# Improvements in the Cellular Automaton 

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## Outline

- Fake rate: xy plane cut, hard pt cut
- Timing: all in one
- Future plans


## Cells creation and connection

- At the moment a Cell is constructed using the legacy Doublet Generator
- given a region ( pT _min, beamspot, LIP, TIP..) it matches two hits from different layers if they are compatible with the region
- in the near future a cell will be constructed when a doublet is found
- When a Cell is created, the compatibility with all the cells in the previous layer pair that are sharing the same outer hit is checked



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## Compatibility

- The area of the triangle ABC in the RZ plane is given by:

$$
A=\left|z_{A}\left(r_{B}-r_{C}\right)+z_{B}\left(r_{C}-r_{A}\right)+z_{C}\left(r_{A}-r_{B}\right)\right|
$$

Hence the tangent of the angle in A is given by:

$$
\begin{aligned}
& \operatorname{tg}(\vartheta)=2 A / \\
& \theta_{0}=\frac{13.6 \mathrm{MeV}}{\beta c p} z_{c h} \sqrt{\frac{t}{X_{0}}}\left[1+0.038 \ln \left(\frac{t}{X_{0}}\right)\right] \\
& \vartheta * \mathrm{p}_{\min }<\mathrm{cut}
\end{aligned}
$$



- If two cells are found compatible they are pushed in each others' outer and inner neighbors vectors


## Compatibility

- Intersection between perpendicular bisectors of the two cells is found.
- Radius of the circle is then found
- No need to know where this circle and the circle given by (center=beamspot, radius $=\mathbf{T I I}$ ) intersect
- They intersect if the distance between the centers $\mathrm{d}(\mathrm{c} 1, \mathrm{c} 2)$ satisfies:

$$
\mathrm{r} 1-\mathrm{r} 2<\mathrm{d}(\mathrm{c} 1, \mathrm{c} 2)<\mathrm{r} 1+\mathrm{r} 2
$$

## - Hard pT cut:

- If the triplet's radius is less than a threshold(pTmin), the triplet is discarded
- default $0 \mathrm{GeV} / \mathrm{c}$



## All-in-one

- The quadruplet generator was taking sets of 4 layers and run a different CA for each layer set

$$
\begin{aligned}
& \text { layerList }=\text { cms.vstring( } \\
& \text { 'BPix1+BPix2+BPix3+BPix4', } \\
& \text { 'BPix1+BPix2+BPix3+FPix1_pos', } \\
& \text { 'BPix1+BPix2+BPix3+FPix1_neg', } \\
& \text { 'BPix1+BPix2+FPix1_pos+FPix2_pos', } \\
& \text { 'BPix1+BPix2+FPix1_neg+FPix2_neg', } \\
& \text { 'BPix1+FPix1_pos+FPix2_pos+FPix3_pos', } \\
& \text { 'BPix1+FPix1_neg+FPix2_neg+FPix3_neg' }
\end{aligned}
$$

This would result in many doublets, checks, evolutions run twice (or more)

## All-in-one ctd.

- In order to run only one CA for all the layer combinations, the hard dependency on the number of layers (as template parameter and in loops), had to be removed
- CAGraph was introduced to store the connections and the ordering between layers
- Given the input string from the Configuration it builds:
- Layer Graph (vertices visitor)
- Layer Pair Graph (edges visitor)
- Applied to out layer list it would result in...

For each hit on the layer, pointers to cells having that his as outer hit
layerList = cms.vstring(
'BPix1+BPix2+BPix3+BPix4',
'BPix1+BPix2+BPix3+FPix1_pos',
'BPix1+BPix2+BPix3+FPix1_neg', 'BPix1+BPix2+FPix1_pos+FPix2_pos', 'BPix1+BPix2+FPix1_neg+FPix2_neg',


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Cells are stored in a CALayerPair and are evaluated once.
Cell construction, matching, evolution is done using a BFS on this graph.


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## Filtering

- Approximate independent fits
- in the R-z plane, straight line + bending corrections
- in the x-y plane, circumference
- Reject quadruplets whose chi2 exceeds a threshold


## Performance

- See other attachment
- A solid and performant version of the CA is now being pushed in the release
- https://github.com/cms-sw/cmssw/pull/15751
- Although it would have been easier to implement graphs and manipulate strings using pointers, std::set, std::maps etc, this is not portable to CUDA. Everything was implemented using integral indeces and std::vectors
- I'll work until the end of the month to port all this new implementation to CUDA, and update the hackaton branch
- show the results at CHEP
- Run this CUDA+CMSSW prototype on different architectures
- Minsky (NVIDIA Pascal P100+ PPC)
- show the results at CHEP

