

Chamber Alignment with globalMuons

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- Issues in global alignment
 - consistent tracker-muon coordinate system
 - magnetic field errors
 - single-scattering in material
 - cross-checks: is it a real alignment?
- Alignment produced for CRAFT analyses (CRAFT_ALL_V9)
 - sample map-plots, values of corrections, final residuals
 - cosmics track-splitting study
- Next steps in alignment
 - barrel improvements
 - alignment of CSCs, using barrel as reference
 - method for combining with hardware data

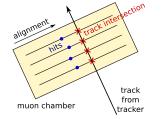
Consistent global coordinates

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- ► For optimal globalMuon resolution, we need to
 - ▶ align muon chambers relative to one another, and
 - put muon system in the same coordinate system as the tracker
- ► Can accomplish both in one step by using the tracker as a reference to align the muon system:



- 1. align the tracker (Alessio's talk, yesterday)
- 2. propagate tracks from tracker to muon layers
- 3. calculate unbiased residuals
- 4. adjust muon chambers to minimize residuals
- ▶ Tracker measurements dominate precision of most tracks anyway (tracks with $p_T \lesssim 200 \text{ GeV}$)
- ▶ Decouples "chicken-and-egg" problem of alignment: track-fitting is independent of geometry updates (no need for global fit or iteration)

Sources of systematic error

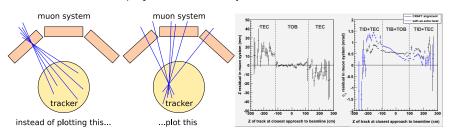
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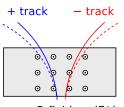
- ► Tracker misalignments: resolution, weak modes
 - use non-projective cosmic rays to look for distortions in tracker



- ▶ left: observation of TEC z misalignment (CRAFT_V4, not latest)
- right: sensitivity study, tracker twist added by hand (blue)
- ▶ Propagation errors: wrong \vec{B} -field, dE/dx, and track scattering
 - ▶ \vec{B} -field and dE/dx errors have distinct dependences on charge and momentum (next slides)
 - scattering yields non-Gaussian outliers, accommodate with fit



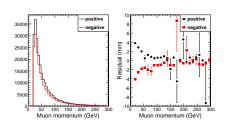
- Residuals from misalignment are independent of the tracks used to measure it
- ▶ Residuals from \vec{B} -field errors flip sign with the charge of the muon and depend on p_T



wrong B-field or dE/dx

Two-bin approach:

Fact: momentum spectra for + and - charges are proportional Fact: wrong \vec{B} -field and dE/dx effects are antisymmetric with q



- ► Find peak of residuals in two charge bins: R₊ and R_−
- ► Average $(R_+ + R_-)/2$ is sensitive to misalignment only
- ▶ Difference is sensitive to \vec{B} error and dE/dx errors only

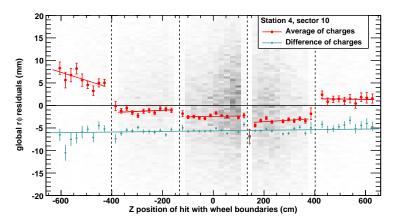
Demonstration in station 4

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- Station 4 has the largest \vec{B} -field errors: plot residuals across barrel
- ▶ The misalignment breaks cleanly at the chamber boundaries
- ▶ The \vec{B} -field error is independent of chamber



grey background is the raw 2-D residuals distribution linear fits are only a guide for the eye: not used in alignment!

Residuals with new $\vec{B}(\vec{x})$ maps Jim Pivarski 7/17





- ► Two new field maps available:
 - scaling corrections from segments (data-based measurement)
 - new TOSCA simulation (consistent field lines)
- ▶ Opportunity to test correctness of new $\vec{B}(\vec{x})$ and insensitivity of alignment measure with tracks propagated through new field
 - left: histogram of bins from the previous plot (for all sectors)
 - right: how each bin changes when new field is applied

statistical errors in bins are $\mathcal{O}(0.5 \text{ mm})$ Station 4, correction from segments Baffield residuals, uncorrected Mean -0.0237 350 RMS 0.5118 alignment residuals, uncorrected 300 alignment residuals 250 200 150 100 50 Mean of residuals in each bin (mm) Bin-by-bin difference (mm)

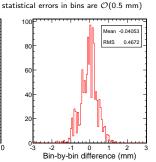
Residuals with new $\vec{B}(\vec{x})$ maps Jim Pivarski 8/17





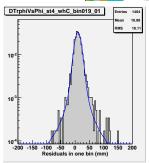
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Station 4, correction from new grid B-field residuals, uncorrected 350 alignment residuals, uncorrected 300 alignment residuals 250 200 150 100 50 Mean of residuals in each bin (mm)





- ► Non-multiple scattering processes have power-law distributions, while experimental resolution is Gaussian
- ► Peak of residuals distribution should not be computed from the mean: it would be pulled by scattered "outliers"
- Model process as Lorentzian-Gaussian convolution:



$$f(x) = \int_{-\infty}^{\infty} \frac{1}{\pi} \frac{\Gamma/2}{(x - \xi - x_0)^2 + (\Gamma/2)^2} \times \frac{1}{\sqrt{2\pi}\sigma} \exp\left(\frac{-\xi^2}{2\sigma^2}\right) d\xi$$

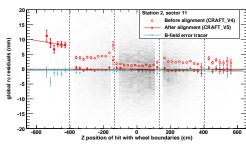
- ightharpoonup Determine peak (alignment correction) from x_0 of unbinned fit
 - regular mean $(\sum x_i/N)$ = center of an unbinned Gaussian fit
 - this is the same thing, but with tails
 - "outliers" contribute far less to f(x) log-likelihood than Gaussian

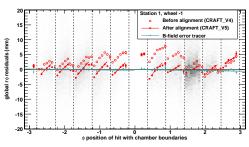
Is it a real alignment?

- ► We want to find the real positions of chambers, not just minimize residuals
- ► To look for biases in the track source, plot residuals more finely than the chamber boundaries
 - bias can change residuals shape inside chambers and across boundaries
 - only misalignments can make discontinuities at chamber boundaries
- ► Cause of linear slopes in $r\phi$ vs. ϕ (bottom) under investigation (DTs stretched in x? tested ϕ_y and z-shift hypotheses...)

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► Complete set of plots: http://indico.cern.ch/conferenceDisplay.py?confId=51267 ("more information")

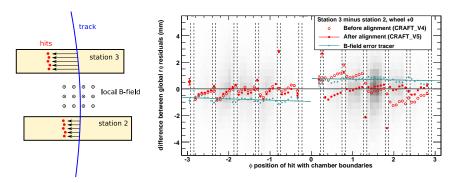
Cross-check: relative positions Jim Pivarski 11/17





- Alignment procedure determines chamber positions relative to tracker
- Chamber positions relative to other chambers is a true cross-check
- Difference of residuals on the same track uses the track as a curved ruler to compare two chambers:

$$difference = (st. 3 track - st. 3 hit) - (st. 2 track - st. 2 hit)$$



Difference distributions are about 4 times narrower than residuals

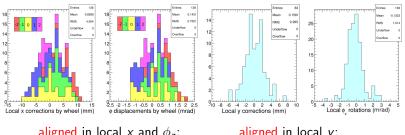
Alignment results

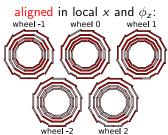
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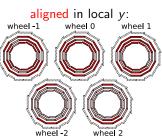




- ▶ The following are alignment corrections used in CRAFT re-processing
 - local x is in the $r\phi$ direction, local y is along the beamline
 - ightharpoonup x re-expressed as ϕ to demonstrate lack of wheel rotations



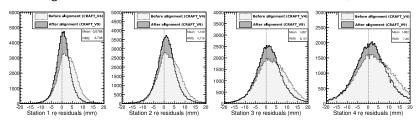




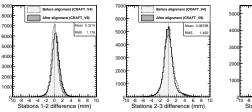
Residuals after global alignment Jim Pivarski 13/17

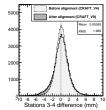


Alignment narrowed and centered residuals distributions, as it must



- ▶ Alignment preserved but didn't improve residuals differences
 - ▶ 1–2 mm *relative* chamber positions before and after alignment





Note smaller scale

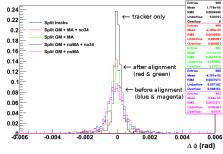
Cosmics track-splitting study

- ➤ Top and bottom half of a cosmic muon should have the same track parameters
- ► GlobalMuon resolution worse than tracker-only for three reasons:
 - 1. global misalignment
 - 2. magnetic field errors
 - tracker given too little weight in global track fit
- ightharpoonup Alignment improves matching of $p_T > 100$ GeV cosmics
 - insensitive to (2)
 - plotted before (3) corrected
- This is another cross-check because top-bottom agreement not used in alignment procedure

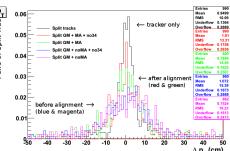
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N. Tran, A. Bonato





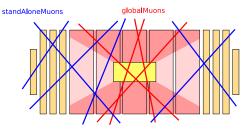


Next steps in alignment

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- 1. Incorporate what we've learned into well-organized alignment package
 - Alignment/MuonAlignmentAlgorithms MuonAlignmentFromReference (CVS tag used for CRAFT alignment: V00-03-01)
 - $\textcolor{red}{\blacktriangleright} \ \texttt{Alignment/CommonAlignmentMonitor} \ \texttt{AlignmentMonitorMuonSystemMap}$
- 2. Solve " $r\phi$ residual vs. ϕ " problem (page 10), if possible
- 2. Re-align accessible chambers of barrel in all 6 degrees of freedom
- Use standAloneMuons to align those DT chambers which are inaccessible to globalMuons
- 4. Use standAloneMuons to align CSCs, using barrel as reference
- 5. Do it all again in CRAFT-2009, but as a push-button procedure

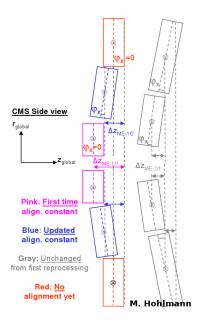


Merging alignment information

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- Straight-line monitors measure bending in endcap disks: z and ϕ_x parameters
 - included in CRAFT alignment
 - can measure x and ϕ_z for some chambers
- ▶ Track-based measurement of x, ϕ_y , and ϕ_z will be more complete and precise
 - ightharpoonup can also measure z and ϕ_x
- ► Combine results parameter-by-parameter:
 - 1. hardware prepares aligned CSCAlignmentRcd, passes it on
 - 2. track-based doesn't update z, $\phi_{\rm x}$
- ► Final result is easier to understand than a combined or constrained fit
- ▶ Redundant measurements (x, z, ϕ_z) reserved for cross-checks



- ▶ First global alignment, connecting muon chambers to the tracker's coordinate system
- Careful to avoid errors due to source of tracks and propagation
- Employed multiple cross-checks:
 - redundant binning shows no distortion inside of chambers
 - aligned in absolute coordinates, checked relative differences
 - track-splitting study showed improvement
- Highly detailed misalignment scenario generated from this information, to be used in CRAFT Monte Carlo production
- Wrapping this all up into a routine procedure
 - to extend alignment into the endcap
 - and be ready for CRAFT-2009