



MAX-PLANCK-GESELLSCHAFT



Optimization of the B meson vertex resolution for the Belle II experiment

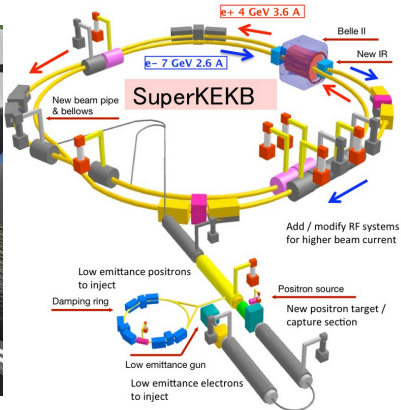
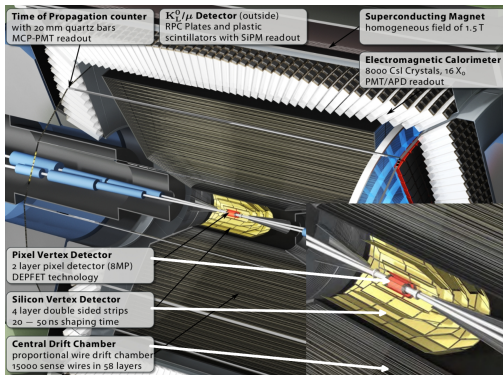
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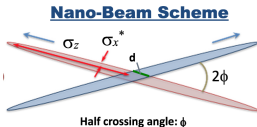
IMPRS YS Workshop , Jul 6-10, 2015

Belle II experiment at SuperKEKB



New Tracking System

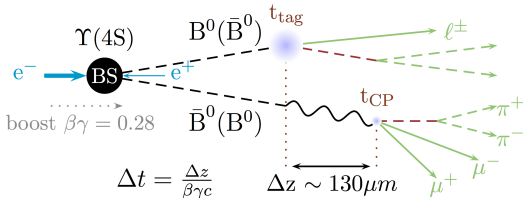
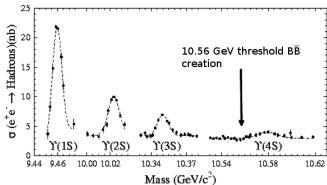
- Pixel Vertex Detector (PXD)
- Silicon Vertex Detector (SVD)
- Central Drift Chamber (CDC)



Instantaneous luminosity

- KEBK:
 $2 \times 10^{34} \text{ cm}^{-2} \text{ s}^{-1}$
- SuperKEKB:
 $8 \times 10^{35} \text{ cm}^{-2} \text{ s}^{-1}$

CP violation and time dependent analysis



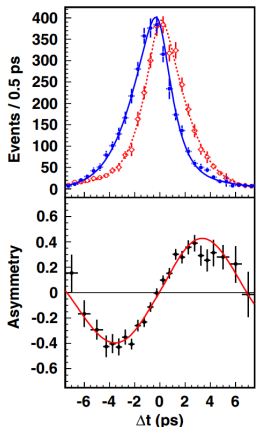
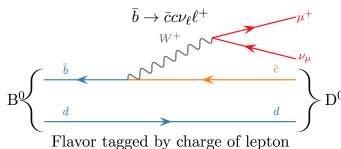
$$\Upsilon(4s) \rightarrow B^0\bar{B}^0 \rightarrow f_{CP}t_{\text{tag}} ; f_{CP} \rightarrow J/\psi K_S, t_{\text{tag}} \rightarrow \text{all modes}$$

Asymmetry in the B/\bar{B} decay to f_{CP}

$$A_{CP} = \frac{\Gamma(\bar{B}^0 \rightarrow f_{CP}, \Delta t) - \Gamma(B^0 \rightarrow f_{CP}, \Delta t)}{\Gamma(\bar{B}^0 \rightarrow f_{CP}, \Delta t) + \Gamma(B^0 \rightarrow f_{CP}, \Delta t)} = A_{CP} \cos(\Delta m_d \Delta t) + S_{CP} \sin(\Delta m_d \Delta t)$$

● A_{CP} : direct C/P ; S_{CP} : mixing induced C/P

- Kinetic reconstruction \rightarrow CP side
- Flavour ID \rightarrow tag side



* PRL 108, 171802 (2012)

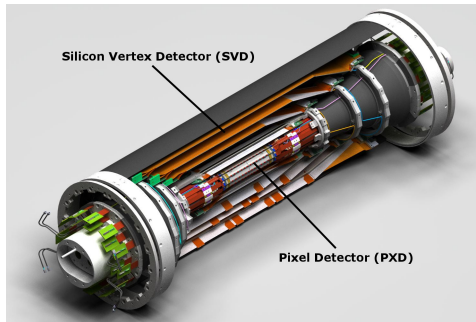
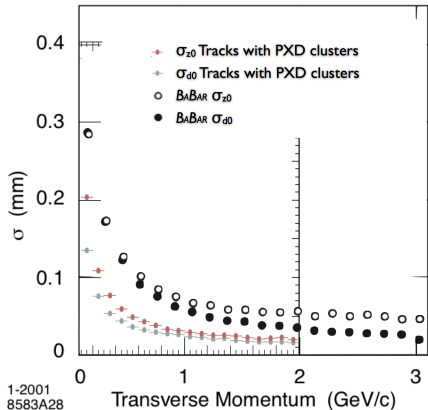
Christian Roca Catalá (MPP-Belle2)

IMPRS YS Workshop

Jul 10, 2015

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Pixel Vertex Detector - Importance of good resolution



What is the contribution of the Vertex resolution?

- **Time-dependent analysis** depend on the good measurement of Δt - obtained from Δz
- Belle: B_{tag} resolution ($89 \mu\text{m}$) $\sim 30\%$ **worse** than B_{CP} resolution ($63 \mu\text{m}$)
- B_{tag} **resolution** depends heavily on the fitting **algorithm**.
- The **contribution** for Δz that needs to be specially **optimized** is the B_{tag} **vertex**

$$B^0(\bar{B}^0) \rightarrow f_{CP} = [J/\psi \rightarrow \mu^+ \mu^-] K_S$$

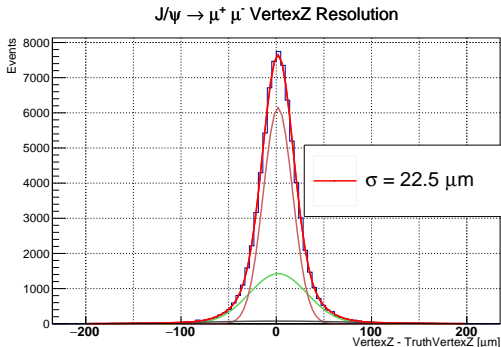
Legend

- Resolution distr. = vertex - MC vertex
- **3 Gaussian fit** with std. dev σ
- Resolution = weighted avg. σ

Observations

- μ tracks ≥ 1 PXD hit
- Resolution **improvement** from Belle ($63 \mu\text{m}$) - **factor 2.7**
- Small **Shift** of $1.8 \mu\text{m}$ (under investigation)

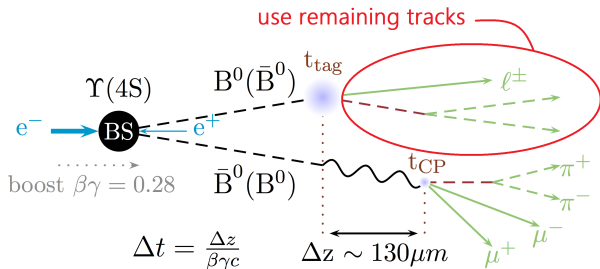
Belle II simulation using PXD



How is the fit performed?

- 1 B_{tag} vertex fit uses the **tracks remaining** after the reconstruction of the CP side
- 2 **No** B_{tag} reconstruction is performed (**loss of statistics**)
- 3 Algorithm: **RAVE** Adaptive Vertex Fit (AVF)* with spatial constraints

* : W. Waltenberger, R. Frühwirth and P. Vanlaer: **Adaptive Vertex Fitting**.
CERN-CMS-NOTE-2008-033 (Jul 2008)



Standard fitting algorithm and spatial constraint

RAVE Adaptive Vertex Fit (AVF) with constraints

All tracks are used and **weighted** following two criteria:

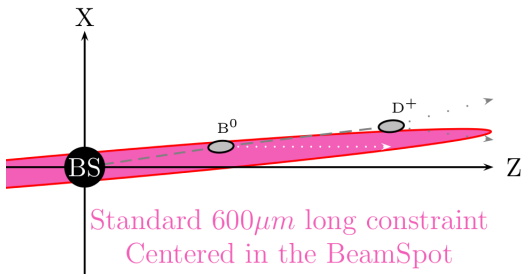
- 1 **Outlying** and isolated tracks are **down-weighted**
- 2 Tracks weighted according to their **position** with respect to the **constraint**

Weighting works **iteratively**

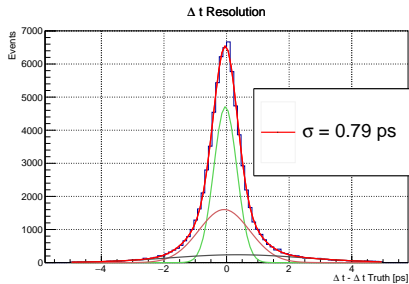
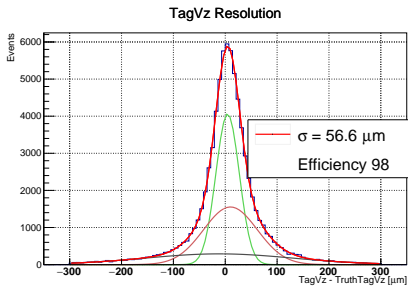
What is the constraint?

A **spatial constraint** is defined within which the B is expected to decay

- **Ellipsoid** of $600\ \mu\text{m}$ long axis
- Centered in the **Beam Spot**
- Along the **boost direction**



Belle II B_{tag} vertex Resolution - Standard algorithm



In comparison with BELLE:

- Resolution = $89 \mu\text{m}$ - factor 1.5

In comparison with BELLE:

- Resolution = 0.92 ps - factor 1.2
- Lower factor \rightarrow Lower Boost!

Improvement in the B_{tag} vertex resolution lower than for B_{CP}

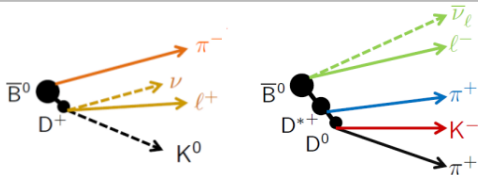
Dominated by the algorithm! PXD precision not fully used.

Can we do better than this? New algorithm

New Algorithm: Using Flavor Tagging Information

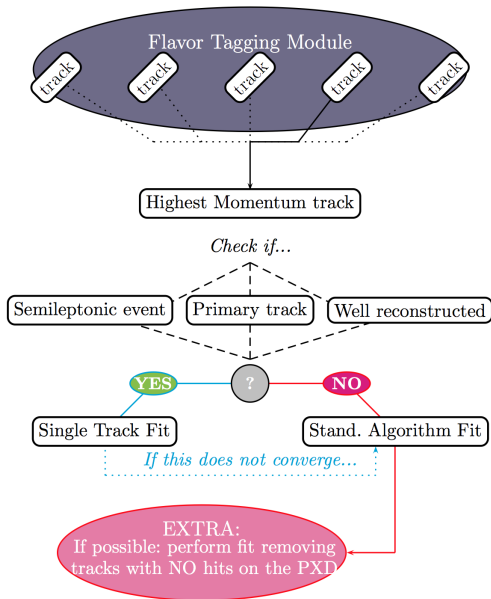
Flavor Tagging Algorithm:

- Take all the **remaining** tracks after the reconstruction of the **CP side**.
- Find the tracks best suited for **Flavor Tagging**
- Extract **probabilistic** information from those tracks
 - Prob.** of being a **daughter** from B_{tag}
 - Prob.** of belonging to a given **decay mode**/category
- Finally return a **parameter** $\in [-1, 1]$ that reflects how good the **flavor** can be identified
- Flavor id. crucial to measure CP**

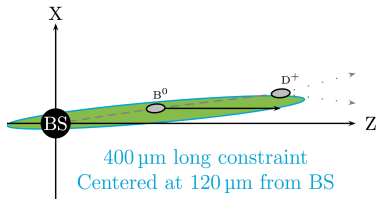


Categories	Targets
Electron	e^-
(Intermediate Electron)	e^+
Muon	μ^-
(Intermediate Muon)	μ^+
KinLepton	e^-
Kaon	K^-
KaonPion	K^-, π^+
SlowPion	π^+
FastPion	π^-
MaximumP	ℓ^-, π^-
FSC	ℓ^-, π^+
Lambda	Λ
Total= 10 (12)	

New Algorithm: Single Track Fit (STF)



Constraint of New Algorithm



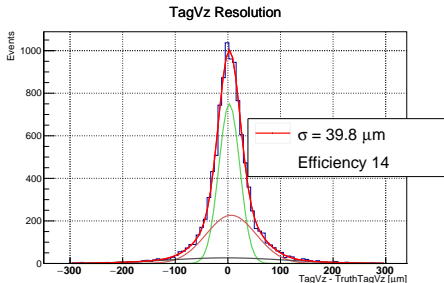
Why do we move the constraint?

Secondary tracks are **not** used in the fit, and therefore:

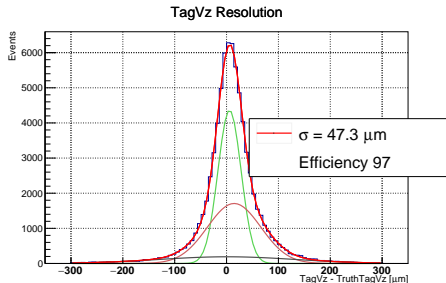
- **Constraint** centred at the BS is no longer down weighting secondary tracks
- B-meson decays in average $\sim 120 \mu\text{m}$
- If the **constraint** remains centred at BS, **large bias** appears ($\sim 15 \mu\text{m}$)

New algorithm: B_{tag} vertex resolution

Events with successful single track fit



All events



In comparison with Belle:

Resolution = 89 μm

- Single Track Fit **Factor 2.2**
- STF + Standard Alg. **Factor 1.9**

In comparison with Standard Algorithm:

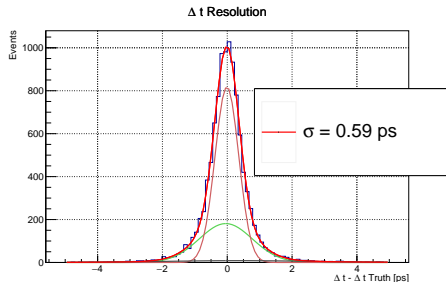
Resolution = 56 μm

- Single Track Fit **Factor 1.4**
- STF+ Standard Alg. **Factor 1.2**

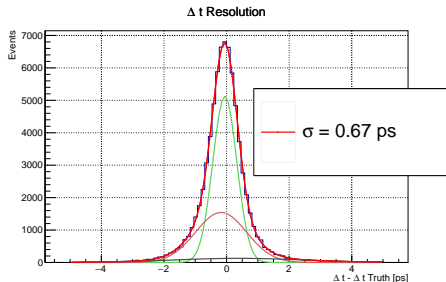
- Single Track Fit **triggered** $\sim 15\%$ of all B_{tag} cases
- Resolution **improvement** with respect to Belle almost reached **CP side**.
 - **CP side** - factor 2.8
 - **Tag side** - factor 2.2 (Single Track Fit) , factor 1.9 (Combined)

New algorithm: Δt distribution

Events with successful single track fit



All events



In comparison with Belle:

Resolution = 0.92 ps

- Single Track Fit **Factor 1.6**
- SFT + Standard Alg. **Factor 1.4**

In comparison with Stand. alg.:

Resolution = 0.78 ps

- Single Track Fit **Factor 1.3**
- SFT + Standard Alg. **Factor 1.2**

- Single Track Fit **triggered** $\sim 15\%$ of all B_{tag} cases
- Resolution **improvement** with respect to Belle almost reached **CP side**.
 - **CP side** - factor 2.8
 - **Tag side** - factor 2.2 (Single Track Fit) , factor 1.9 (Combined)

Vertex resolution and optimization using PXD

- **B_{CP} Vertex resolution** (23 μm): improved a **factor 2.7** with respect to Belle (63 μm).
- **B_{tag} Vertex resolution with Standard Algorithm** (57 μm): improved a **factor 1.5** with respect to Belle (89 μm).
- Improvement in **Tagged side** do not scale as in **Reconstructed side**
- **New algorithm** performs a **single track fit** with higher resolution (39 μm) with 15% efficiency, improvement by a **factor 2.2** with respect to Belle.
- **B_{tag} Vertex resolution with Standard Algorithm + Single Track Fit** (47 μm): improved a **factor 1.9** with respect to Belle (89 μm).

Outlook

- Understand the small **shift** on the **CP side** vertex
- Improve the **single track selection criteria** in order to increase the efficiency.
- **Improve** the B_{tag} vertex fit **constraint's** parameters to reduce the **bias** in the **resolution**

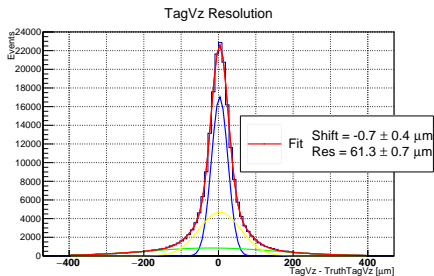
THANKS FOR YOUR ATTENTION!



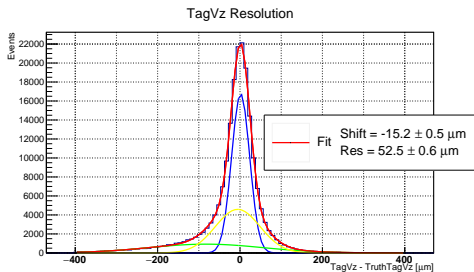
$$B_{tag} \rightarrow \mu^- \bar{\nu}_\mu D^{(*)+}$$

and conjugate

USING ALL TRACKS



USING MC MATCHED MUONS



$$B_{tag} \rightarrow \text{generic}$$

Purity analysis

- Perform several **cuts** on the variables of the **tracks**
- Compare with **Monte Carlo** information
- Aim: Kill the **bad** ones and keep the **good** ones!
- **High purity** acquainted after selection

Normalized distribution of Purities

