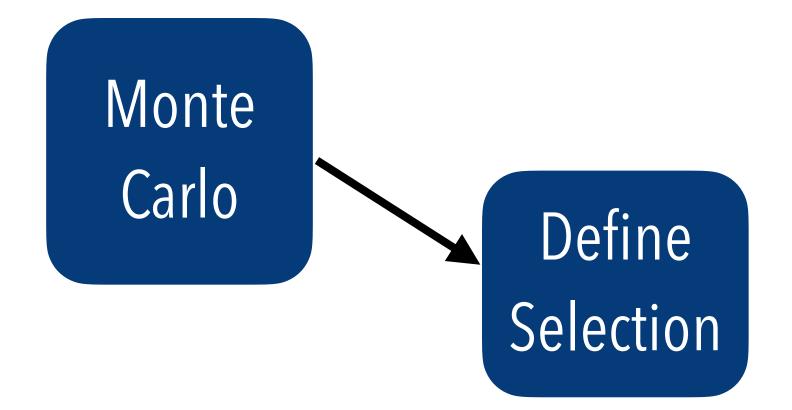
via Analysis Production

Starterkit lesson about it

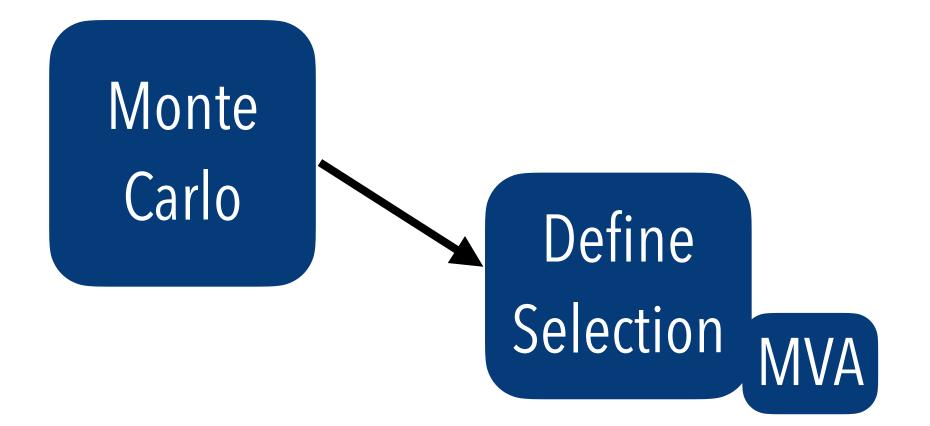


Ntuples Monte Carlo

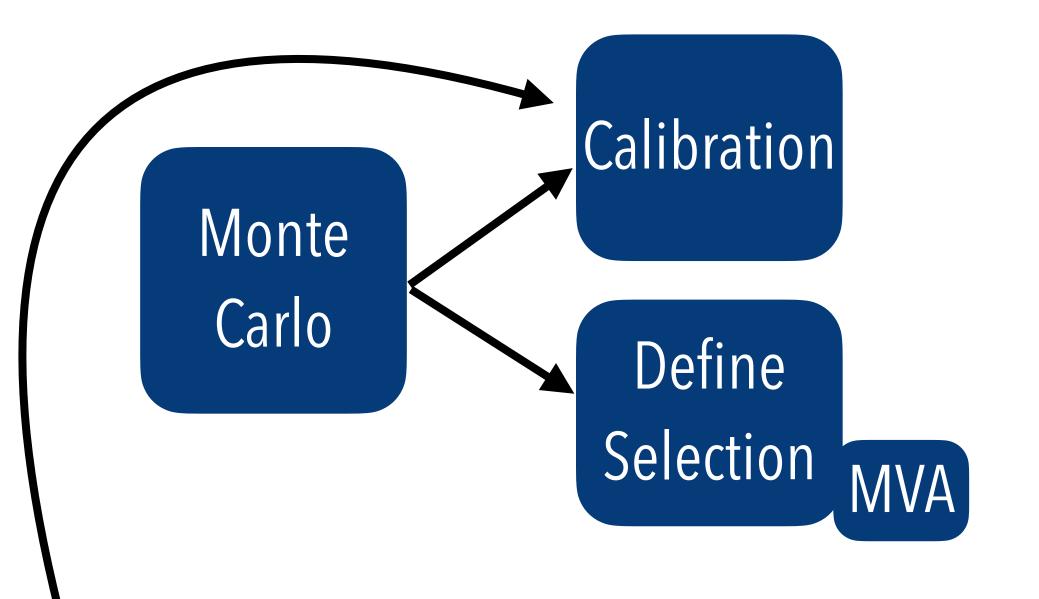






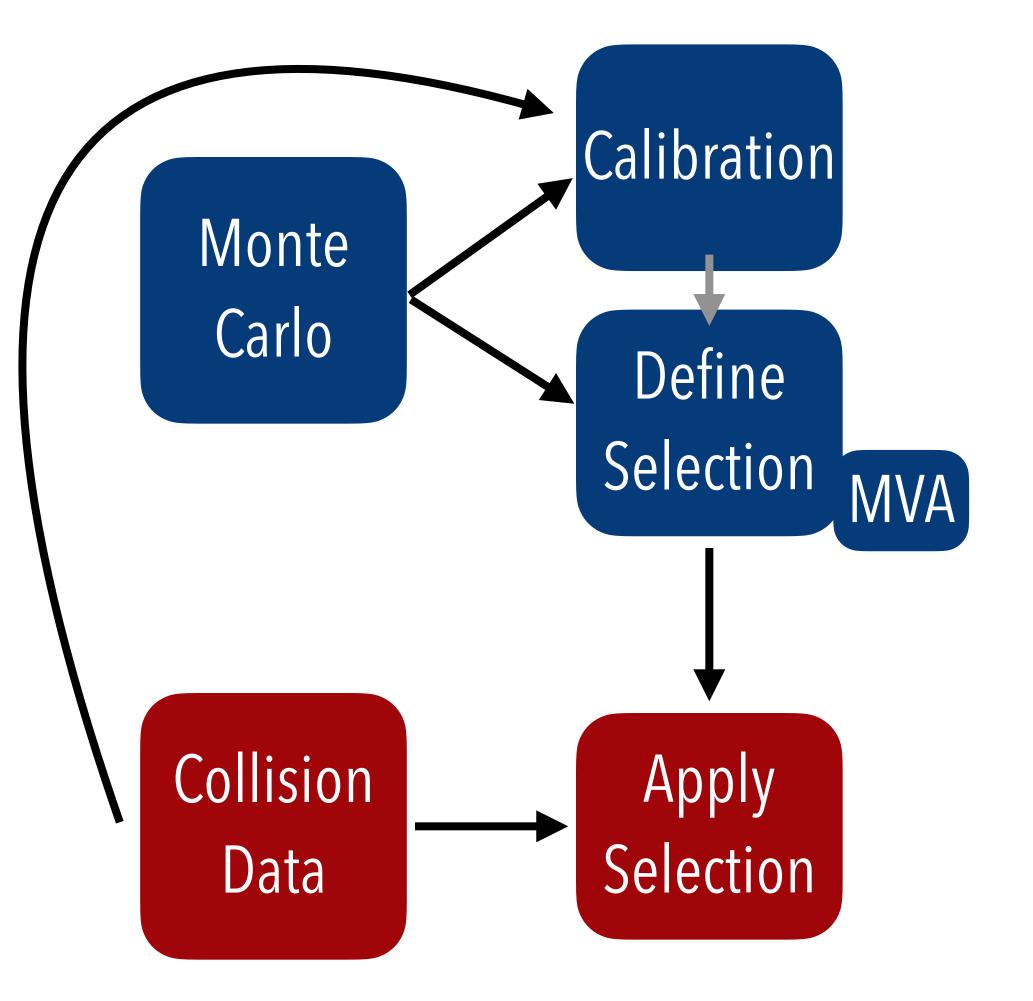


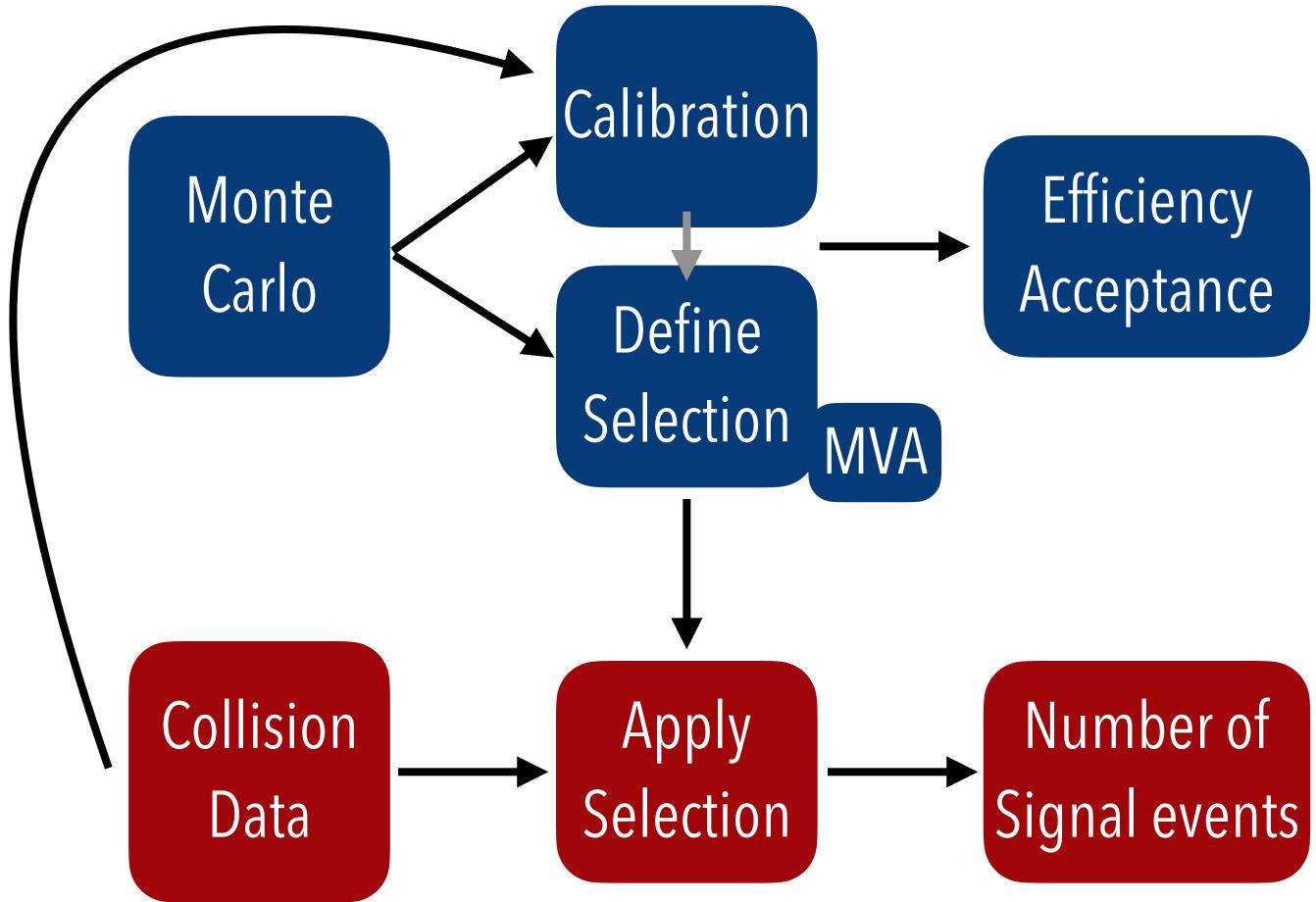


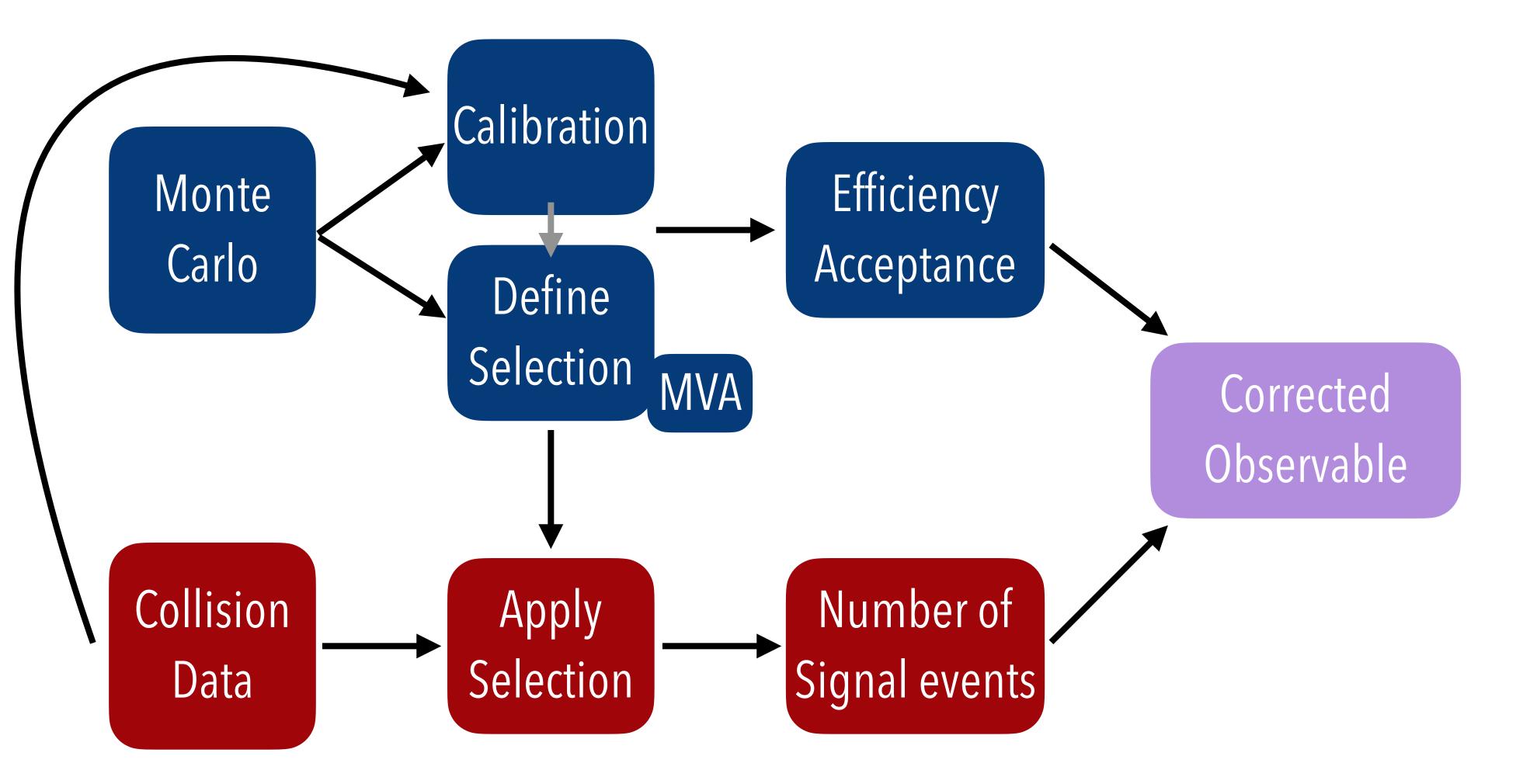


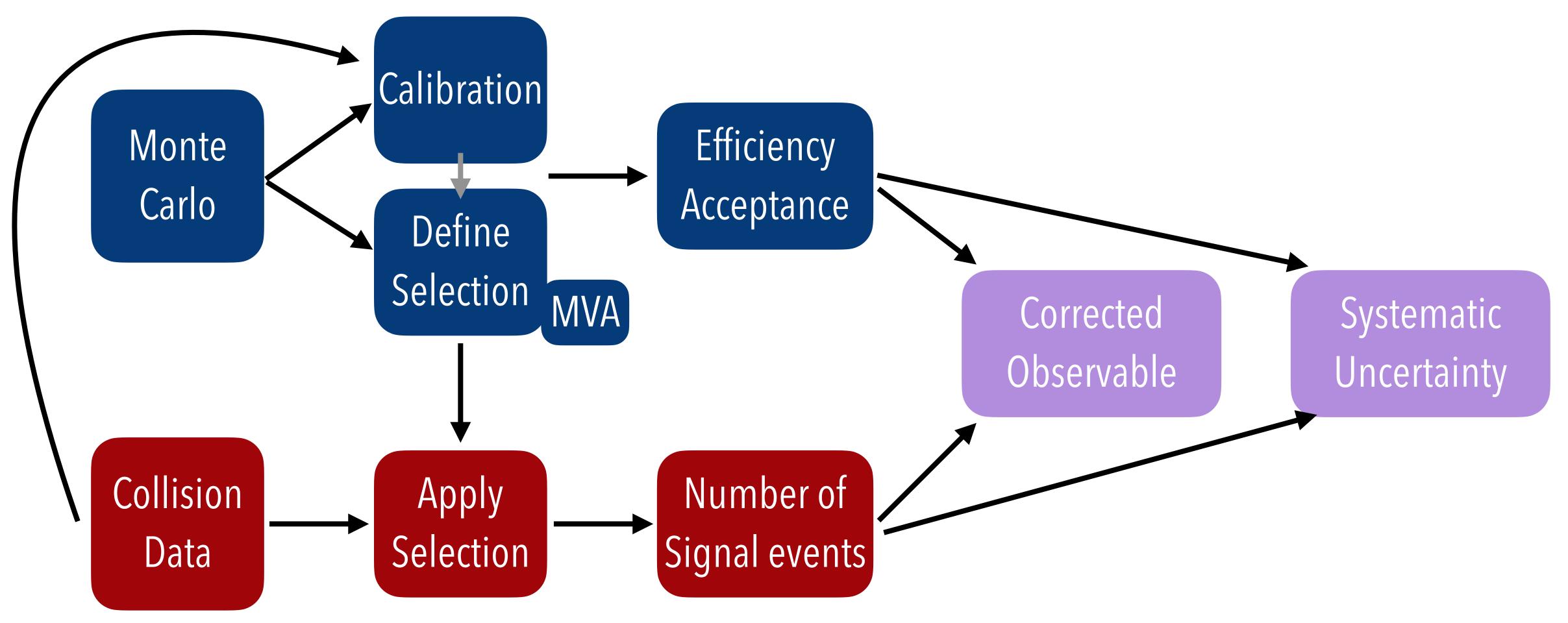


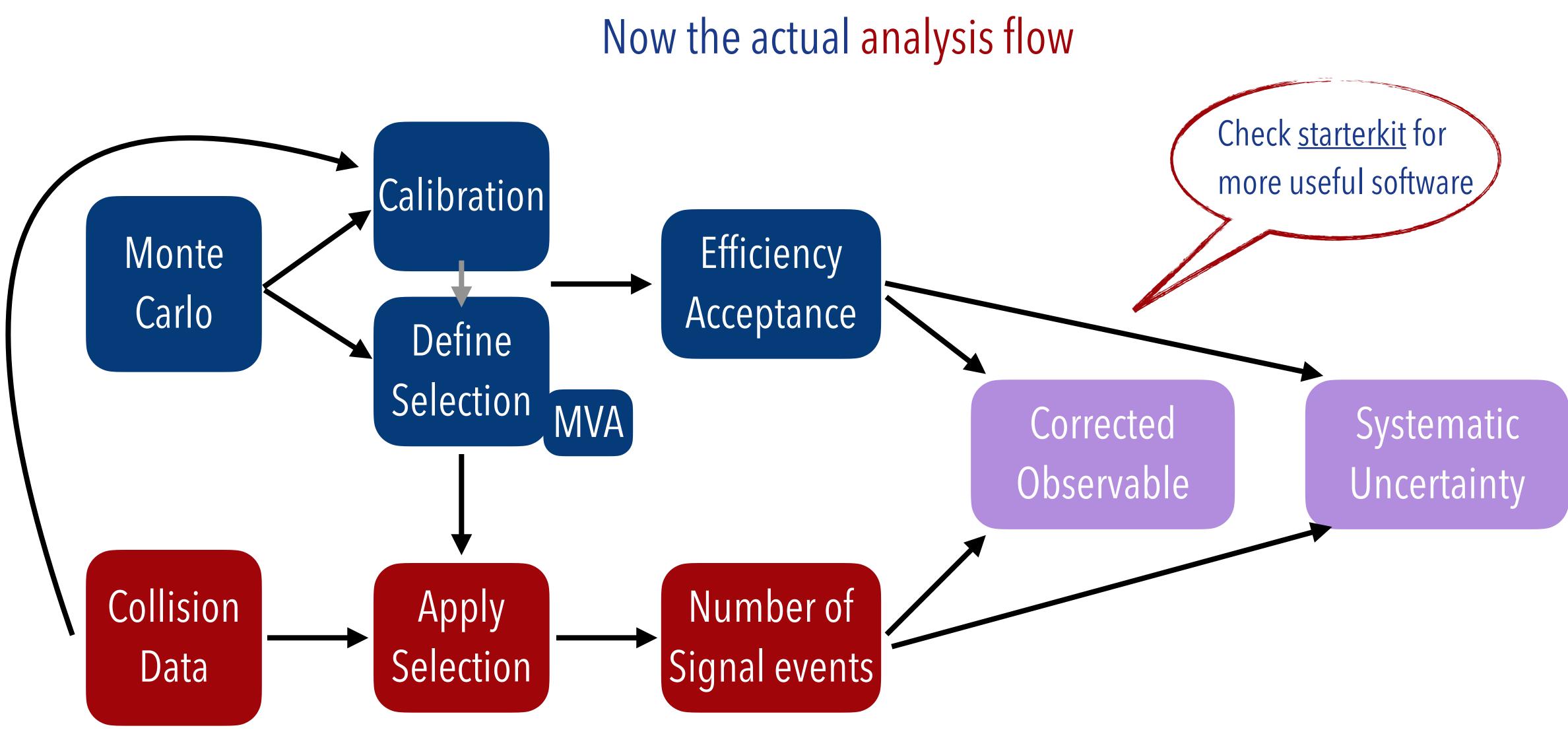
LHCb software: PIDCalib, TrackCalib,...

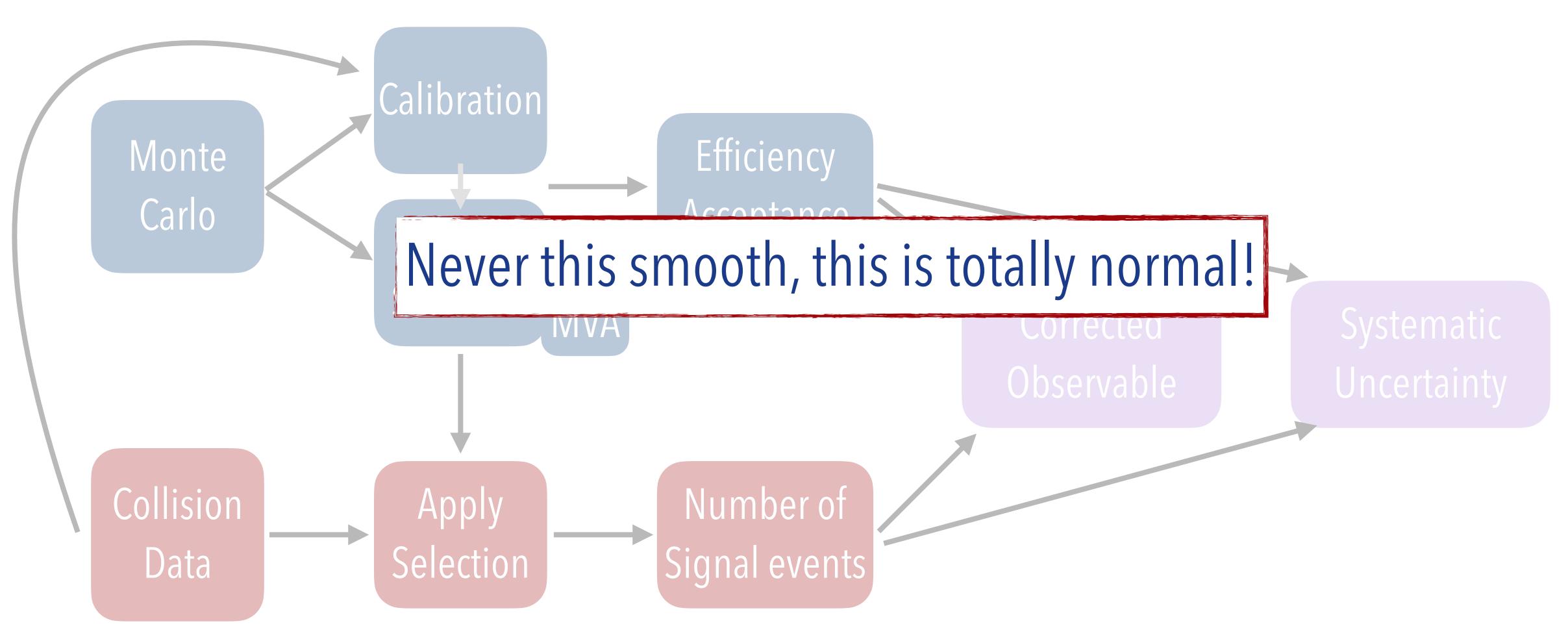












Analysis now in principle done:

• Analysis note: Contains all studies, documentation of your analysis, published on <u>CDS</u> at the end



Analysis now in principle done:

- after answering all questions (from talks and readers) WG approval talk

• Analysis note: Contains all studies, documentation of your analysis, published on <u>CDS</u> at the end • Working group review: AnaNote in good state, working group reader, WG pre-approval talk,



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 Working group review: Analysis in good state, working group reader. WG pro approval talk
- Working group review: AnaNote in good state, working group reader, WG pre-approval talk, after answering all questions (from talks and readers) WG approval talk
- Physics reviewer: Will be assigned upon request of WG convener, same procedure as with WG, update of AnaNote, give ok to unblind with WG convener (if necessary)



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 - Reviewing paper draft, if happy request Editorial Board Reviewer



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- Collaboration wide review process: Approval to go to paper talk, institute reviewers, two rounds, all comments need to be addressed, followed by all reviewer and physics coordinator



Analysis now in principle done:

- Working after ansv
- Physics re update of
 - And ask your colleagues.
 - In the end we are a collaboration! :)

• Reviewing paper draft, if happy request Editorial Board Reviewer all comments need to be addressed, followed by all reviewer and physics coordinator

• Analysis note: Contains all studies, documentation of your analysis, published on <u>CDS</u> at the end

al talk, Helpful are the LHCb guidelines for the preservation, the flowchart for review steps, the Publishing FAQ with WG,

- Collaboration wide review process: Approval to go to paper talk, institute reviewers, two rounds,



ow to

Analysis preservation needs to things

- If you don't use the lb-conda environment, preserve the package versions of the software e.g. with your own <u>conda environment</u> or a <u>docker container</u>
- Analysis code need to be accessible on <u>Gitlab</u>, the use of snakemake can make it easier to make your workflow reproducible



s of the software Never heard of snakemake? Analysis code No problem, tutorial about it on Thursday make it easier