VtxKalman: a decay chain fit for Babar

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Outline:

Part I: decay chain fit

• Part II: Δz -fit

For up-to-date slides: www.slac.stanford.edu/~hulsberg/talks/vtxkalman.pdf

Part I: A decay chain fit

- Fits a complete decay chain 'in one go'
- Input:
 - reconstructed particles (+pid hypothesis)
 - decay tree hypothesis
 - evt: mass constraints, beamspot, beamenergy, lifetimes
- Output:
 - momenta, vertex positions
 - full covariance matrix

Why?

- 'GeoKin' fit only updates parameter of daughters, not of 'granddaughters'
- → parameters not optimal if
 - the decay tree is more than one layer deep (e.g. $B \rightarrow DD$), and
 - the fit is strongly non-linear
 - 'GeoKin' fit does not compute the correlation between daughters
 - only technical problem for 'one-layer'
 - not possible for granddaughters'
 - a global decay chain fit
 - propagates constraints through complete tree
 could improve resolutions in a non-linear fit
 - has all correlations
 - \Longrightarrow could be interesting for $B \to DD$ analyses

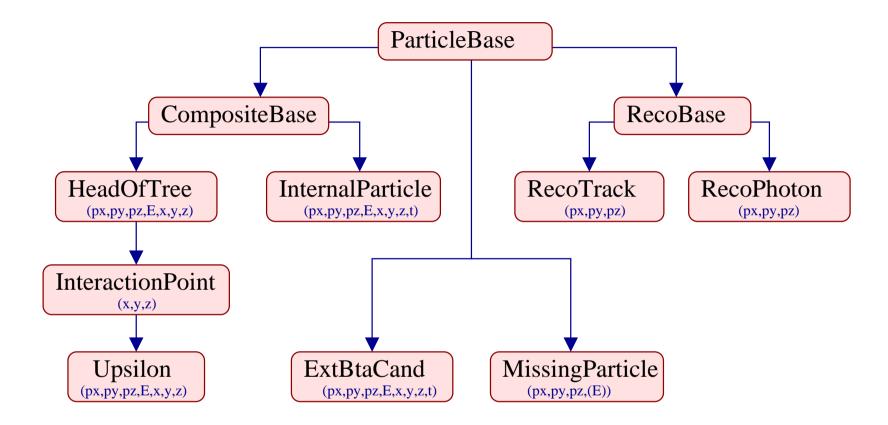
Modeling the decay chain

- decay chain broken down into 'particles'
 - 3-momentum or 4-momentum
 - decay vertex
 - lifetime

type	parameters	constraints
composite	$(x, y, z, \tau, p_x, p_y, p_z, E)$	geo+kin (+mass) (+lifetime)
resonance	(px, p_y, p_z, E)	kin (+mass)
photon	(p_x, p_y, p_z)	$(ec{x}_{ m cl}, E_{ m cl})$
track	(p_x, p_y, p_z)	$(d_0, \phi_0, z_0, \omega, \tan(\lambda))$

- mass constraints of external particles are 'built in'
- internal mass, geometric and kinematic constraints enforced via Lagrange multipliers in the fit

Enter C++



Fitting

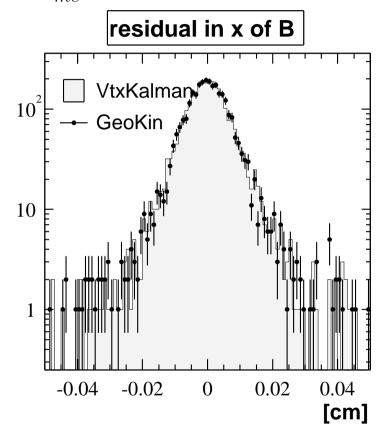
- decay chain model leads to large number of parameters with many correlations
- example: the decay tree of $B^0 \to D^+D^-$ with $D \to K\pi\pi$ has 40 parameters
- hard to solve with a global least χ^2 fit based on Newton-Rapson iteration since it requires inversion of the full covariance matrix
- therefore, use (extended) Kalman fit
 - add constraints one-by-one, updating pars and cov in each step
 - iterate non-linear constraints (*i.e.* mass)
 - iterate complete fit until failure or convergence
- Kalman filtering
 - not new for vertexing (e.g. FastVtx)
 - new for complete decay chain (I think)
 - new for exact ('Lagrange') constraints (I think)

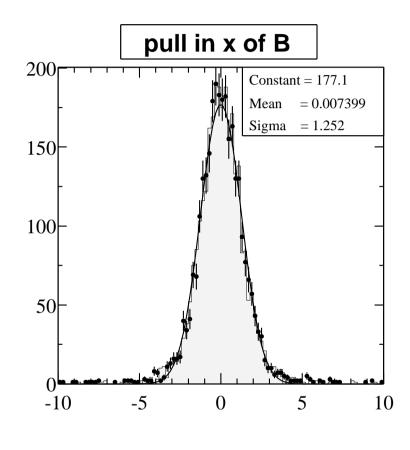
Some differences wrt GeoKin

- GeoKin for photons:
 - take cluster position and estimated origin
 - translate into four-vector (p_x, p_y, p_z, E)
 - use four vector in vertex fit clearly doesn't work for $K_s \to \pi^0 \pi^0$
- use cluster position and energy instead of 'constructed' four-vector
 - same for charged tracks: helix parameters instead of four-vector
 - in GeoKin the 'beamspot' is an xy-constraint on decay vertex
 - doesn't makes sense for B, or B decay products
 - GeoKin uses special 'beamspot-B-flight' assigns uncertainty in y of 30 μ m
 - very non-gaussian
- ⇒ define as constraint on *production* vertex ...

Comparison of position resolution

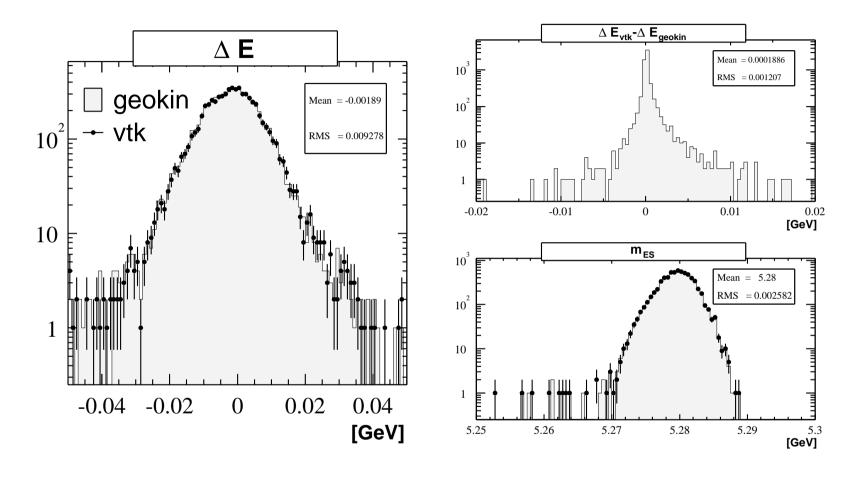
• $x - x_{mc}$ of B^0 in $B^0 \to D^+D^-$





- no difference in overall parameter resolution
- no difference in pulls

Comparison of ΔE resolution



- event-by-event difference of the order of 0.2σ
- no difference in overall parameter resolution
- no difference in pulls

Why no improvement?

- 1. is the problem too linear to see a difference?
- 2. is the Extended Kalman Filter non suitable for propagating the non-linear constraints all through the tree?
 - EKF only inherits the parameters of the previous iteration, not its covariance
 - I am thinking of an alternative . . .

CPU time consumption

The disclaimer

- calculating correlations takes time
- ullet CPU consumption scales roughly with n^2
 - expect VtxKalman to be slow for large decay trees
- time consumption predominantly in matrix manipulation
- current limitation: accessing data in HepSymMatrix, HepMatrix

CPU (2)

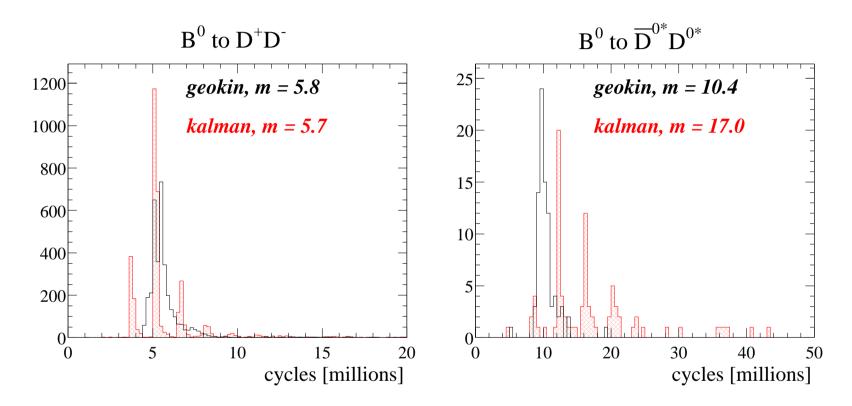
Measured with 'clock()', on a desktop

- a single-level decay chain $D^+ \to K\pi\pi$)
 - 16 parameters
 - factor ~ 1.5 faster than GeoKin (0.67 ms vs 1.07 ms)
- a 2-level decay chain $B^0 \to D^+(\to K\pi\pi)D^-(\to K\pi\pi)$
 - 41 parameters
 - 2 mass constraints
 - equally fast as GeoKin
- a multi-level decay chain $B^0 \to D^{*0} \bar{D^{*0}}$ with $D^{*0} \to \pi^0(\gamma\gamma) D^0(K\pi)$
 - 63 parameters
 - 6 mass constraints
 - factor 2 slower than GeoKin

Note: VtxKalman *can* fit any tree in layers *for the same information, it is always faster*

CPU (3)

CPU time ('Gerhard's cycles') on MC



VtxKalman timing proportional to number of iterations

Fitting $K_S \to \pi^0 \pi^0$

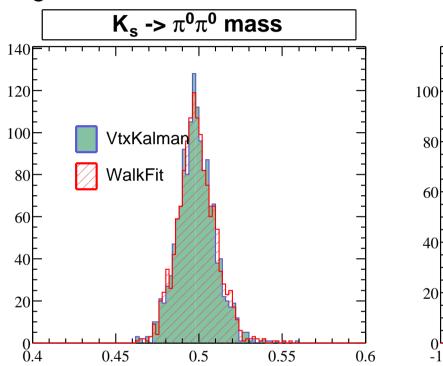
- GeoKin uses separate fitter for $K_S \to \pi^0 \pi^0$:
 - uses clusters (like VtxKalman)
 - needs direction constraint for Ks (not optimal)
 due to missing implementation for production vertex constraint
 - not completely finished (and doesn't work now)
- default algorithm is 'VtxKsToPi0Pi0WalkFit'
 - uses production vertex constraint
 - very slow
- VtxKalman can fit this decay, given
 - mass constraints on the π^0 s
 - production vertex

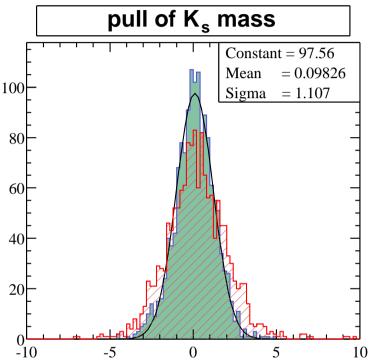
$K_S \rightarrow \pi^0 \pi^0$: VtxKalman versus 'WalkFit'

- $K_S \to \pi^0 \pi^0$ from $B^0 \to J/\psi K_S$ \Longrightarrow relatively hard
- K_S tree build using MC truth \Longrightarrow no composition sequence involved
- ullet photon energy larger than 20 MeV and within 4 σ from MC truth
- use beamspot for production vertex
 B decaylength irrelevant here
- use π^0 mass constraints
- fit has 2 degrees of freedom

$K_S \to \pi^0 \pi^0$: Comparing mass resolution

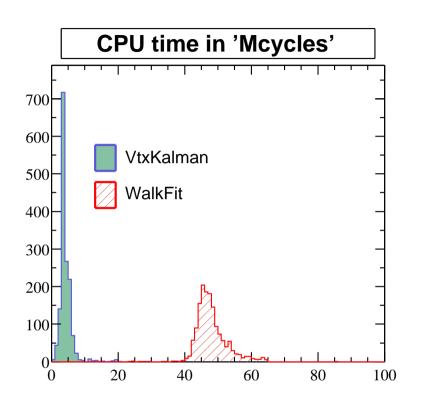
For *good* fits:

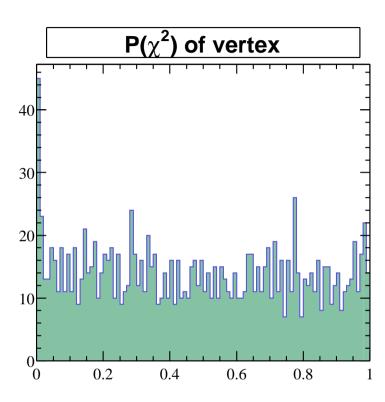




- comparable resolution
- VtxKalman has much better pull

$K_S \to \pi^0 \pi^0$: CPU time, χ^2

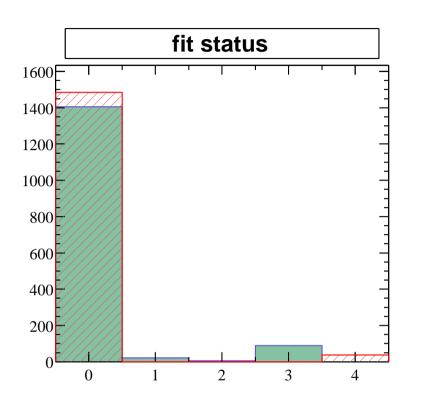


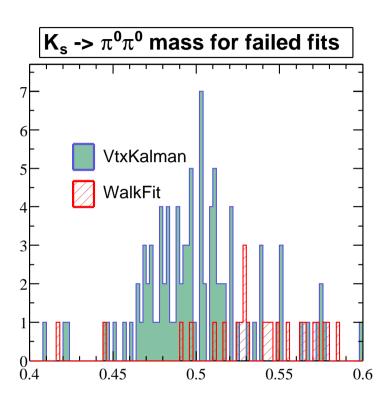


- both fits are slow
- VtxKalman approximately 10 times faster (~ 2 ms vs ~ 20 ms)
- VtxKalman χ^2 distribution looks 'perfect' (no χ^2 from the walk fit)
- note: since (almost) all particles from beam spot:, can interprete χ^2 as due to π^0 mass constraints

$$\Longrightarrow$$
 bad χ^2 means ≥ 1 bad π^0

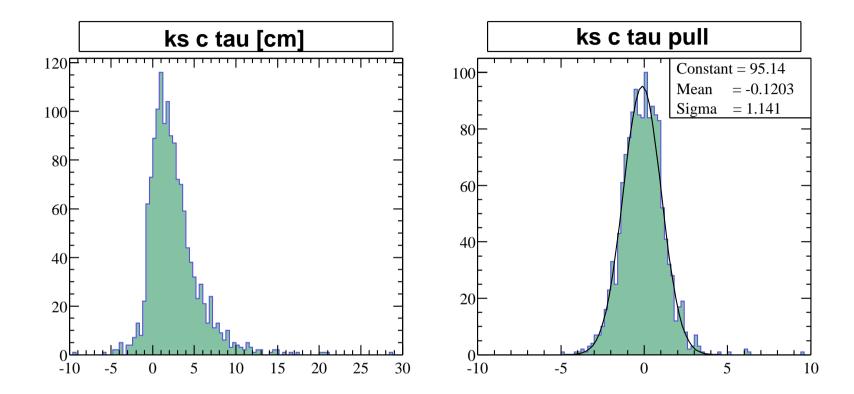
$K_S \to \pi^0 \pi^0$: Failed fits





- VtxKalman fit 'fails' a bit more often
- much worse resolution for failed fits, but still signal
 must find what that is
- (however, on the same events, WalkFit mass is *just as bad*)

$K_S \to \pi^0 \pi^0$: decay length



- ullet c au distribution looks good
- → could use for background suppression

Problem 1: the order of constraints

- in a kalman fit constraints are applied in a specific order
- in non-linear fits the result is sensitive to this order
- example: $X \to Y + Z + \pi^0$ with a mass constraint π^0 \Longrightarrow geometric constraint on π^0 must be applied *before* mass constraint, otherwise the π^0 'walks away'
- order of constraints
 - all independent 'external' constraints (tracks, photons, beamspot, fitted composites)
 - kinematic constraints (four-vector addition)
 - geometric constraints (mother-to-daughter)
 - mass constraints

scheme is not yet completely satisfactory

Problem 2: initialization

- kalman fits very sensitive to initialization (both pars and cov)
- parameters are initialized using TrkPoca, four vector addition, pdt lifetimes etc
 - it was hard to come up with a scheme that always works
 the 'initialization' routine is the most complicated and fuzzy part of the package
- covariance matrix initialition is basically result of fine-tuning
- I keep hitting decays for which things do not work out well will this converge?

Problem 3: machine accuracy

 the expression for the updated covariance matrix in the kalman fit is

$$C_k = (1 - K_k H_k) C_{k-1} (1 - K_k H_k)^T + K_k V_k K_k^T$$
 (1)

identical to the simpler expression

$$C_k = (1 - H_k K_k) C_{k-1} (2)$$

⇒ much faster, but . . .

- the latter is known to be more sensitive to machine accuracy . . .
- first time I observe sensitivity to this effect
- managed to code expression (1) such that it is 'only' factor 2 slower than (2)

Problem 4: interfacing to Beta

Main problems:

- 1. some information cannot be stored in BtaFitParams
 - correlation between daughters
 - lifetimes
 - only accessible if explicitely calling 'vtxkalman::VtkVertex'
- 2. fitting not independent of the way the tree is built example: $B^0 \to D^{*+}\pi^-$
- 3. current behaviour of vertexers wro updating daughter parameters is confusing

Problem 4c: updated daughters

- by default, copying a BtaCandidate yields a shallow copy
 - the underlying data structure, the BtaCandBase, is not copied
 - decay trees share daughter parameters
- currently, vertexers make only shallow copies
 - only update the top layer of the decay tree
 - never update daughter parameters
 - clearly, at odds with the idea of a decay tree fit
- to access 'daughter' parameters the daughter must be changed!
 - currently steered by 'fitAll' flag (but users do not understand this)
 - this daughter is changed for ever
- it is a nightmare that the output could depend on the order in which the composition sequences are called

Summary I: pros/cons of new fit

- + correlations in complete decay tree
- + consistent treatment of mother constraints (e.g. for $B^0 \to D^*X$)
- + photons treated as ecal clusters (e.g. for $K_S \to \pi^0 \pi^0$)
- + tracks treated as helices
- + beam spot constraint on production vertex
- + missing particles, lifetime constraints
- + applicable to practically all decay chains (can get rid of several dedicated fitters)
- no kinematic fit (but do we really need that?)
- kalman filter still 'fails' more often than global fit
- decay chain fitting slightly at odds with 'fitting layers'

Summary II: status

- implementation roughly complete (5863 lines), first version in babar-cvs (VtxKalman)
- parameter pulls look fine, resolutions equivalent to GeoKin
- interfaced via VtxFitterOper (not in cvs)
- implemented 'Fast' mode and 'SingleTrack' mode; however, purely kinematic fitting not supported
- tested as replacement of GeoKin in the CompositionFactory
 - tested on different MC samples
 - tested on B-flavor sample (UpsilonQA)
 - small loss in yield ($\sim 2\%$) . . .
 - both background and signal → probably failed fits
- a write-up of the Kalman-math is in progress: www.slac.stanford.edu/~hulsberg/notes/vtxkalman.ps

Summary III: plans

- try to recover remaining loss of signal events on B-flavor sample
- come up with a safe solution for 'updated daughter parameters'
- → needs some redesign in VtxBase/VtxFitter

• . . .

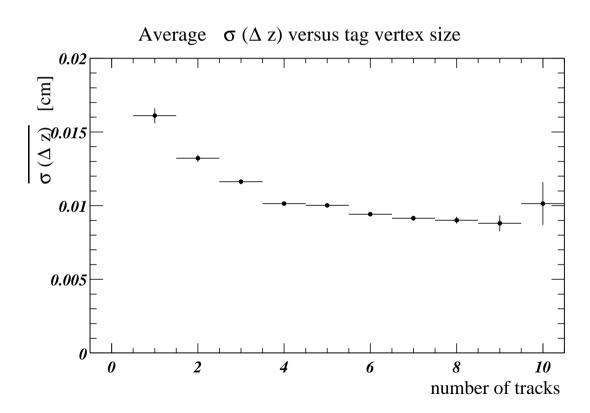
Part II: Δz Fitting

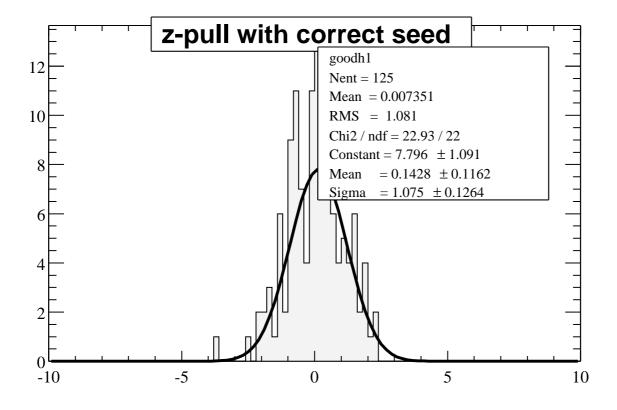
- common recipe:
 - remove tracks from reco-B (or B-CP)
 - estimate trajectory of tag-B from reco-B and beamspot/energy constraint
 - reconstruct vertex for tag-B from remaining tracks and the seed
- in GeoKin:
 - \bullet combine all tracks into one vertex (after removing conversions and substituting $K_S \text{tracks}$ with mother)
 - remove incompatible tracks \implies tracks with χ^2 contribution > $\chi^2_{\rm max}$
- alternative scheme:
 - start with a good seed
 - add compatible tracks \implies tracks with $\Delta \chi^2 < \chi^2_{\rm max}$

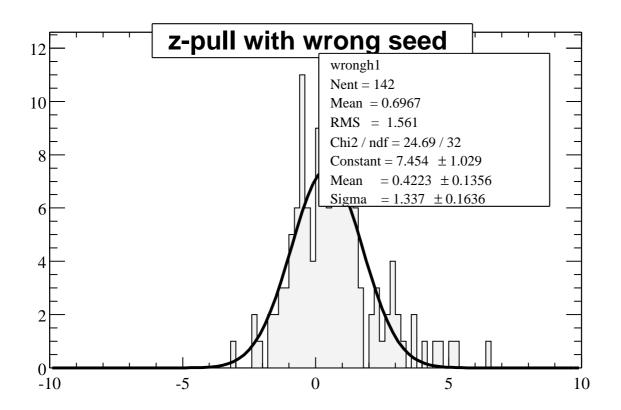
Tag vertex reconstruction

Obvious conflict:

- the more tracks, the smaller the statistical uncertainty
- the more tracks, the larger the bias due to D daughters







Seed how-to

It is important to choose a good seed, but how?

Looked at several observables

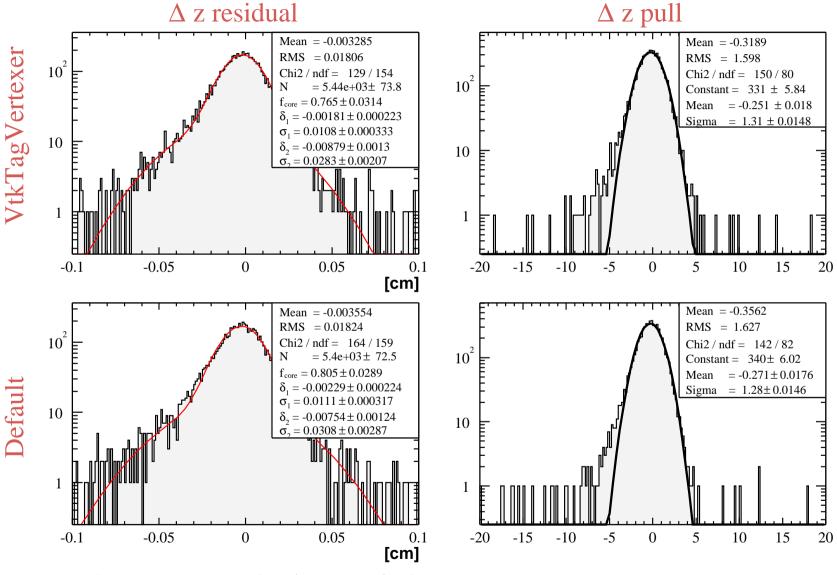
- $\chi^2 \Longrightarrow$ no good: slow tracks have large errors
- $p_{\perp} \Longrightarrow$ no good: *most* B do not yield 'one' high p_{\perp} track
- best choice: likelihood 'density'

$$-2\log \mathcal{L} = \chi^2 + \log[\det(V)]$$

where V is the covariance of the 'pseudo-B + seed-track' vertex

• (in fact, using σ_z^2 instead of full V makes little difference)

Results: Δz resolution



- ullet no improvement in Δz resolution
- maybe, very small hint for reduced bias

Conclusions

- there exist an alternative vertex fit
 - full covariance matrix of a decay chain fit
 - better model of photons
 - no improvement in parameter resolution
 - small improvement in CPU time consumption
- there exists an alternative Δz fit
 - slightly more 'advanced' choice of tag vertex tracks
 - no improvement in Δz resolution
 - ullet possibly, tiny reduction of Δz bias
- is it worth pursuing either of these projects?

Deep copies and refitted daughters (II)

Two solutions

- 1. only deep copies in decay trees
 - safe
 - not useful for vertexers that do not calculate updated daughters
 - severe penalty in memory consumption/CPU
 I would say that this is not a viable solution
- 2. shallow copies in decay trees, but access to daughter parameters only via deep copies
 - requires only small modifications in current design