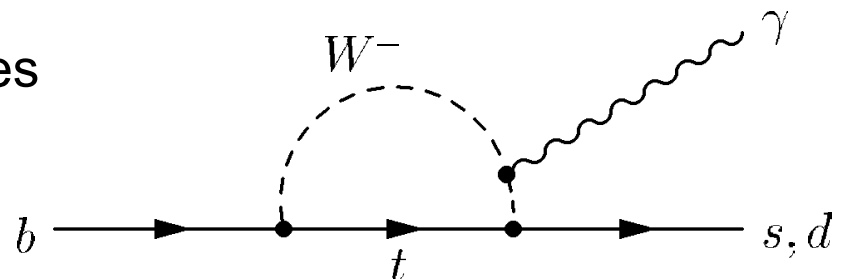


***Inclusive BR($b \rightarrow s \gamma$)
with a fully
reconstructed B***

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Why study $b \rightarrow s\gamma$?

- Possibility of new non-SM particles entering in the loop.



- We are unable to measure the parton level decay rate for $b \rightarrow s\gamma$, however:

$$\text{HQET} \Rightarrow \Gamma(B \rightarrow X_s \gamma) = \Gamma(b \rightarrow s \gamma) + \Delta^{\text{nonpert}}$$

- Next to leading order calculations for $\text{BR}(B \rightarrow X_s \gamma)$:

$(3.29 \pm 0.33) \cdot 10^{-4}$	Kagan&Neubert
$(3.60 \pm 0.30) \cdot 10^{-4}$	Gambino&Misiak

- Theoretical uncertainty $\sim 10\%$, mainly from contribution of higher order diagrams in the expansion.

The signal model

We select signal photons and measure the integral of the **photon energy spectrum** to determine $\text{BR}(B \rightarrow X_s \gamma)$.

Due to the strong force the b quark fragments into B meson, and X_s into a spectrum of K^* resonances.

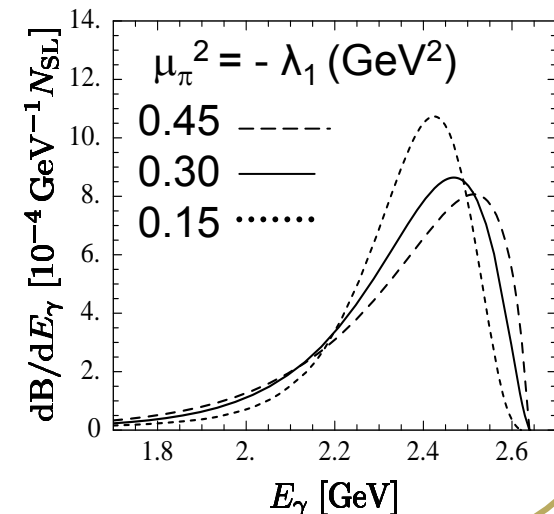
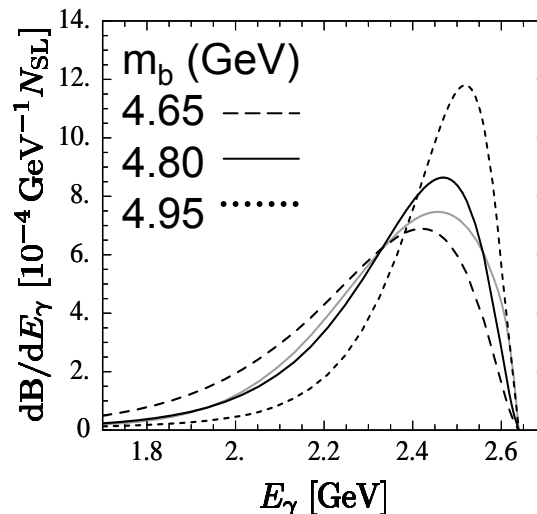
The b quark interacts with the spectator quark \Rightarrow **scale dependence in m_b** .
It also has **Fermi momentum p_F** from its confinement in the meson.

The shape function depends on m_b and p_F .

Spectrum is smeared with

$$\langle E_\gamma \rangle \approx m_b / 2$$

The width (first moment, λ_1) depends on p_F



Advantages of a reconstructed B

Breco (Brecoil) kinematics are well known.

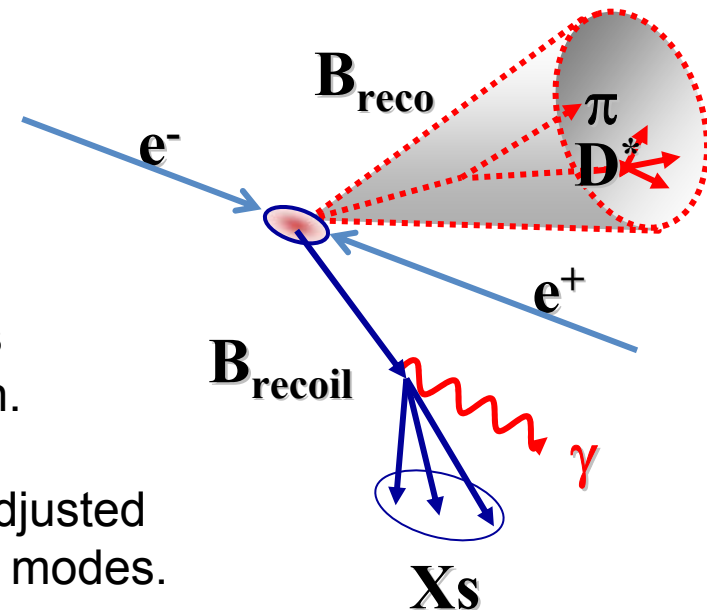
E_γ spectrum is measured in the B rest frame.

Luminosities of samples and B reconstruction efficiencies are not needed, as normalization is taken from # reconstructed B's before selection.

The purity of the Breco sample can be easily adjusted by selecting a sub-sample of the reconstructed modes.

Continuum events can be estimated and subtracted by performing a fit to M_{ES} .

Fully hadronic reconstruction of one B determines tagging of charge and B flavour.
⇒ We can measure $BR(B \rightarrow Xs\gamma)$ and α_{CP} in B^0 and $ChgB$ separately.



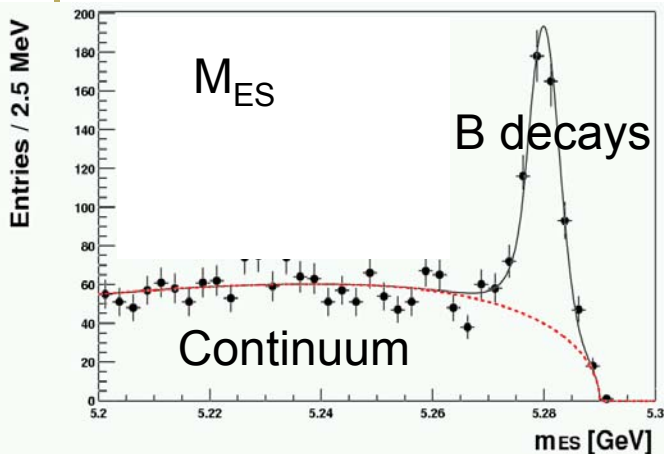
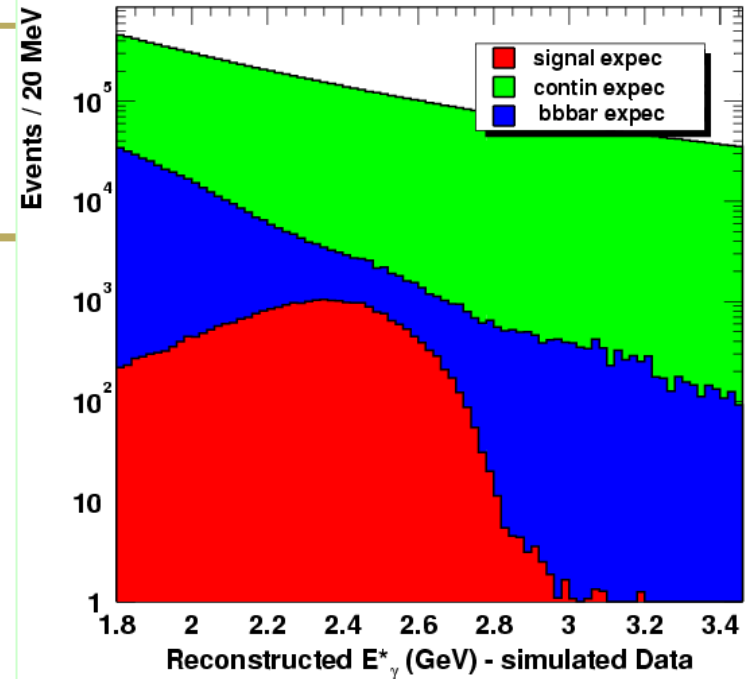
Disadvantage: Small B reconstruction efficiency $\sim 0.4\%$

Analysis concept

Signal: A fully reconstructed B decay &
A high energetic photon in the event.

Backgrounds: from other B decays &
continuum background
($u\bar{u}, d\bar{d}, s\bar{s}, c\bar{c}$)

- Selection criteria and a Fisher Discriminant are designed to suppress background.
- Continuum is estimated from a fit to M_{ES} of the Breco and subtracted.

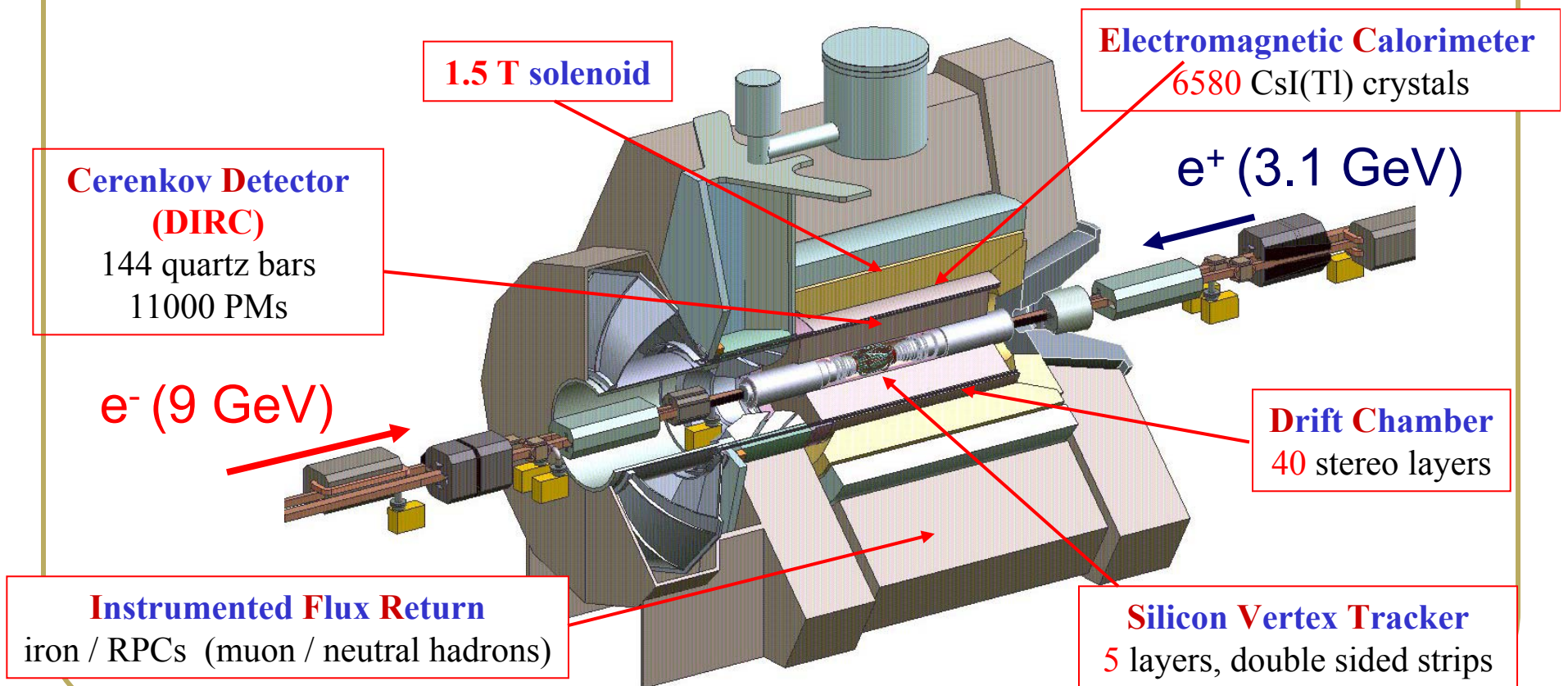


$$M_{ES} = \sqrt{(\sqrt{s} / 2)^2 - P_B^{*2}}$$

- The # BBbar background events in the same peaking component as signal are estimated from MC
- The signal yield is extracted from a binned χ^2 fit to E_γ distribution of events that pass the selection.

The Data

82 fb-1 of data (1999-2002) collected at the BaBar detector



Selection criteria

We require: 1 fully reconstructed B decay (Breco)

Summary of selection Criteria	
B_{reco} candidate	int-pur > 0.25
Minimum photon energy	$E_\gamma > 1.3 \text{ GeV}$
Quality cut	LAT < 0.45
Bump isolation cut	Bump separation > 40cm
π^0 veto	$115 \text{ MeV}/c^2 < M_{\gamma\gamma} < 155 \text{ MeV}/c^2$
η veto	$508 \text{ MeV}/c^2 < M_{\gamma\gamma} < 588 \text{ MeV}/c^2$
Fisher	$F > 0.2$
ρ veto	$620 \text{ MeV}/c^2 < M(\pi^0, \pi) < 920 \text{ MeV}/c^2$

70% of the π^0 background comes from $B \rightarrow D^* \rho$, $\rho \rightarrow \pi^0 \pi^+$, $\pi^0 \rightarrow \gamma \gamma$

For multiple candidates, we select the most energetic photon
(only 0.2% with $E_\gamma > 1.9 \text{ GeV}$)

Fisher Discriminant

(A linear discriminant technique taking care of correlations between variables).

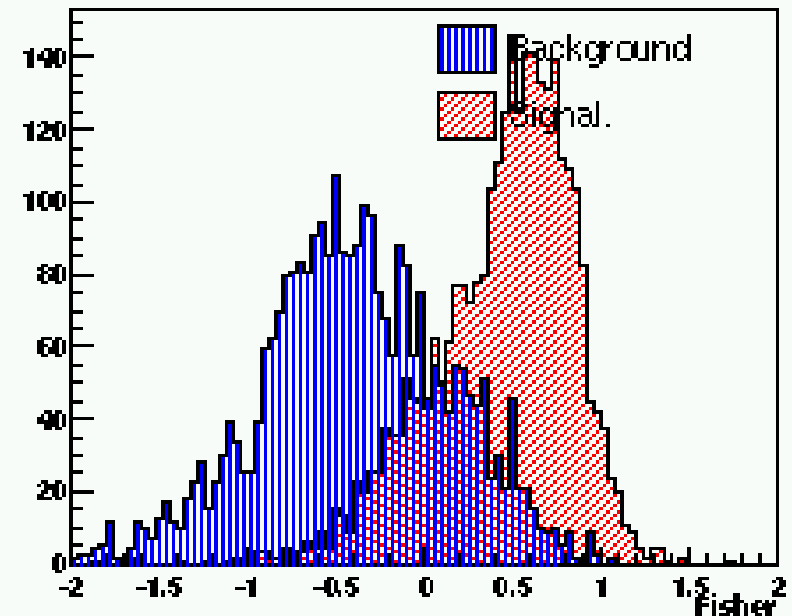
We use event topology to discriminate between signal and continuum background.

- Continuum events are jet like
- $B\bar{B}$ events (incl. signal) are isotropic

28 Fisher Variables:

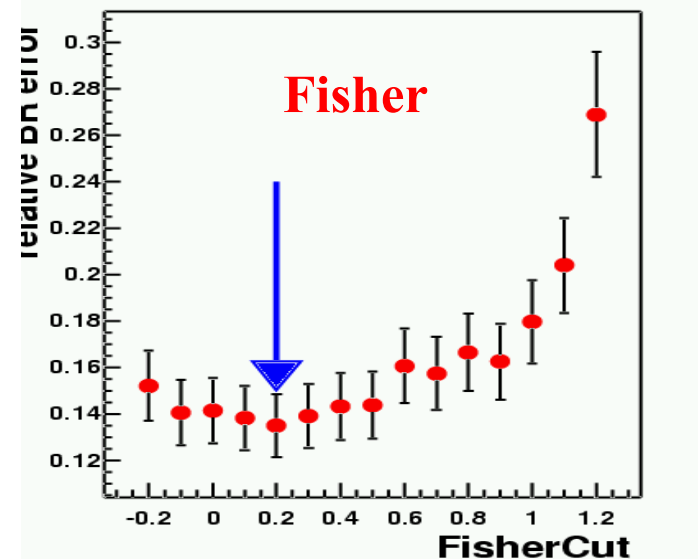
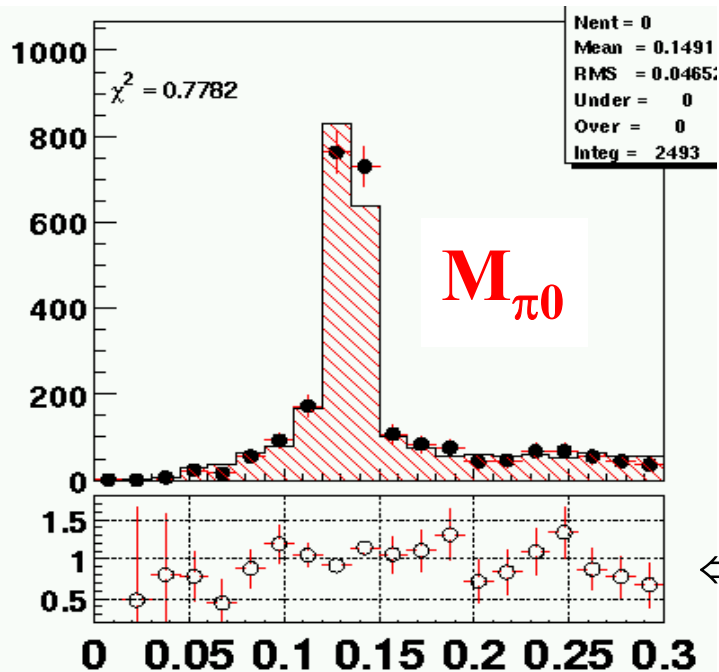
CosTBB, CosGamTrecos(il),
R2,R2Neu, Thrust,ThrustNeu,
ThrustBreco, Energy cones

linear correlation with $E_\gamma < 20\%$,
and with charged and neutral multiplicity
of the Xs $< 15\%$.



The variables

Selection criteria are optimized by varying the cut and minimizing the relative branching fraction error, using MC samples.



Selection efficiencies:

Signal 40%

BBbar 0.02%

← Good Data – MC agreement for variables

Extraction of $BR(B \rightarrow X_s \gamma)$

The branching ratio is extracted using the E_γ distribution

$$Br(B \rightarrow X_s \gamma) = \frac{N_{TRUE}^{sig}}{N_{TRUE}^B} = \frac{N^{sig}}{N^{Breco} \cdot \epsilon^{sel}} \cdot \frac{\epsilon_{tag}^{all}}{\epsilon_{tag}^{sig}}$$

N^{Sig} is the signal contribution to be extracted for $E_\gamma > 1.9 \text{ GeV} \rightarrow \text{fit}$

N^{Breco} is the number of Reco B before any cut $\rightarrow m_{ES}$ fit on data

ϵ^{sel} is the selection efficiency for signal events

$\epsilon^{All(sig)Tag}$ is the Breco tagging efficiency in a generic(sig) } $\rightarrow \text{MC}$

decay.

Extraction of Nsig

Samples are divided into bins of E_γ .

For each bin, the M_{ES} distribution of the Breco is fitted with Argus & Crystal Ball functions and the combinatorial background is subtracted.

N^{Sig} extracted from a binned χ^2 fit:

$$\chi^2(C_s, C_b) = \sum_{i=0}^{\#bins} \left(\frac{N_i^{meas} - C_s N_i^{MC(b \rightarrow s\gamma)} - C_b N_i^{MC(bkgd)}}{\sqrt{\delta N_i^{meas^2} + \delta N_i^{MC^2}}} \right)^2$$

C_s and C_b are the normalizations of Sig and Bkgd components (free parameters)

The last bin contains events with [E \$\gamma\$ >1.9 GeV](#)

$$N_{last}^{sig} = N_{last}^{meas} - C_b \cdot N_{last}^{MC(bkgd)}$$

$b \rightarrow d\gamma$

The $B \rightarrow X_d\gamma$ component is subtracted.

According to the SM expectation $BR(B \rightarrow X_d\gamma)$ and $BR(B \rightarrow X_s\gamma)$ are in the ratio $|V_{td}/V_{ts}|^2$ assuming same efficiency for both components.

BR reduced by $(4.0 \pm 1.6)\%$

Fit Validation

Generic BBbar MC is used instead of data ($\sim 240 \text{ fb}^{-1}$).

Result : $Br = (3.16 \pm 0.43(\text{stat})) \cdot 10^{-4}$

in **good agreement** with the input generator value ($Br = 3.29 \cdot 10^{-4}$).

A check on data was performed using a Pi^0 control sample

(π^0 Veto inverted & no Bump isolation cut)

Fit result is consistent with a *no signal hypothesis*: $NSig = 0.5 \pm 10$

Things to do

Plans for improvement:

- Add more data
- Improve selection (Fisher Discriminant and π^0 & Eta vetos)
- Re-optimize cuts
- Study systematics
- Extract result.

We will have a result by the summer!!!