## Tales of Riemann.

## F2F tracking meeting - Pisa



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F HELMHOLTZ
ASSOCIATION

> Two dimensional fitting
> Three dimensional fitting
> Planes
> Two dimensional fitting

## Three dimensional fitting

Planes

## Versions of the Riemann fit

## Distance measure - fits to points

$$
\epsilon=n_{0}+n_{1} \cdot x+n_{2} \cdot y+n_{3} \cdot r^{2}
$$

(Frühwirth)

## Distance measure - fits to drift circles of radius I

This requires some prior knowledge of the right left passage hypotheses a as seen by the particle.


This difference is not neglectable especially for rather short segments.
Only using the drift length and the correct right left passage hypotheses yields accurate results.

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## CDC measurements are drift circles

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## Segment fits to wire position - display



## Segment fits to wire position - detailed display



Figure 1: Segment fits have too high curvature, more information needed to straighten trajectories.

## Segment fit to wire position - estimation quality

Scatter plot estimates versus truths


## Segment fits to drift circles - display



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## Segment fits to drift circles - detailed display



Figure 2: Using drift circles fits to axial segment yields right curvature. Fits to stereo segments have differing curvature. Here the curvature sign is opposite (blue vs. red).

## Axial segment fit to drift circles - estimation quality



## Difference between axial and stereo layers

## Axial layers

$>$ Projection carried out by the wires is the xy projection.
$>$ Curvature in xy projection is the real curvature.
> Unbiased fit in projection

## Stereo layers

$>$ Projection carried out by the wires changes with z position.
$>$ A particle trajectory can gradually pick up additional displacement as it is moving forward/backward.
$>$ Curvature in the projection and $\tan \lambda$ are interwined.
$>$ This generally leads to a bias.

## Bias in curvature residuals - stereo superlayer 1

## Curvature residuals in super layer 1

Continuous distribution


## Cause of the bias

## Stereo effect influences curvature in the $x y$-projection

Curvature residual versus tan lambda in super layer 1


## Extending the Riemann fit to estimate covariance matrices

## Covariance estimation

$$
V_{i j}^{-1}=\sum_{k} w_{k} \frac{\mathrm{~d} \epsilon_{k}}{\mathrm{~d} n_{i}} \frac{\mathrm{~d} \epsilon_{k}}{\mathrm{~d} n_{j}}
$$

## Application to Riemann fit

$>$ Simple in normal parameters $n$, since they enter the distance $\epsilon$ linearly.
$>$ Slightly complicating translation from four $n$ parameters to three perigee parameters.

## Covariance estimation quality

## Distribution of absolute curvature pull (clipped)



## Auxiliary implementation details.

$>$ Fitting methods interfacing to other code easily do able.
$>$ Transport / extrapolation of perigee covariances to new reference point.
$>$ Helix class also doing closest approaches to points.
$>$ Parameter and covariance matrix estimation yield reasonable results for axial layers.
$>$ Fitting in stereo layers alone leads to a bias.
$>$ On segment level fitting the wire position is not enough.
$>$ Active use of drift length is not optional.
$>$ The latter is requires the correct right left passage hypotheses.
$>$ Comparision of the right left passage to the Monte Carlo truth would be desireable.

## Two dimensional fitting

> Three dimensional fitting

Planes

## Changed logic in the second stage

## Pairs instead of triples

$>$ Use pairs of segments as cells for the cellular automaton instead of triples

## Benefits

$>$ Needs only one filter in the creation step instead of two.
$>$ Tracks are allowed to end in stereo layers.
$>$ Fewer left over segments
$>$ Still enough information to construct a helix with from combination of axial and stereo information

## Combining the axial and stereo fit

## Kalmanesk combination of parameters

Combined helix covariance

$$
V^{-1}=H_{a}^{T} \cdot V_{a}^{-1} \cdot H_{a}+H_{s}^{T} \cdot V_{s}^{-1} \cdot H_{s}
$$

Combined helix parameters

$$
x=V \cdot\left(H_{a}^{T} \cdot V_{a}^{-1} \cdot p_{a}+H_{s}^{T} \cdot V_{s}^{-1} \cdot p_{s}\right)
$$

Residuals

$$
r_{a / s}=p_{a / s}-H_{a / s} \cdot x
$$

Combined chi square

$$
\chi^{2}=r_{a}^{T} \cdot V_{a}^{-1} \cdot r_{a}+r_{s}^{T} \cdot V_{s}^{-1} \cdot r_{s}
$$

with perigee parameters $p_{a / s}$, perigee covariances $V_{a / s}$ and segment ambiguity matrix $H_{a / s}$ of axial and stereo segment respectively.

## Ambiguity matrix

## For axial segments

$$
H_{a}=\frac{\mathrm{d} p}{\mathrm{~d} x}=\left(\begin{array}{ccccc}
1 & 0 & 0 & 0 & 0 \\
0 & 1 & 0 & 0 & 0 \\
0 & 0 & 1 & 0 & 0
\end{array}\right)
$$

## For stereo segments

$$
H_{s}=\frac{\mathrm{d} p}{\mathrm{~d} x}=\left(\begin{array}{ccccc}
1 & 0 & 0 & 0 & 0 \\
0 & 1 & 0 & \zeta & 0 \\
0 & 0 & 1 & 0 & -\zeta
\end{array}\right)
$$

where

$$
\zeta=\frac{1}{\# \text { hits }} \sum_{\text {hits }} \frac{\text { wire vector }_{x y} \cdot \text { normal to trajectoy }_{x y}}{\text { wire vector }}
$$

## Fit results

## Distribution of $\tan \lambda$ pull (clipped)

Continuous distribution


# > Two dimensional fitting 

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## Things that need to be done

## Obvious

$>$ Eventually use the fit output in proper rejection of cells and neighborhoods.
$>$ Integrate the new track model (RecoTrack).

## Not so obvious

$>$ In superlayer segment merging
$>$ Break the forward backward symmetry.
$>$ Merge left over segments in superlayer 0 with VXD.
$>$ Reconstuct cycles in the cellular automaton as curlers.

