

CSC Overlaps Alignment: Review and Conclusions

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- ▶ We'll be talking about a discrepancy that we've previously described as an indication of an error in our track-based alignment
- Since then, we've significantly cleaned up the analysis, understanding and correcting track-based errors, and the discrepancy has reduced by a factor of 3
- By comparing final alignment results with photogrammetry, we have very strong circumstantial evidence that the remaining discrepancy is not due to misalignment or track-based issues
- ▶ We have two remaining hypotheses that we propose here to get expert input. If what we're suggesting is not possible, given what is known about the chamber geometry, then we (collectively) will have to scratch these out and find another explanation



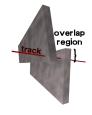
- Motivation and overview of the CSC Overlaps procedure
- Developments since CMS Week, including
 - narrow and uniform residuals distributions
 - ightharpoonup agreement with photogrammetry at the level of 300 μ m
- ▶ However, the ring still doesn't close (the "discrepancy" from page 1)
- ► Conclusions: what might be causing the lack of closure, and what has been ruled out



- ▶ Baseline alignment procedure shown to require 10–100 pb⁻¹ for a few hundred micron precision
- Quicker alternative:
 - relative alignment of chambers in each ring (CSC Overlaps)
 - \blacktriangleright align whole ring relative to tracker with ${\sim}1000s$ of quality tracks
- Can be done with CRAFT data

Overlaps procedure:

- select tracks that pass through overlap of chambers in a ring
- require consistency in pair of segments: slope and intercept
- 3. solve system





System is over-constrained: must be consistent with a circle ("closure")

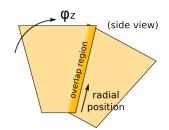


- ▶ Select good tracks, refit segments to 1-D straight lines: $\phi(z) = a + bz$
- lacktriangle Align $arphi_y$ angles by requiring segments to have equal slopes ($\Delta b=0$)



- Now intercept residual (Δa) tells us about misalignment in the $r\phi$ direction: align ϕ positions with fixed r (arcs centered on beamline)
- ightharpoonup Linear trend in intercept residual versus radial position is due to $arphi_z$ misalignment

Parameters decouple when corrected in this order: first φ_y , then $r\phi$, then φ_z (repeated alignment yields zero corrections)



Closure conditions

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▶ Four types of residuals, each measures a different parameter:

	segment slopes	intercepts
evaluated at center of chambers	$arphi_y$ angles	$r\phi$ positions
linear trend versus radial position	chamber twist	$arphi_z$ angles

- No significant "twist" observed (shear of alignment pins)
- Residuals measure chamber-by-chamber differences: in a ring, they must add to zero
- lacktriangle All types of residuals add to zero ("close") except $r\phi$
- $ightharpoonup r\phi$ residuals fail to close if chambers are the wrong width in some sense:
 - ▶ if the chambers are literally too wide or narrow
 - if the average distance to the beamline (radius) is wrong (potentially measurable by closure, if we can rule out other effects)
 - lacktriangle if chambers are rotated in $arphi_{
 m y}$ (discussed in last DPG presentation)
- Last effect is second-order in φ_y , but we have a direct first-order measurement of φ_y angles

Status: last CMS Week

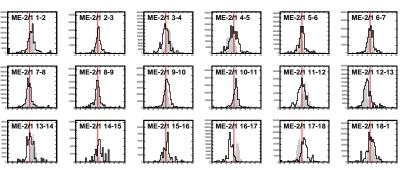
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- ▶ Only aligned φ_y and $r\phi$ (and some φ_y corrections were questionable)
- lacktriangle Wide residuals, presumed due to misaligned $arphi_z$

Hollow: aligned (ME-2/1 & -3/1 combined), grey: unaligned, red line: ME-2/1-only fit



- ► ME-2/1, ME-3/1 whole-ring underclosure: 18 mm, 20 mm
- Cross-checked overlaps alignment results against other track-based measurements (which are less precise than overlaps)

Status: now

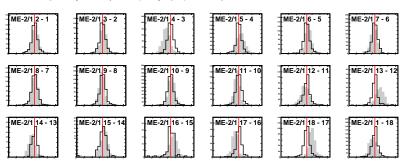
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▶ Full φ_v , $r\phi$, φ_z alignment; intercept residuals are all 1.2 mm wide

Hollow: aligned (ME-2/1-only fit), grey: unaligned, red line: common mean



- ▶ Used photogrammetry to set chamber-by-chamber distances from beamline: reduced underclosure by 20%, but still significant
- ightharpoonup ME-2/1, ME-3/1 whole-ring underclosure: 16 mm, 14 \pm 0.4 mm
- ▶ Interpreted as radial corrections: 2.5 mm, 2.3 mm inward
- Cross-checked track-based alignment results against photogrammetry

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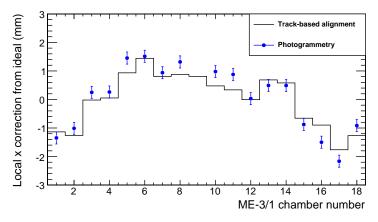
9/15





- The method is sound: Monte Carlo closure is consistent with zero
- In data, all residual types have zero closure except $r\phi$
- $r\phi$ and φ_z constants compare favorably with photogrammetry

Overlay of $r\phi$ translations, track-based and photogrammetry relative to ideal



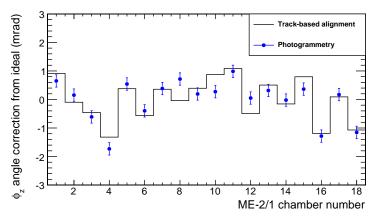
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10/15



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Overlay of φ_z rotations, track-based and photogrammetry relative to ideal

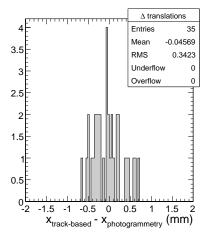


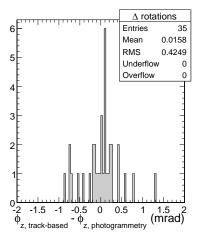
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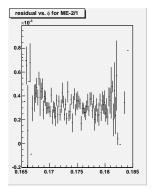
- \triangleright RMS difference between track-based and PG: 340 μ m, 0.42 mrad
- Photogrammetry $r\phi$ uncertainty is $(300/\sqrt{2}) \mu m = 210 \mu m$
- $\sqrt{340^2-210^2}=270~\mu \text{m}$ errors in track-based method alone
- ▶ Statistics from one large beam-halo run (62232, 9 minutes)

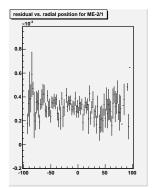


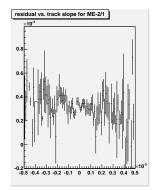




- ► High-precision agreement with photogrammetry gives us confidence in alignment results (with closure error distributed uniformly)
- ▶ Therefore, closure error is due to a systematic effect which is the same for each chamber
- \triangleright Are there trends versus ϕ , radial position, or track slope? No.











- ► Hypothesis 1: layers are *narrower* in real life than in software by 0.8 mm at the center (0.087% error in total width, or 10 μm per strip)
 - Maybe "narrowing" is due to bowing, e.g. if layer surface pinched to fit frame? No: sagitta = 9.6 mm, bow angle $= 4.8^{\circ}$
- ▶ Hypothesis 2: chamber centers are *closer* to the beamline by 2.5 mm in ME-2/1 and 2.3 mm in ME-3/1 (\pm 0.06 mm each)
 - We've verified the alignment pin positions in ϕ , and can therefore be certain that they are correct in r (photogrammetry errors must be isotropic in x and y!)
 - Therefore, we're not disputing the global pin positions, but the chamber centers relative to those pin positions (from mf.xml)
 - Also, this is not the chamber center as measured by the wires (the normal way of measuring radial position), but measured through angle of strips

Ideas to search for the effect

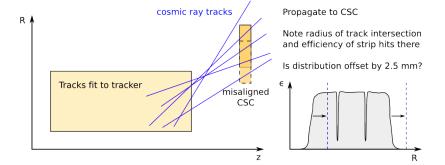
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14/15



- ▶ It was a *different* 2.5 mm offset that we found a few months ago by looking at the drawings (ME2/2 and 3/2, and in opposite direction)
- ▶ We didn't see an analogous effect in other rings, but there might be a different drawing/DDD discrepancy, e.g. distance from pins to center, rather than bottom of chamber frame to center
- ► Track-based method to cross-check hypothesis #2 (radial shift): wire residuals or strip efficiency turn-on curve (more direct)





- \triangleright Our alignment results are correct, to desired precision (300 μ m)
- \blacktriangleright However, $r\phi$ residuals very clearly do not close when summed around rings ME-2/1 and ME-3/1
- ▶ The problem is uniform across all these chambers: if it were not, we would not be able to reproduce photogrammetry
- Real chambers are than in simulation
 - ▶ 0.8 mm narrower across the center
 - 2.4 mm closer to the beamline.
 - ▶ pinching layers such that they bow by 4.8° more
 - ► 50-70° C colder
 - or . ?