Belle note 1486 Measurement of the decay $B^+ \to K^+ K^- \ell^+ \nu_\ell$ with B2BII

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Changelog

- 4. May: First submission of the note.
- 16. May: Added ROE validation section.
- $\bullet\,$ 7. July: update of whole BNOTE, B2BII chapter, finishing control fits.
- 19. August:
 - More details on the fit process and background splitting.
 - Adding sources of systematics
 - Ready for box opening
- 5. October:
 - Added fits to data
 - Added all sources of systematics
 - New model for uncertainty study
 - $-q^2$ calculation description
 - efficiency as a function of m_{KK} and q^2
 - Fit yields in bins of m_{KK} and q^2

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Chapter 1.

Introduction

Particle physics is an established branch of physics with a rich history in theory and experiments ever since the beginning of the $20^{\rm th}$ century. So far the experimental and theoretical research have shown us hand in hand that the universe consists of particles and interaction carriers. Particles of matter, or elementary particles, are divided in two groups – quarks and leptons. The quarks that we know today are called u (up), d (down), s (strange), c (charm), b (bottom) and t (top). Leptons are further split in charged leptons; e (electron), μ (muon), τ (tau lepton), and their corresponding neutrinos; ν_e (electron neutrino), ν_μ (muon neutrino) and ν_τ (tau neutrino). Interaction carriers are known as gauge bosons and they are γ (photon), g (gluon), W^{\pm} (charged weak bosons) and Z^0 (neutral weak boson). Theoretical calculations also predicted the recently discovered Higgs boson (H), which is responsible for the mass of all particles. Some of the particles above also have mirrored versions of themselves, called antiparticles, which exhibit somewhat different properties compared to their un-mirrored versions.

Combinations of quarks such as $q_1q_2q_3$ (hadrons) or $q_1\bar{q}_2$ (mesons) make up heavier particles. Examples of such particles are not only protons and neutrons, but also heavier particles which can be produced in processes involving high enough energies. Such heavy particles are unstable and decay into lighter ones. Together with the elementary particles and interaction carriers, three (out of four) of these interactions are joined in a theoretical model called the Standard Model (SM) [1, 2, 3, 4] (see Figure 1.1). Standard Model describes the electromagnetic, weak nuclear and strong nuclear interaction. General relativity – the theory of gravity – is not included in the Standard Model, since the two are incompatible on a mathematical level. However, due to its low coupling constant, gravity does not play a significant role in the world of subatomic particles. Experimental studies of particle processes give an insight into the mechanisms of basic interactions between them. By doing so, we are able to learn the secrets of the universe and how it all began.

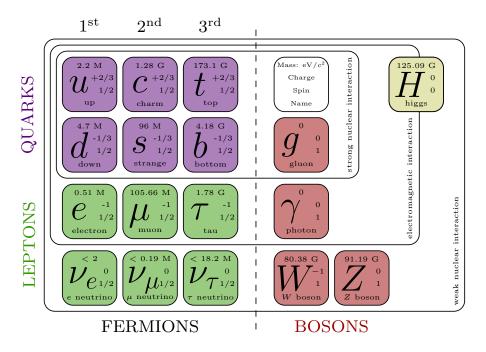


Figure 1.1.: A schematic representation of particles in the Standard Model.

This analysis revolves around decays of the so-called B mesons, which are particles consisting of a b quark and a light u or a d quark. The charged B^+ and the neutral B^0 meson have a structure of (\bar{b}, u) and (\bar{b}, d) , respectively, while the antiparticle B mesons are $B^-(b, \bar{u})$ and $\bar{B}^0(b, \bar{d})$. Perhaps one of the most surprising features of nature that can be studied with decays of B mesons is the CP symmetry violation (CP). CP symmetry is a combination of the C symmetry (charge conjugation) and the P symmetry (spatial inversion). A conservation of the CP symmetry would mean that the mirrored processes, in which all particles are exchanged by the corresponding anti-particles, proceed in exactly the same manner as the original processes. Today we know that this does not hold true for all cases and we, in fact, find processes which violate the CP symmetry. We also know that CP is related to the weak nuclear interaction. Here lies our motivation for studying decays of B mesons, since they exhibit a rich spectrum of decays which proceed via the weak nuclear interaction.

One of the most important properties of the weak interaction is the fact that it can change the flavor of particles. Such processes are forbidden for the electromagnetic and the strong nuclear interaction, but not for the weak one. Information about the quark transition probabilities is merged into a form of a complex matrix called the Cabibbo-Kobayashi-Maskawa (CKM) matrix [5, 6]

$$V_{CKM} = \begin{bmatrix} V_{ud} & V_{us} & V_{ub} \\ V_{cd} & V_{cs} & V_{cb} \\ V_{td} & V_{ts} & V_{tb} \end{bmatrix}.$$
 (1.1)

The CKM matrix is a unitary matrix and has only four parameters, which are free parameters of the theory and hence must be experimentally determined. The unitarity of the CKM matrix provides us with several mathematical identities, out of which the most relevant one for B meson physics is

$$V_{ud}V_{ub}^* + V_{cd}V_{cb}^* + V_{td}V_{tb}^* = 0. (1.2)$$

It can be represented by a triangle in the complex plane, called the unitarity triangle, shown in Figure 1.2. The sides and the angles of the unitarity triangle are related to the free parameters of the CKM matrix. All measurements of weak interaction processes involving B mesons depend on the four free parameters of the CKM matrix. Results of such measurements hence determine the sides and angles of the unitarity triangle. The goal is to then combine all such measurements and overconstrain the sides and angles of the unitarity triangle to check if all the sides meet. By improving such measurements one can check whether the SM is consistent, or if there are some contributing physics processes that we do not yet understand. Such processes are commonly referred to as "new physics" (NP). The measurements of the sides and angles of the triangle are done using different decays, with the most important input from B meson decays. This fact represents another motivation to study the B meson decays.

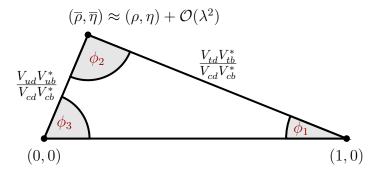


Figure 1.2.: The unitarity triangle in the Wolfenstein parametrization [7].

In this analysis, we focus on the V_{ub} CKM matrix element, which corresponds to $b \to u$ quark transitions. It has the smallest absolute value of all the CKM matrix elements and is currently determined with the largest uncertainty. Such quark transitions are present in charmless semileptonic B meson decays of the form

$$B^+ \to X_u^0 \ell^+ \nu_\ell, \tag{1.3}$$

where X_u^0 represents a charmless hadron with a u quark, and ℓ is one of the charged leptons, e, μ or τ . Measuring the decay rate of the B meson in such decays paves the way for the CKM matrix element determination. Decay rates are directly connected to the V_{ub} element as

$$d\Gamma \propto G_F^2 |V_{ub}|^2 |L^{\mu} \langle X_u | \bar{u} \gamma_u \frac{1}{2} (1 - \gamma_5) b |B\rangle|^2, \qquad (1.4)$$

where Γ is the decay width, G_F is the Fermi coupling constant, L^{μ} is the leptonic current and the expression in the Dirac brackets is the hadronic current. The factor $|V_{ub}|^2$ is the CKM element describing the $b \to u$ quark transition. Measurement of the V_{ub} CKM matrix element can be performed using two general approaches, with the exclusive or inclusive method, which are described below. Both methods

require an application of different experimental and theoretical techniques, so they provide largely independent determinations of $|V_{ub}|$. Currently, both methods also have comparable accuracies.

In the exclusive method, one studies the decays of B mesons to a specific charmless hadronic final state, such as $B \to \pi \ell \nu$. Clean determination of the $|V_{ub}|$ is possible due to precise experimental measurements along with reliable theoretical calculations. However, theoretical calculations are more challenging for decays to a specific final state, since hadronization of quarks has to be taken into account. There are also two main experimental challenges in this method. One has to reduce the abundant background from $B \to X_c \ell \nu$ processes since the $b \to c$ quark transition is much more probable than the $b \to u$ transition. The second experimental challenge is to separate B meson decays into a specific charmless hadronic final state from other $B \to X_u \ell \nu$ decays, since they populate roughly the same regions of the phase-space as the signal decay.

In the inclusive method, one studies the decays of B mesons to any charmless hadronic final state $B \to X_u \ell \nu$. In this case, the total decay rate for $b \to u \ell \nu$ can be calculated accurately since hadronization does not have to be taken into account. The greater challenge with this method is again the experimental measurement of the total decay rate due to the $B \to X_c \ell \nu$ background. Experimental sensitivity to V_{ub} is highest where $B \to X_c \ell \nu$ decays are less dominant. Theory and experiment have to compromise and limit the $|V_{ub}|$ determination to a region of phase-space where the signal-to-background ratio is good. Theoretical calculations take this into account by calculating the partial decay rate $\Delta\Gamma$, which is more challenging to determine than the total decay rate. One possible and often used approach to reduce $b \to c$ background is to reject all events with kaons present in the final particle selection. The procedure is called a K-veto. Kaons consist of an s quark, which is mainly produced in the dominant $b \to c \to s$ transition chain. This means that if a kaon is found in an event, it is very likely that it originates from a particle with a c quark, indicating the $b \to c$ process.

If $|V_{ub}|$ is determined with both of these methods, the values can be compared and potentially combined. It turns out that the consistency between the two results is only marginal, the difference is at a level of 3σ . The current world averages [8] of the exclusive (from $B^0 \to \pi^- \ell^+ \nu$) and inclusive (GGOU collab. [9]) methods are

$$|V_{ub}|_{\text{excl.}} = (3.65 \pm 0.09 \pm 0.11) \times 10^{-3},$$

$$|V_{ub}|_{\text{incl.}}^{\text{GGOU}} = (4.52 \pm 0.15 ^{+0.11}_{-0.14}) \times 10^{-3},$$
(1.5)

$$|V_{ub}|_{\text{incl.}}^{\text{GGOU}} = \left(4.52 \pm 0.15 \, {}^{+0.11}_{-0.14}\right) \times 10^{-3},$$
 (1.6)

where the first and the second uncertainties are the experimental and the theoretical, respectively. We see that inclusive measurements prefer higher values than exclusive ones. This is known as the V_{ub} puzzle. It is necessary to make further research as to why this difference occurs. The reason could be an unknown experimental or theoretical error, or it is even possible that some NP contributions occur. This analysis will focus on a possible reason that could be hidden in the selection mentioned before. By performing a K-veto, one discards all events with kaons in the final state in order to suppress $b \to c$ contributions. We focus on the charged $B \to KK\ell\nu$ decay, which is very similar to the $B \to \pi\ell\nu$, except for a production of an $s\bar{s}$ quark pair, which then combines with final state quarks to form kaons, as shown in Figure 1.3. In this case, we have kaons in the final state where the Bmeson decayed via a $b \to u$ process. Such decays were discarded in previous $|V_{ub}|$ determinations with the inclusive method, but in principle, they contribute to the result and should be taken into account. The results of this analysis should help us take a step closer towards solving the V_{ub} puzzle.

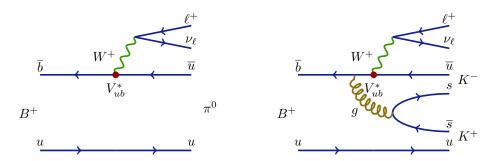


Figure 1.3.: Feynman diagrams for the $B^+ \to \pi^0 \ell^+ \nu_\ell$ decay (left) and the $B^+ \to K^- K^+ \ell^+ \nu_\ell$ decay (right).

Specifically, we will be focusing on the decays of the charged B mesons of the form $B^+ \to K^+ K^- \ell^+ \nu$, since it includes two charged kaons, as opposed to the case of the neutral B meson decay. The reason for this is a simpler decay chain and a higher reconstruction efficiency. All further occurrences of $B \to K K \ell \nu$ imply decays of the form $B^+ \to K^+ K^- \ell^+ \nu$ and its charge conjugated counterpart.

Chapter 2.

Data and Monte-Carlo Samples

The Belle detector acquired a dataset of about $L_0 \approx 710 \text{ fb}^{-1}$ of integrated luminosity in its lifetime at the $\Upsilon(4S)$ energy of 10.58 GeV, which corresponds to about $771 \times 10^6 \ B\bar{B}$ meson pairs. Additionally, several streams of Monte-Carlo (MC) simulated samples were produced, where each stream of MC corresponds to the same amount of data that was recorded the detector. The main focus of this work is to study a rare signal decay that is not necessarily produced abundantly or at all in the existing MC samples. In such cases, it is a common practice to produce specific samples of signal MC, where the abundance of the signal decay is much larger, enabling us to study its properties in greater detail.

The following samples were used in analysis

• data

- Belle on-resonance dataset of about L_0 integrated luminosity, measured at $\Upsilon(4S)$ resonance energy,
- Belle off-resonance dataset of about $1/10 \times L_0$ integrated luminosity, measured at 60 MeV below $\Upsilon(4S)$ resonance energy,
- signal MC, corresponding to about $400 \times L_0$,

• other MC

- generic on-resonance, 10 streams of B^+B^- and $B^0\bar{B}^0$ (denoted as charged and mixed) and 6 streams of $q\bar{q}$ produced at $\Upsilon(4S)$ resonance energy, where each stream corresponds to L_0 ,
- generic off-resonance, 6 streams of $q\bar{q}$ produced at 60 MeV below $\Upsilon(4S)$ resonance energy, where each stream corresponds to $1/10 \times L_0$,
- $-B \rightarrow X_u \ell \nu$ (denoted as ulnu), not included in previous MC samples, equal to an amount of $20 \times L_0$,
- other rare B meson decays (denoted as rare), not included in previous MC samples, equal to an amount of $50 \times L_0$.

2.1. Signal MC Production

The signal MC sample of $B^+ \to K^+ K^- \ell \nu_\ell$, including the charge conjugated B^- mesons, was produced using the mcproduzh [10, 11] package for producing Belle MC. The package accepts a decay file, which describes the decays to be generated. The decay file used for signal MC generation was the same as for the ulnu sample, since it includes the decays of interest. An additional MC skim was applied in order to select only events of interest with at least 2 kaons and a light lepton, all coming from the same B meson. This decreases the CPU consumption during the detector simulation and reconstruction.

The relevant processes which contribute to our signal decay are

- $B^+ \to a_{00} \ell^+ \nu_{\ell}$,
- $B^+ \to a_{20}\ell^+\nu_\ell$,
- $B^+ \to f_2 \ell^+ \nu_\ell$,
- $B^+ \to f_0 \ell^+ \nu_\ell$,
- $B^+ \to X_u^0 \ell^+ \nu_\ell$,

where a_{00} , a_{20} , f_2 , and f_0 are light, unflavored states which include further decays into a K^+K^- pair. The X_u^0 state represents a generic $u\bar{u}$ quark pair, which further hadronizes based on the PYTHIA quark hadronization model [12]. Figure 2.1 shows the invariant mass of the KK pair from various contributions of the MC generator. The light unflavored states have small contributions with resonant structures, while KK pairs from the X_u^0 state are more abundant and follow a wider and smoother distribution.

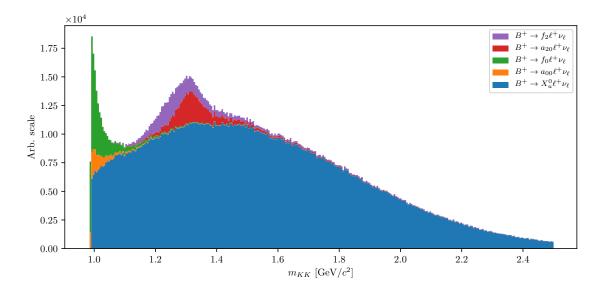


Figure 2.1.: Invariant mass of the KK pair from various contributions of the MC generator. The light unflavored states have small contributions with resonant structure, while KK pairs from the X_u^0 state are more abundant and follow a wider and smoother distribution.

The produced signal MC sample contains decays of the form $B \to KK\ell\nu$, as well as $B \to KKX\ell\nu$, where X can be any hadron as long as it satisfies all the selection rules of the decay. It is possible to calculate the MC branching fractions for each channel by making combinations of the particles directly from the generator output. Table 2.1 shows some of the most prominent $B \to KKX\ell\nu$ channels and their relative fractions. It is clear that our signal decay is the most abundant one, with a relative contribution of about 28 %, while other channels contribute only up to about 8 % or less. Additionally, our signal decay is the cleanest, while other decays include neutral particles like π^0 , which are harder to reconstruct and suffer from a decrease in efficiency due to reconstruction effects.

Channel	Ratio [%]	Channel	Ratio [%]
K^+K^-	28.14	$K^+K^-\rho^0$	1.93
$K^+K^-\pi^0$	8.94	$K^+ \bar{K}^0 ho^-$	1.84
$K^+ ar K^0 \pi^-$	8.71	$K^0K^-\rho^+$	1.83
$K^0K^-\pi^+$	8.70	$K^0 \bar{K}^0 ho^0$	0.00
$K^+K^-\pi^+\pi^-$	4.15	$K^{+}K^{-}\pi^{0}\pi^{0}$	0.86
$K^0ar{K}^0$	3.32	$K^{+}K^{-}\pi^{+}\rho^{-}$	0.69
$K^0ar{K}^0\pi^0$	3.26	$K^+K^-\rho^+\pi^-$	0.68
$K\bar{K}$ pair with η	7.08		
$K\bar{K}$ pair with ω	5.33		
Other	14.53		

Table 2.1.: Relative branching fractions of $B \to KKX\ell\nu$ decays by channel.

We generate about 1.3×10^9 events of the form $B \to X_u \ell \nu$, which corresponds to an integrated luminosity of about $L = 400 \times L_0$, where this value was obtained by normalizing the signal MC to the amount of signal candidates in the ulnu MC sample. This amounts to a total of about 9.37×10^6 generated signal events, and to a branching fraction

$$\mathcal{B}\left(B^{+} \to K^{+}K^{-}\ell^{+}\nu_{\ell}\right)_{GEN} = 1.53 \times 10^{-5},$$
 (2.1)

where ℓ is e or μ . During the analysis, the abundant signal MC sample is scaled down to correspond to the amount of data taken with the Belle detector.

2.2. Control Decay

In this analysis we are also define another B meson decay which occupies almost the same phase-space as our signal decay. This process can be used for the monitoring of our analysis steps, which are applied to both measured and simulated data. Any kind of difference between the two might indicate our procedure to be fine-tuned to simulated data, or some other similar problem.

We define the control decay of the form

$$B^+ \to \bar{D}^0 \ell^+ \nu$$
, $D^0 \to K^+ K^-$,

which is much more abundant and, most importantly, easy to suppress, since it only populates a very narrow region in the kaon invariant mass spectrum. Due to no extra particles in the D^0 decay, the kaon invariant mass is equal to $m_{KK} \approx m_{D^0}$ up to very good precision. By excluding this narrow region, we discard the majority of the control candidates, while discarding only a small amount of the signal candidates. A more quantitative description of suppressing control and other background candidates is written in Chapter 6.

Chapter 3.

Belle to Belle II Format Conversion

3.1. Conversion Procedure

The Belle experiment finished its data-taking run of 10 years at the end of 2010, after collecting a dataset of about 1 ab⁻¹. That year the Belle detector was shut down, the Belle II experiment started in its place and the focus moved to the construction of the Belle II detector and the development of the Belle II Analysis Framework (BASF2) [13]. However, Belle analyses are still on-going and Belle data is still being used today. BASF2 software, with its modular structure, has a more intuitive approach to performing analyses, however, since it was rewritten completely from scratch, it was designed for the incoming Belle II data, therefore usage of Belle data is outside of its scope.

In the Belle Collaboration, a task force was created in order to convert Belle data into Belle II format (B2BII) [14]. The B2BII package was developed as a part of BASF2 in order to convert the data and MC of the Belle experiment and make it available within BASF2. In addition to the convenience of the Belle data being processed in the more intuitive and advanced BASF2 framework, B2BII allows for estimation and validation of performances of various advanced algorithms being developed for Belle II. The conversion itself, however, is considered non-trivial. Although the conversion of the raw detector data would be possible, the reconstruction algorithms of BASF2 are optimized for Belle II and cannot be effectively applied to Belle data. To bypass this problem, reconstructed objects from PANTHER tables, a custom solution of the Belle collaboration based on C/C++ and Fortran, are mapped to their corresponding representations in BASF2. In this analysis, we use the developed converter package in order to analyze Belle data with the Belle II software.

The conversion in the B2BII package is divided into three BASF2 modules. The first module opens the Belle input files and reads the events into memory in the form of PANTHER tables. This module consists predominantly of reused BASF code. The second module applies various calibration factors, such as experiment- and run-dependent factors, to the beam energy, particle identification information, error matrices of the fitted tracks, etc. The module also applies some low-level selection criteria to reproduce removing background events as done within BASF. The actual conversion and the mapping of reconstructed objects are done in the last module. For more information see [15].

3.2. Validation

In order to make sure that the conversion was successful, a thorough validation is performed. This is done by comparing histograms of all physical quantities of the reconstructed objects on simulated and recorded events, processed with BASF and BASF2.

Our signal decay mode consists of three charged tracks, so track conversion should perform flawlessly. Additionally, energy measurement is also important in our analysis. In order to successfully determine the missing four-momentum in the event, we also need a correct conversion of the ECL clusters for photons and π^0 particles. Figures 3.1 to 3.3 show the basic physical properties of converted tracks, photons and π^0 particles, obtained with BASF and BASF2, and their difference, which is (up to numerical precision) equal to 0. The plots indicate that the conversion is successful in all aspects, so we can proceed with the analysis in the framework of BASF2.

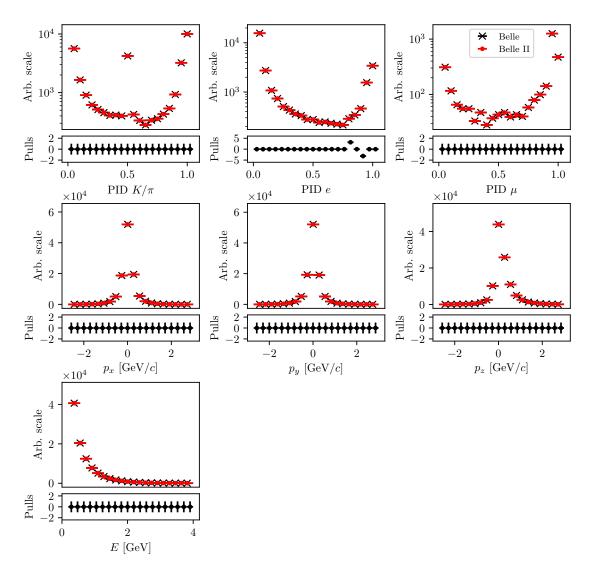


Figure 3.1.: Some of the more important physical properties of tracks for Belle and Belle II in the conversion process.

The histograms seem to overlap and the conversion is assumed to be successful.

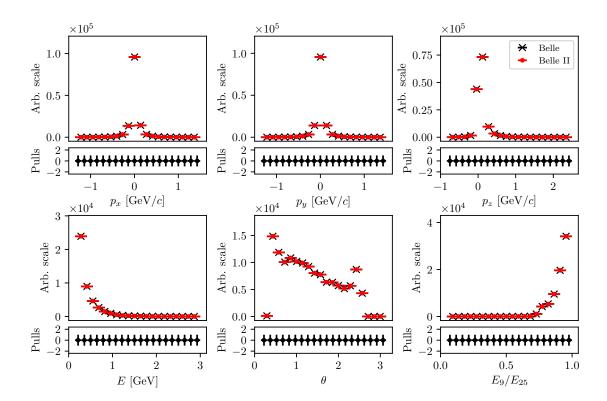


Figure 3.2.: Some of the more important physical properties of photons for Belle and Belle II in the conversion process.

The histograms seem to overlap and the conversion is assumed to be successful.

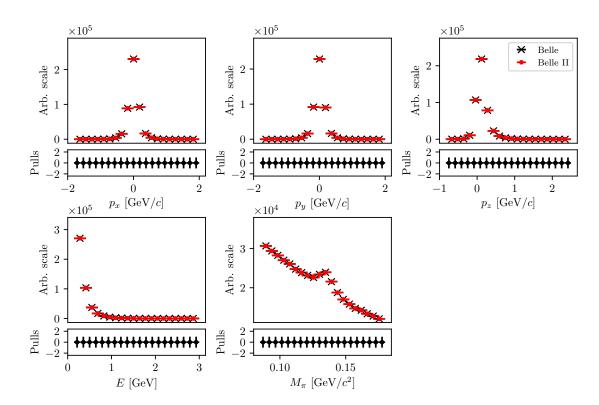


Figure 3.3.: Some of the more important physical properties of π^0 particles for Belle and Belle II in the conversion process. The histograms seem to overlap and the conversion is assumed to be successful.

Chapter 4.

Event Reconstruction

In this chapter, the procedure for event reconstruction of the B meson decay $B \to KK\ell\nu$ is shown, starting with final state particle selection and then combining them to obtain the B meson candidates.

4.1. Final State Particles Selection

The Belle detector is not able to detect all kinds of particles. Neutrinos are one such example, since they escapes detection, therefore we can only reconstruct the charged tracks in the decay, which are the two charged kaons (K) and the light lepton (e or μ). These are some of the particles which are commonly referred to as the final state particles (FSP). Final state particles have a long lifetime and are usually the particles that we detect when they interact with the material in the detector.

At this point in the analysis, we do not apply any specific particle selections yet, which results in a large number of available particles and their combinations, and, consequently, the computation time. In order to minimize this effect, we perform this part of the study on a smaller subset of the available generic MC, experiments no. 23 and 31, which correspond to an integrated luminosity of 6.273 fb⁻¹ and 17.725 fb⁻¹, respectively. We chose these two experiments to approximate the appropriate ratio of SVD1 and SVD2 data in the full Belle MC.

Leptons

Figures 4.1 and 4.2 show the impact parameters d_0 and z_0 , the momentum in $\Upsilon(4S)$ center-of-mass system (CMS), and the PID information for true and fake electrons and muons from any source, where true electrons/muons from the signal B meson decay are shown separately. True and fake implies that the particles are correctly or wrongly reconstructed, with respect to the generated MC truth. The difference between the true leptons from any source and those from signal decays is due to the distinct kinematics of the parent's decay.

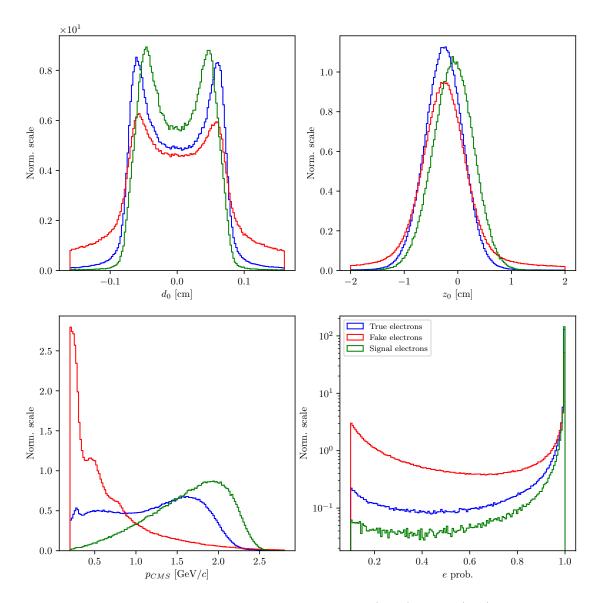


Figure 4.1.: Normalized properties of true (blue), fake (red) electrons from any source, and true electrons from signal B candidates (green).

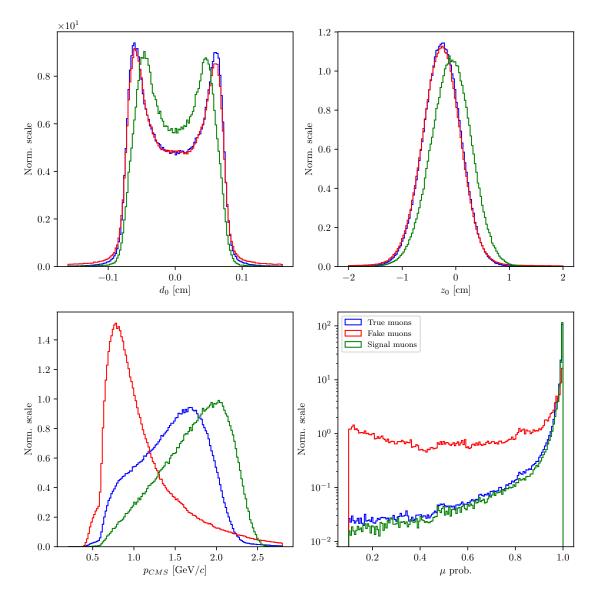


Figure 4.2.: Normalized properties of true (blue), fake (red) muons from any source, and true muons from signal B candidates (green).

Based on these distributions, we can define a selection criteria

- $|d_0| < 0.1 \text{ cm}$,
- $|z_0| < 1.5 \text{ cm},$
- $p_{CMS} \in [0.4, 2.6]$ GeV/c for electrons,
- $p_{CMS} \in [0.6, 2.6]$ GeV/c for muons.

After this we can determine the optimal PID selection for electrons and muons, where we optimize the criteria by maximizing the standard definition of figure of $merit\ (FOM)$, defined in Eq. (4.1)

$$FOM = \sqrt{\mathcal{EP}} \propto \frac{S}{\sqrt{S+B}},$$
 (4.1)

where the argument in the square root is the product of the efficiency (\mathcal{E}) and the purity (\mathcal{P}) . The definitions of signal (S) and background (B) are somewhat fluid throughout the analysis and need to be defined for each FOM separately. In this section we define two representations of S and B. In FOM_1 the signal S represents correctly reconstructed final state particles, while in FOM_2 the signal S represents correctly reconstructed final state particles coming from correctly reconstructed S meson candidates. In both cases the background represents all other particle candidates which do not satisfy the conditions of S.

The FOM plots are shown in Figures 4.3 and 4.4. The selection criteria are based on PID cuts used for PID efficiency calibration. The optimal value for the PID cuts is equal to the largest available value for true leptons (FOM_1) , as well as for true leptons from signal B decays (FOM_2) , so selections via both methods are the same. The optimal PID selection criteria for leptons are then

- e prob. > 0.9 for electrons,
- μ prob. > 0.97 for muons.

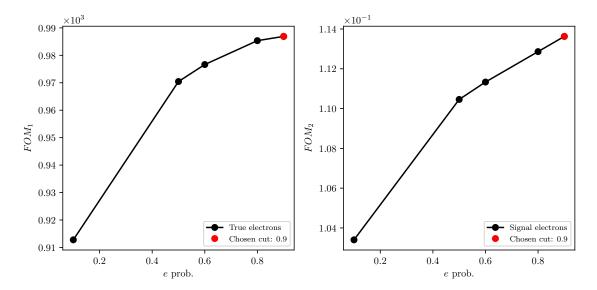


Figure 4.3.: FOM optimizations of the PID selection for true electrons (left) and true electrons from signal B candidatess (right).

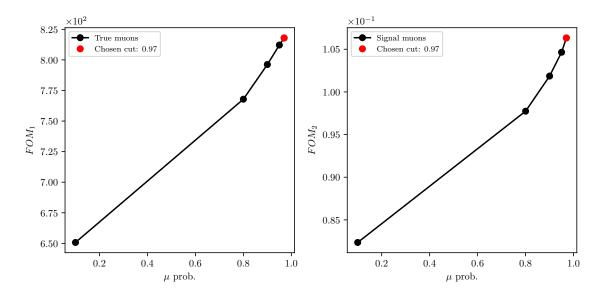


Figure 4.4.: FOM optimizations of the PID selection for true muons (left) and true muons from signal B candidates (right).

Kaons

We repeat the procedure for both kaons. Figure 4.5 shows the impact parameters d_0 and z_0 , the momentum in $\Upsilon(4S)$ center-of-mass system (CMS), and the PID information for true and fake kaons, where true kaons from the signal decay are shown separately.

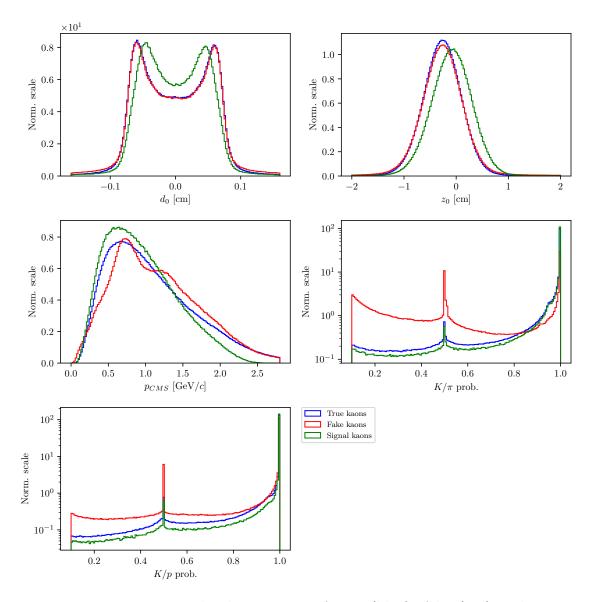


Figure 4.5.: Normalized properties of true (blue), fake (red) and true kaons (green) from signal B candidates.

We define the kaon selection criteria in the same manner as in the case for leptons

- $|d_0| < 0.15$ cm,
- $|z_0| < 1.5 \text{ cm},$
- $p_{CMS} \in [0, 2.5] \text{ GeV}/c.$

The PID optimization, in this case, is taken in two steps. First, we optimize the selection on K/π , and after that, the K/p separation probability. Figure 4.6 shows the optimization procedure for PID cuts on kaon candidates.

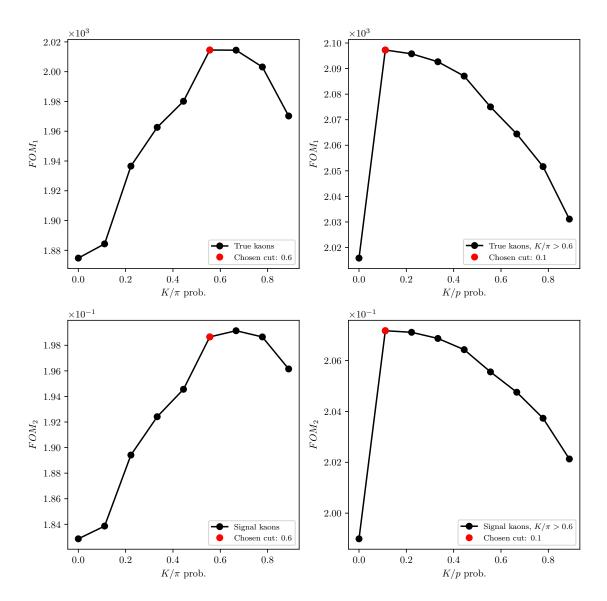


Figure 4.6.: FOM optimizations of the PID selection for true kaons (top) and true kaons from signal B candidates (bottom). The plots on the left show the optimization of the first step for the K/π probability cut, and the plot on the right the K/p probability cut.

The optimal PID selection for kaons is

- $K/\pi > 0.6$,
- K/p > 0.1.

4.2. Pre-selection of First B Meson Candidates

In this section, we use the charged particle candidates from the previous section to make particle combinations, which correspond to B meson candidates. When a B meson candidate is selected, additional features can be calculated and used for background rejection. Since we are still performing this part of the study on a

smaller subset of the full available MC sample, we will perform an under-optimized selection based on the FOM optimizations, in order to optimize them later on the full Belle MC sample.

Since the missing neutrino escapes detection, we reconstruct the B mesons using the following final state particles

$$B^+ \to K^+ K^- e^+,$$

 $B^+ \to K^+ K^- \mu^+,$

and similarly for B^- . When an arbitrary combination is obtained, we perform a vertex fit of the three tracks in order to discard combinations with a low probability of tracks coming from the same point. B mesons have a relatively long lifetime and decay along the z axis of the detector in the direction of the boost, so the vertex fit is enforced with an IPTUBE constraint, which constrains the vertex to an elongated ellipsoid along the beam direction. We require that the fit converged and apply a selection on the minimal fit probability. The fit probability for signal and background B meson candidates is shown in Figure 4.7 (left). We perform an FOM optimization of this selection, which is shown in Figure 4.7 (right) for the subset of the Belle MC sample. In this, and in the following cases, the definition of S from Eq. (4.1) are correctly reconstructed B meson candidates with a missing neutrino which are not coming from a $b \to c$ transition.

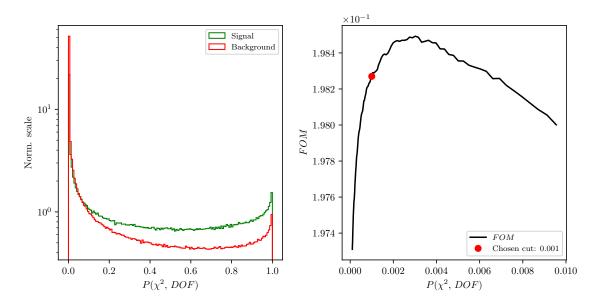


Figure 4.7.: Normalized vertex fit probability distribution for signal and background B meson candidates in the logarithmic scale (left) and the FOM optimization of the vertex fit probability (right) for the subset of the full Belle MC sample.

The chosen pre-selection on the fit probability is

•
$$P(\chi^2, NDF) > 1.0 \times 10^{-3}$$
.

With the neutrino being the only missing particle on the reconstructed side, it is possible to determine the angle between the direction of the reconstructed B (denoted as $Y \to KK\ell$) and the nominal B, as

$$p_{\nu} = p_B - p_Y, \tag{4.2}$$

$$p_{\nu}^{2} = m_{\nu}^{2} = m_{B}^{2} + m_{Y}^{2} - 2E_{B}E_{Y} + 2\vec{p}_{B} \cdot \vec{p}_{Y} \approx 0, \tag{4.3}$$

$$\cos(\theta_{BY}) = \frac{2E_B E_Y - m_B^2 - m_Y^2}{2|\vec{p}_B||\vec{p}_Y|},\tag{4.4}$$

where p denotes a scalar, \vec{p} a vector, and p a four-vector. All the energy and momenta above are calculated in the CMS frame. The mass of the neutrino is equal to 0 up to a very good precision, so we use it in Eq. (4.3). Additionally, we substitute the unknown energy and momentum magnitude, E_B and $|\vec{p}_B|$, of the B meson in Eq. (4.4), with quantities from the well known initial conditions

$$E_B = E_{CMS}/2, (4.5)$$

$$|\vec{p}_B| = p_B = \sqrt{E_B^2 - m_B^2},$$
 (4.6)

where E_{CMS} is the total energy of the e^+e^- collision in the CMS frame and m_B is the nominal mass of the B meson. This improves the resolution of $\cos(\theta_{BY})$, which leads to better signal-to-background discrimination.

For the correctly reconstructed candidates, this variable lies in the [-1,1] region, though only to a certain precision, due to the finite detector resolution. Background candidates, however, populate also the non-physical regions, as shown in Figure 4.8 (left).

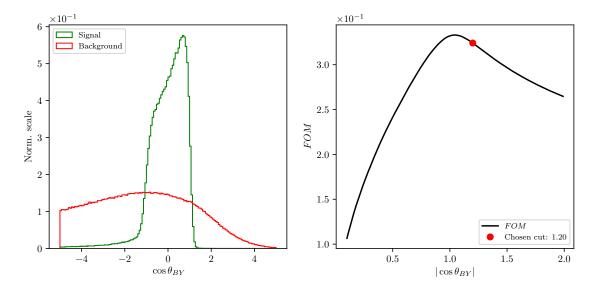


Figure 4.8.: The normalized $\cos \theta_{BY}$ distribution for signal and background B meson candidates (left) and the FOM optimization of the $\cos \theta_{BY}$ variable (right) for the subset of the full Belle MC.

We again impose an under-optimized selection on this variable from Figure 4.8 (right) to discard a large amount of background on this subset of the full Belle MC

•
$$|\cos(\theta_{BY})| < 1.20.$$

4.3. Loose Neutrino Reconstruction

We are not able to directly determine the four-momentum of the missing neutrino. However, due to the detectors geometry, which almost completely covers the full solid angle, and due to the well known initial conditions of the $\Upsilon(4S)$ meson, it is possible to determine the kinematics of the missing neutrino indirectly. Specifically, this is performed by reconstructing the companion B meson via summing up the four-momenta of all the FSP particles in the event, which were not used in the reconstruction of the signal side B meson. This is known as the untagged method, since we are not using any kind of tagging method to reconstruct the companion B meson. The particles used in the indirect companion B meson reconstruction are also said to belong to the rest of the event (ROE).

Due to the beam background in the detector, material interactions, or other processes, random tracks and clusters enter our event and get reconstructed as part of the physics process we want to study. In order to remedy this, we perform an extensive clean-up of the tracks and clusters in the ROE side before calculating the four-momentum of the missing part of the event. The clean-up procedure is performed separately on tracks and clusters, and uses multiple steps with multivariate analysis (MVA) algorithms in order to separate good tracks and clusters from the background ones, which also populate the ROE. Then, for each ROE object, a ROE mask is created for tracks and clusters, which narrates the use of this object in the final calculations of the ROE four-momentum. From this point on, we assume the ROE to be efficiently cleansed of extra tracks and clusters. A more detailed description of the ROE clean-up can be found in Chapter 5.

The total missing four-momentum in the event can be determined as

$$p_{miss} = p_{\Upsilon(4S)} - \sum_{i}^{\text{Event}} (E_i, \vec{p}_i), \qquad (4.7)$$

$$p_{miss} = p_{\Upsilon(4S)} - \left(p_Y - \sum_{i}^{\text{Rest of event}} (E_i, \vec{p_i})\right), \tag{4.8}$$

where the summation runs over all charged and neutral particles in the defined set with

$$\mathbf{p}_i^{\text{neutral}} = (p_i, \vec{p_i}) \quad \text{and} \quad \mathbf{p}_i^{\text{charged}} = \left(\sqrt{m_i^2 + p_i^2}, \vec{p_i}\right)$$
 (4.9)

We assume all neutral particles to be massless. For charged tracks in the ROE, a mass hypothesis needs to be defined in order to determine the energy of the track. After the ROE clean-up, we make the following procedure of choosing the mass hypothesis

- 1. e, if e prob. $> \mu$ prob. and e prob. > 0.9,
- 2. otherwise μ , if μ prob. > e prob. and μ prob. > 0.97,
- 3. otherwise K, if K/π prob. > 0.6,
- 4. otherwise π .

We calculate the square of the missing mass, m_{miss}^2 , which is consistent with zero, if signal-side neutrino is the only missing particle in the event. The m_{miss}^2 distribution is shown in Eq. (4.11).

$$p_{\nu} = p_{miss} = (E_{miss}, \vec{p}_{miss}),$$
 (4.10)

$$p_{\nu} = p_{miss} = (E_{miss}, \vec{p}_{miss}),$$
 (4.10)
 $m_{miss}^2 = p_{miss}^2 = p_{\nu}^2 = m_{\nu}^2 \approx 0.$ (4.11)

Since the detector performance is not perfect, the distribution of the m_{miss}^2 variable has a non-zero width. Additionally, a tail is introduced due to missing particles like neutrinos, other neutral undetected particles such as K_L^0 , or simply missing tracks, due to detection failure. Figure 4.9 shows the distribution of m_{miss}^2 , as defined with the missing four-momentum in Eq. (4.10). Correctly reconstructed candidates, which come from events, where the other B meson decayed via a hadronic decay mode, peak at zero. If the companion B meson decayed (semi-)leptonically, candidates are shifted to larger values of this variable, even if the event in question is a signal event. For this purpose, we define a subset of all signal candidates, which come from events where the companion B meson decayed hadronically and all of its particles were taken into account correctly. We only allow for missing photons, since they are frequently radiated due to bremsstrahlung effects from final-state electrons and they typically do not have a big impact on the four-momentum of the final candidate. We denote this subset as the *perfect* signal. This subset is used to correctly define the clean-up parameters and is not used in any reconstruction steps, since we cannot know in data which neutral particles are actually missing.

Due to this fact, we impose a selection on the m_{miss}^2 variable, in order to partially discard candidates with spoiled properties, even if it was in principle a correct combination of FSP particles on the signal side. The selection was chosen based on the optimization of the FOM, where in this case the definition of S were perfectly reconstructed signal candidates. The chosen selection is

•
$$|m_{miss}^2| < 3.0 \text{ GeV}/c^2$$
,

which is also under-optimized at this point, due to the same reasons as in the cases above.

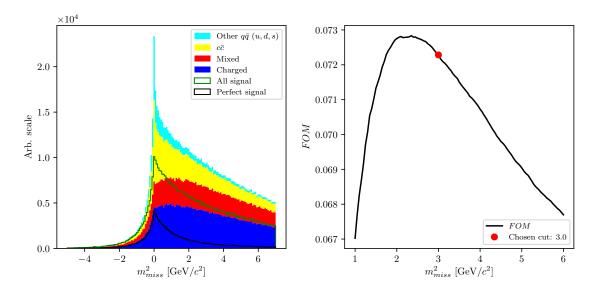


Figure 4.9.: m_{miss}^2 distribution for signal and various types of background. All signal (green) and perfect signal (black) are scaled up equally.

The main uncertainty of the neutrino four-momentum, defined in Eq. (4.10), comes from the energy uncertainty. It is a common practice to substitute the missing energy with the magnitude of the missing momentum, since the momentum resolution from the measurement is much better,

$$p_{\nu} = (|\vec{p}_{miss}|, \vec{p}_{miss}).$$
 (4.12)

This also fixes the neutrino mass to $0 \text{ GeV}/c^2$. The newly defined neutrino four-momentum can be added to the four-momentum of the $Y(KK\ell)$ candidate to obtain the full B meson four-momentum and calculate the traditional M_{BC} and ΔE variables

$$\Delta E = E_B - E_{CMS}/2,\tag{4.13}$$

$$M_{BC} = \sqrt{(E_{CMS}/2)^2 - |\vec{p}_B|^2}.$$
 (4.14)

Since the final fit will be performed over ΔE and M_{BC} , we define the fit region

- $M_{BC} \in [5.1, 5.295] \text{ GeV}/c^2$,
- $\Delta E \in [-1.0, 1.3] \text{ GeV}.$

Figure 4.10 shows the distributions of ΔE (left) and M_{BC} (right) for signal and major types of background after the pre-selection. Both signal components (all signal and perfect signal) are scaled up with respect to the background components but are in proper scale one to another. The effects of missing particles are clearly seen based on the shape difference between full and perfect signal.

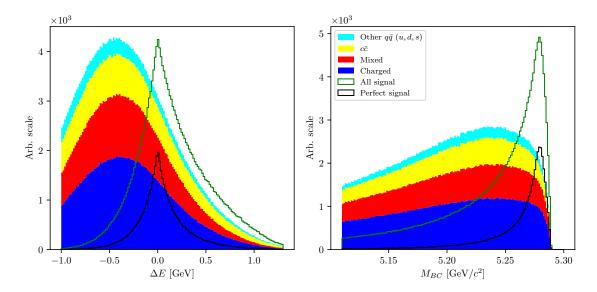


Figure 4.10.: Distributions of ΔE (left) and M_{BC} (right) for signal and major types of background after the pre-selection. Both signal components are scaled up with respect to the background components, but are in proper scale one to another. The perfect signal has a much better resolution in both distributions, since the event is perfectly reconstructed.

4.3.1. q^2 calculation

 q^2 is the squared Lorentz invariant of the four-momentum, also known as momentum transfer squared, which is transferred from the B meson to the W boson. There are several possible calculations of this variable which offer a different resolution. In this analysis, we follow the calculation procedure of q^2 from [16], which yields the best resolution.

For correctly reconstructed events, Eq. (4.13) satisfies the condition $\Delta E \approx 0$ within precision. It is possible to rescale the neutrino energy in such way that we fix ΔE to zero, meaning

$$\Delta E' = (E_Y + \alpha E_\nu) - E_{CMS}/2 = 0. \tag{4.15}$$

and calculate a corrected value of M_{BC}

$$M'_{BC} = \sqrt{(E_{CMS}/2)^2 - |\vec{p}_Y + \alpha \vec{p}_\nu|^2}.$$
 (4.16)

The neutrino momentum resolution dominates the ΔE uncertainty, so the correction factor α reduces this effect.

A second correction can be applied by rotating the direction of the neutrino momentum by a small angle with respect to the reconstructed one. Such an angle is chosen in order fix the value of M'_{BC} to the nominal mass of the B meson, m_B .

The corrected neutrino momentum is then fixed to expected values of ΔE and M_{BC} , and is solely used for the q^2 calculation. With p_{ℓ} as the reconstructed lepton four-momentum, we define the q^2 as

$$q^2 = q^2 = (p_\ell + p_\nu)^2$$
. (4.17)

The q^2 distributions and the corresponding resolutions are shown in Figure 4.11, along with other methods of q^2 determination.

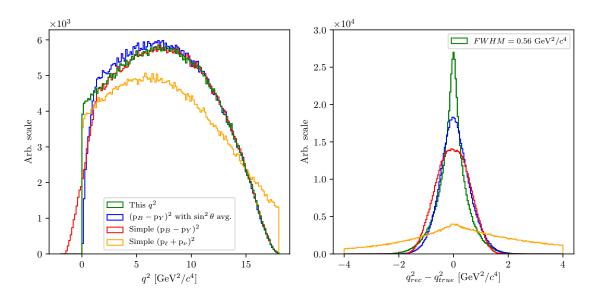


Figure 4.11.: Distributions of q^2 (left) and q^2 resolution (right) for various methods of q^2 calculation. The green distribution follows the procedure in [16], the blue distribution takes into account the weighted average of the B meson direction [17], and the red and orange distributions are straight-forward calculations with available information in the reconstruction. The q^2 calculation in red assumes a resting B meson in the CMS frame, and the calculation in orange uses the neutrino four-momentum from Eq. (4.10).

One must bear in mind that even though the chosen calculation yields the most precise result, this does not affect the correctness of the q^2 model, which was used in MC simulation (ISGW2 [18]). Since the signal decay has not been observed yet, we do not have a good description of the decay model, and we treat this as a source of systematic uncertainty in this analysis.

4.4. Final Stage Optimization

With the charge particle selection and a rough selection of the B meson particles in place, we can now afford to run the reconstruction procedure over all of the 10 streams of the full available Belle generic MC. After obtaining the full reconstructed sample, the first task is to optimize the under-optimized selection criteria from the previous stage. Repeating the procedure on the full sample results in the FOM shapes shown in Figure 4.12, with the optimal selection criteria

- $P(\chi^2, NDF) > 6.0 \times 10^{-3}$,
- $|\cos(\theta_{BY})| < 1.05$.

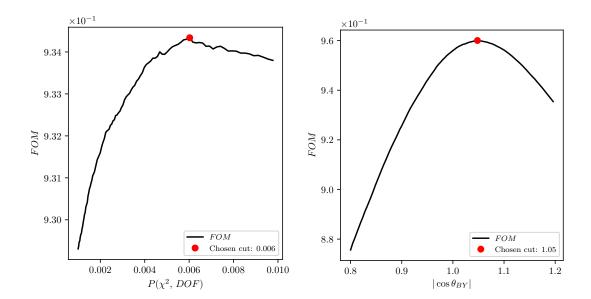


Figure 4.12.: The FOM optimization of the vertex fit probability (left) and the $\cos \theta_{BY}$ variable (right) for the full Belle MC sample.

With further optimizations, we will be fine-tuning the signal-to-background ratio. The most prominent and distinguishable part of our signal is the perfect signal. For this purpose, we define a signal region in ΔE and M_{BC} , where most of our perfectly reconstructed candidates lie. We use this region for all of the following optimization steps in this chapter and also in the background suppression (Chapter 6), since this allows us to better improve the signal-to-background ratio. The 2D FOM optimization of the optimal ΔE and M_{BC} signal region is shown in Figure 4.13.

The signal region is defined as

- $M_{BC} > 5.271 \text{ GeV}/c^2$,
- $|\Delta E| < 0.126$ Gev.

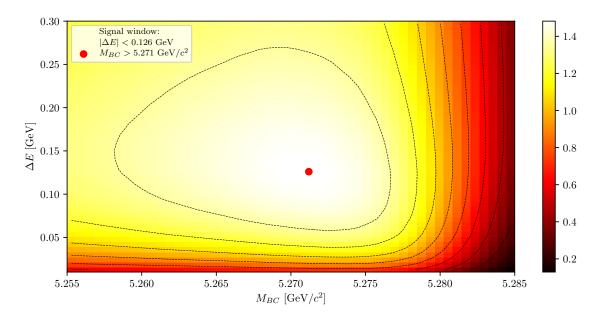


Figure 4.13.: 2D FOM optimization of the signal region definition, where most of the perfectly reconstructed candidates are located.

With the signal window defined, we can tighten the selection on m_{miss}^2 , which we intentionally left loose before the signal categorization. With the FOM optimization of perfectly reconstructed candidates inside the signal region, shown in Figure 4.14, the optimal selection of m_{miss}^2 range is

• $|m_{miss}^2| < 0.975 \text{ GeV}/c^2$.

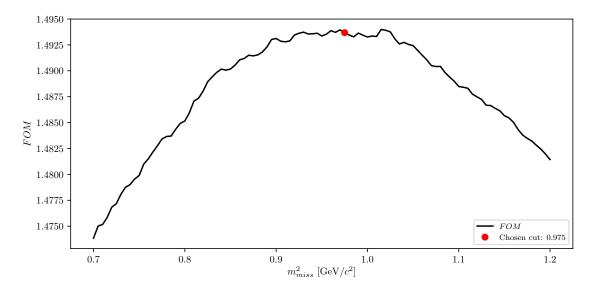


Figure 4.14.: FOM optimization of the m_{miss}^2 selection in the signal region.

4.5. Charge product categorization

The missing information, due to an escaping neutrino in our reconstructed channel, is replaced by the information from the companion B meson. Since this is an untagged reconstruction, the quality of the companion B meson affects the properties of the signal candidate. Perfect reconstruction of a hadronically decayed companion B meson results in pronounced peaks at $\Delta E \approx 0$, $m_{miss}^2 \approx 0$, and $M_{BC} \approx m_B$, while imperfect reconstruction, due to any kind of missing particles, produces tails, shift, or simply a worse resolution of the mentioned distributions. These effects are undesired, since they make it harder to separate signal from background.

To remedy this, we look at the charge product of the reconstructed B meson and the ROE object. For correctly reconstructed events, this should have a value of

$$q_{B^{\pm}}q_{B^{\mp}} = -1, (4.18)$$

however, this value is distributed due to missing, or additional background charged particles in the ROE. Figure 4.15 shows various signal distributions of ΔE and M_{BC} in arbitrary (left) and normalized (right) scales. We find the relative ratios of 67.86 % and 32.14 % for correct and wrong values of the charge product. Correctly reconstructed events represent the majority of the signal candidates and also have the best resolution in ΔE and M_{BC} , hence we proceed with the analysis by imposing the selection in Eq. 4.18.

While this selection introduces a drop in the signal efficiency of about 32.14 %, it improves the resolution of our signal ΔE and M_{BC} distributions and also the signal-to-background ratio, where the latter changes from 0.95×10^{-3} to 1.09×10^{-3} .

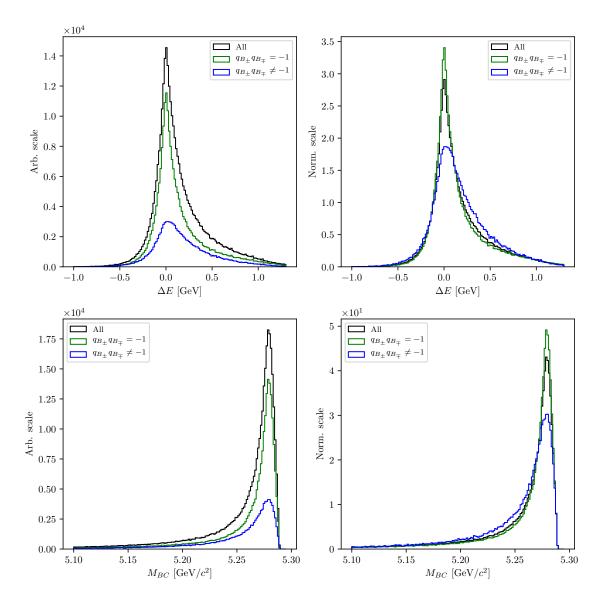


Figure 4.15.: Signal distributions of ΔE and M_{BC} based on the charge product of both B mesons in the event. The plots on the left show the distributions in an arbitrary scales, while the plots on the right show the normalized distributions.

4.6. Selection Summary

In this section, one can find the full summary of final selection criteria in the event reconstruction, from FSP particles up to the B meson.

• FSP particles:

- electrons: $|d_0| < 0.1 \text{ cm}, |z_0| < 1.5 \text{ cm}, p > 0.6 \text{ GeV}/c,$ $p_{CMS} \in [0.4, 2.6] \text{ GeV}/c, eID > 0.9,$
- muons: $|d_0| < 0.1$ cm, $|z_0| < 1.5$ cm, $p_{CMS} \in [0.6, 2.6]$ GeV/c, $\mu ID > 0.97$,
- kaons: $|d_0| < 0.15$ cm, $|z_0| < 1.5$ cm, $p_{CMS} < 2.5$ GeV/c, $K/\pi~ID > 0.6, K/p~ID > 0.1,$

\bullet B meson candidates:

- standard selection: $P(\chi^2, DOF) > 6 \times 10^{-3}, |\cos \theta_{BY}| < 1.05, |m_{miss}^2| < 0.975 \text{ GeV}/c^2,$
- fit region selection: $\Delta E \in [-1.0, 1.3]$ GeV, $M_{BC} \in [5.1, 5.295]$ GeV/ c^2 ,
- signal region selection: $|\Delta E| < 0.126 \text{ GeV}, M_{BC} > 5.271 \text{ GeV}/c^2$,
- charge categorization: $q_{B^{\pm}}q_{B^{\mp}}=-1$.

Chapter 5.

Rest of Event Clean-up

Continuing from Section 4.3, the description of the ROE clean-up process is described here.

Training the MVA classifiers follows the same recipe for all the steps in this chapter. For each step, we run the B meson reconstruction on Signal MC with a generic companion B meson. For every correctly reconstructed signal B meson candidate we save the necessary information for each MVA step (e.g. properties of ROE clusters). Only correctly reconstructed B candidates are chosen here, to prevent leaks of information from the signal side to the ROE side.

5.1. Machine Learning Setup

Throughout the analysis we use the Fast-BDT algorithm as the main machine learning model, in order to perform the multivariate analysis steps, unless explicitly stated otherwise. The Fast-BDT algorithm [19] is an optimized version of the boosted decision trees (BDT) algorithm, which is one of the most commonly used in the world of machine learning. As the name suggests, it is based on the method of decision trees. The decision tree algorithm generates decision trees from the data to solve classification and regression problems, where each decision tree splits the data in at least two groups. A schematic of such a decision forest is shown in Figure 5.1, where the forest consists of several trees, each with a certain depth.

The number of trees in the classifier and the depth level of a single tree are two hyper-parameters of the machine learning classifier, which we optimize in each case of applying the MVA step. More information about the MVA training, hyper-parameter optimization and feature importance for each MVA step in this chapter can be found in Appendix A.

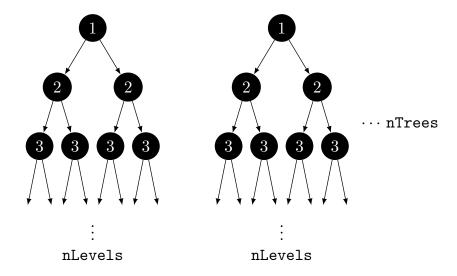


Figure 5.1.: A schematic representation of decision trees in a decision forest.

5.2. Clusters Clean-up

Photons originate from the IP region, travel to the ECL part of the detector in a straight line, and produce a cluster. The direction of the photon is determined via the location of the cluster hit in the ECL and the energy of the photon is directly measured via the deposited energy. This way the four-momentum of photons is determined and used in Eq. (4.8).

Most of the photons in events with B mesons come from processes such as $\pi^0 \to \gamma\gamma$ decays, which are interesting for physics. However, a lot of hits in the ECL are also created by photons coming from the beam-induced background or secondary interactions with the detector material, which we are usually not interested in, except in cases involving material studies. Photons of the first kind should be taken into account when calculating the missing four-momentum, while photons, which are not directly related to the collision, add extra energy and momentum to the event. In the first step of the clusters clean-up, we train an MVA which recognizes good π^0 candidates and apply this information to the daughter photons. This represents a sort of a π^0 origin probability, which peaks at or is equal to 0 for photons not coming from π^0 particles, and peaks at 1 otherwise. This information is used as an additional classifier variable in the next step of the clean-up, where we train to recognize good photons in an event.

5.2.1. π^0 MVA Training

The training dataset of π^0 candidates contains

- 183255 target candidates,
- 200000 background candidates,

where the definition of a target is that both photon daughters, which were used in the reconstruction of the π^0 , are actual photons and real daughters of the π^0 particle. We use π^0 candidates from the converted Belle particle list and select those with the invariant mass in the range of $M \in [0.10, 0.16]$ GeV. After that, we perform a mass-constrained fit on all π^0 candidates, keeping only the ones for which the fit converged.

The input variables used in this MVA are

- p and p_{CMS} of π^0 and γ daughters,
- fit probability of the mass-constrained fit, invariant mass and significance of mass before and after the fit,
- angle between the photon daughters in the CMS frame,
- cluster quantities for each daughter photon
 - $-E_9/E_{25}$
 - theta angle,
 - number of hit cells in the ECL,
 - highest energy in cell,
 - energy error,
 - distance to closest track at ECL radius.

The classifier output variable is shown in Figure 5.2.

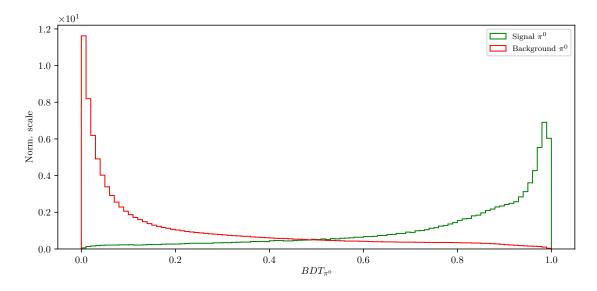


Figure 5.2.: Classifier output of the π^0 training for signal and background π^0 candidates.

5.2.2. γ MVA Training

In this MVA training, we take the π^0 classifier output of the previous training as an input in order to train a classifier to distinguish between good and bad photons. The π^0 probability information from the previous step is applied to all photon pairs, which pass the same π^0 selection criteria, as defined in the previous step. Since it is possible to have overlapping pairs of photons, the π^0 probability is overwritten in the case of a larger value, since this points to a greater probability of a correct photon combination. On the other hand, some photon candidates fail to pass the π^0 selection. These candidates have a fixed value of π^0 probability equal to zero.

The training dataset of γ candidates contains

- 171699 target candidates,
- 177773 background candidates,

where the definition of a target is that the photon is an actual photon, which is related to a primary MC particle. This tags all photon particles from secondary interactions as background photons. We use the converted γ candidates from the existing Belle particle list.

The input variables used in this MVA are

- p and p_{CMS} of γ candidates,
- π^0 probability,
- cluster quantities
 - $-E_{9}/E25,$
 - theta angle,
 - number of hit cells in the ECL,
 - highest energy in cell,
 - energy error,
 - distance to closest track at ECL radius.

The classifier output variable is shown in Figure 5.3.

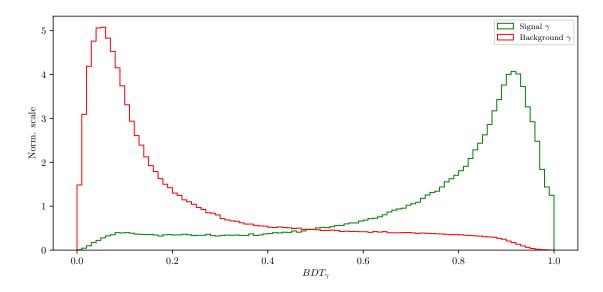


Figure 5.3.: Classifier output of the γ training for signal and background γ candidates.

With the classifier output at hand, we apply a selection to the photon particle list. The selection optimization is shown in Figure 5.4 (left), with the optimal selection on the γ classifier output at

• $BDT_{\gamma} > 0.5045$.

Figure 5.4 (right) shows the LAB frame momentum of the photons in the logarithmic scale, before and after the selection. The signal efficiency and background rejection of this clean-up step are

• Signal efficiency: $\epsilon_{SIG} = 83.2 \%$,

• Background rejection: $1 - \epsilon_{BKG} = 81.2 \%$.

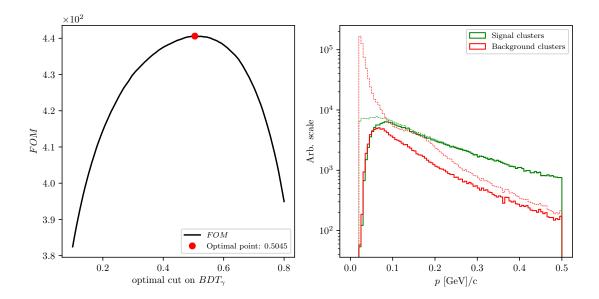


Figure 5.4.: The *FOM* of the classifier output optimization (left) and momentum magnitude in the LAB frame (right) of signal and background photon candidates before (dashed) and after (solid) the optimal selection.

The event is now considered to be clean of extra clusters.

5.3. Tracks Clean-up

Charged particles leave hits in the detector, which are then grouped into tracks by advanced tracking algorithms. The track is fitted and the track momentum is determined. With the help of particle identification information (PID), we are able to make an intelligent decision about the mass hypothesis of the particle and thus reconstruct the four-momentum of the charged particle, which is then taken into account in the loop in Eq. (4.8).

Most of the quality (good) tracks, which come from physics event of interest, come from the IP region, where the collisions occur. Cleaning up the tracks is a more complex procedure than cleaning up the clusters. The following facts need to be taken into account

- (a) good tracks can also originate away from the IP region, due to decays of long-lived particles, such as $K_S^0 \to \pi^+\pi^-$,
- (b) charged particles from background sources produce extra tracks or duplicates,
- (c) low momentum charged particles can curl in the magnetic field and produce multiple tracks,
- (d) secondary interactions with detector material or decays of particles in flight can produce "kinks" in the flight directory, resulting in multiple track fit results per track.

Schematics of all the cases mentioned above are shown in Figure 5.5.

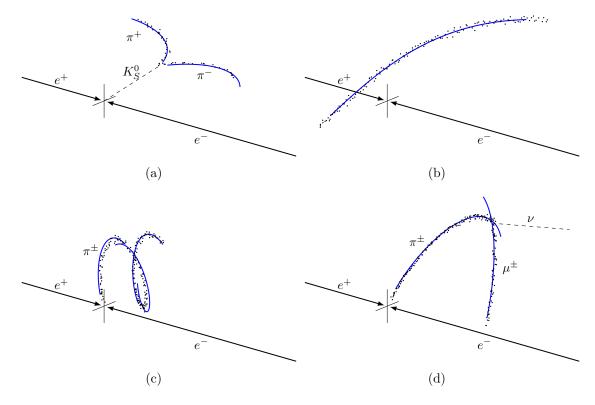


Figure 5.5.: (a) Tracks from long-lived neutral particles, which decay away from the IP region, (b) random reconstructed background tracks, (c) low-momentum particles, which curl in the magnetic field, (d) in-flight decays of particles, which produce a "kink" in the trajectory.

It is obvious that tracks from the same momentum source should only be taken into account once, or, in the case of background tracks, not at all. Such tracks will, from this point on, be denoted as *extra* tracks, because they add extra four-momentum to our final calculations in Eq. (4.8). At the same time, we have to take care that we do not identify *good* tracks as *extra* tracks. Both of these cases have negative impacts on the final resolution of all variables, which depend on information from ROE.

5.3.1. Tracks from Long-lived Particles

The first step in the tracks clean-up is taking care of the tracks from long-lived particles, such as $K_S^0 \to \pi^+\pi^-$, $\gamma \to e^+e^-$ and decays of Λ baryons. Here we only focus on K_S^0 , since they are the most abundant. This step is necessary because of the π^{\pm} particles, coming from the K_S^0 decays, have large impact parameters, which is usually a trait of background particles. In order to minimize confusion from the MVA point-of-view, these tracks are taken into account separately.

We use the converted K_S^0 candidates from the existing Belle particle list and use a pre-trained Neural Network classifier in order to select only the good K_S^0 candidates. Figure 5.6 shows the distribution of the K_S^0 invariant mass for signal and background candidates, before and after the selection cut on the classifier output. The momentum of selected K_S^0 candidates is added to the ROE, while the daughter tracks are discarded from our set.

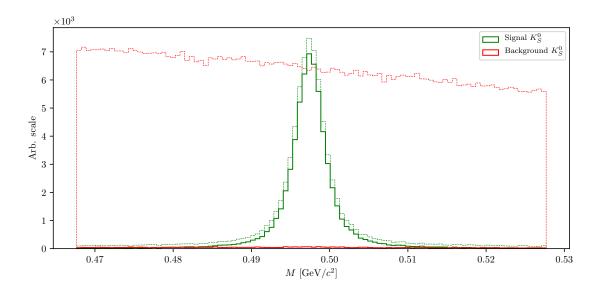


Figure 5.6.: Invariant mass of the K_S^0 candidates before (dashed) and after (solid) the selection on the Neural Network classifier for signal (green) and background candidates (red). Signal peaks at nominal K_S^0 mass, while background covers a wider region.

The signal efficiency and background rejection for K_S^0 candidates after this selection and on the full range are

• Signal efficiency: $\epsilon_{SIG} = 80.7 \%$,

• Background rejection: $1 - \epsilon_{BKG} = 99.4 \%$.

5.3.2. Duplicate Tracks

All good tracks at this point should be coming from the IP region, since we took care of all the good tracks from long-lived particle decays, therefore we apply a selection on impact parameters for all the remaining tracks

• $|d_0| < 10$ cm and $|z_0| < 20$ cm

and proceed with the clean-up of track duplicates.

Defining a duplicate track pair

In this step, we wish to find a handle on secondary tracks from low momentum curlers and decays in flight. The main property for these cases is that the 3D opening angle between such two tracks is very close to 0° or 180°, since the tracks deviate only slightly from the initial direction, but can also be reconstructed in the opposite way. Figure 5.7 shows the distribution of the angle between two tracks in a single pair for random track pairs and duplicate track pairs, where the latter were reconstructed as two same-sign or opposite-sign tracks.

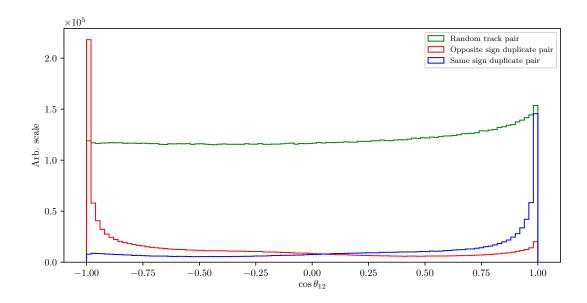


Figure 5.7.: Distribution of the angle between two tracks in a single pair for random track pairs (green) and duplicate track pairs, where the latter were reconstructed as two samesign (blue) or opposite-sign tracks (red).

If the particle decayed mid-flight or produced multiple tracks due to being a low-momentum curler, then, as the name suggests, these particles most likely a had low momentum in the transverse direction, p_T . Since both tracks originate from the same initial particle, the momentum difference should also peak at small values. Figure 5.8 shows the momentum and momentum difference of tracks which belong to a random or a duplicate track pair.

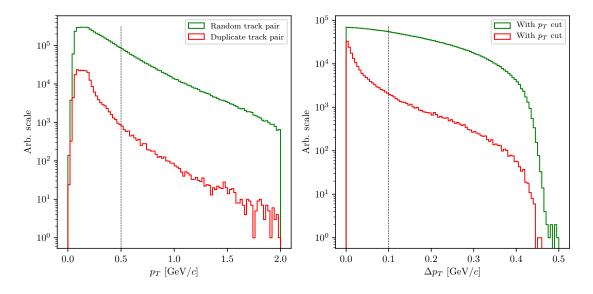


Figure 5.8.: Distribution of transverse momentum p_T (left) and transverse momentum difference Δp_T (right) for all tracks coming from random (green) or duplicate track pairs (red). The plot on the right already includes the selection on p_T from the plot on the left.

We impose a selection of

- $p_T < 0.5 \text{ GeV}/c$,
- $|\Delta p_T| < 0.1 \text{ GeV}/c$,

in order to minimize the number of random track pairs, while retaining a high percentage of duplicate track pairs. After applying the selection criteria defined in this chapter, the final distribution of the angle between two tracks is shown in Figure 5.9.

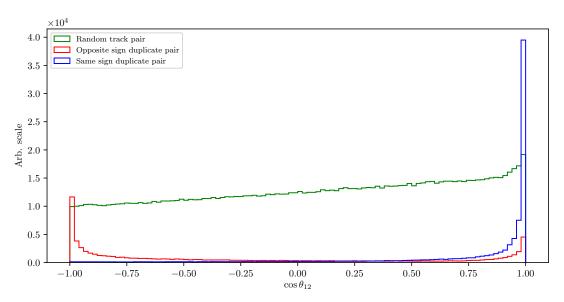


Figure 5.9.: Distribution of the angle between two tracks in a single pair after applying the selection defined in this section. The distributions are shown for random track pairs (green) and duplicate track pairs, where the latter were reconstructed as two same-sign (blue) or opposite-sign tracks (red).

Training the duplicate track pair MVA

This final sample of track pairs is used in the MVA training to recognize duplicate track pairs. The training dataset contains

- 113707 target candidates,
- 190314 background candidates,

where the definition of a target is that the track pair is a duplicate track pair. The input variables used in this MVA are

- angle between tracks,
- track quantities
 - impact parameters d_0 and z_0 ,
 - transverse momentum p_T ,

- helix parameters and helix parameter errors of the track,
- track fit *p*-value,
- number of hits in the SVD and CDC detectors

The classifier is able to distinguish between random and duplicate track pairs in a very efficient manner, as shown in Figure 5.10.

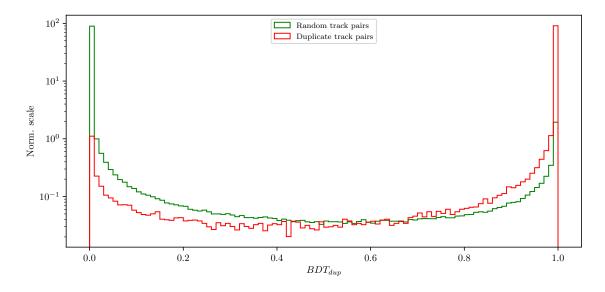


Figure 5.10.: Classifier output of the track pair training for random track pairs and duplicate track pairs.

The FOM function for optimal selection is shown in Figure 5.11 (left), along with the angle between the two tracks before and after the optimal selection (right). The optimal duplicate track selection is

• $BDT_{duplicate} > 0.9985$.

The signal efficiency and background rejection for duplicate pair candidates after this selection is

- Signal efficiency: $\epsilon_{SIG} = 87.2 \%$,
- Background rejection: $1 \epsilon_{BKG} = 98.8 \%$,

where signal and background represent duplicate and random track pairs, respectively.

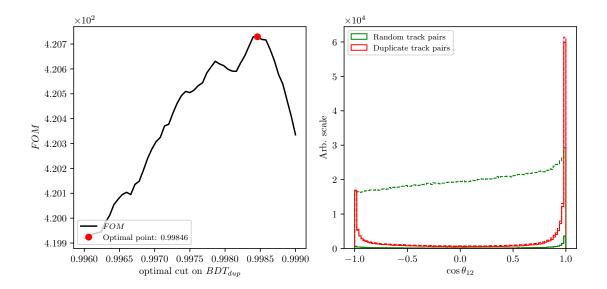


Figure 5.11.: The optimization of the *FOM* function for the cut on classifier output (left) and distribution of the angle between two tracks in a single pair before (dashed) and after (solid) applying the optimal cut on the output classifier for random and duplicate track pairs (right).

Defining duplicate tracks

What remains now is to decide which track from the duplicate track pair to keep and which to discard. For this purpose, we apply duplicate pair-level information to each track in the pair in the form of

$$\Delta f = f_{this} - f_{other},\tag{5.1}$$

where f is an arbitrary variable from the list of track quantities in Section 5.3.2. From the point-of-view of *this* track, a track is more *duplicate*-like if the following is true

- $\Delta d_0, \Delta z_0 > 0$ (this track further away from the IP region),
- $\Delta p_T, \Delta p_Z < 0$ (this track has lower momentum),
- ΔN_{SVD} , $\Delta N_{CDC} < 0$ (this track has less hits in the SVD and CDC),

Additionally we define an MC truth variable

$$\Delta \chi^2 = \chi_{this}^2 - \chi_{other}^2, \quad \chi^2 = \sum_{i=x,y,z} \frac{(p_i - p_i^{MC})^2}{\sigma(p_i)^2},$$
 (5.2)

where we compare all components of track momentum to the true values. If the condition $\Delta \chi^2 > 0$ is satisfied, then *this* track is probably a duplicate track and should be discarded.

However, it turns out that solving this problem is not as simple as discarding one track and keeping the other one. An additional complication is that we can have

more than one extra track from the same initial particle, which leads to track pairs, where both tracks are track duplicates. For example, if we have the following case

```
t_1: good track,

t_2: extra track,

t_3: extra track,

pair<sub>1</sub>: (t_1,t_2),

pair<sub>2</sub>: (t_1,t_3),

pair<sub>3</sub>: (t_2,t_3),
```

where t_1 is the original track and t_2 and t_3 are extra tracks, with t_3 being even more duplicate-like with respect to t_2 . Here tracks t_2 and t_3 should be discarded while t_1 should be kept. We can achieve this, if we overwrite existing pair-level information in the tracks for cases, where the variable difference Δf is more duplicate-like. If we follow the same example, we could fill information about the property f in six different orders.

```
1. (t_1,t_2*) \rightarrow (t_1,t_3*) \rightarrow (t_2*,t_3*),

2. (t_1,t_2*) \rightarrow (t_2*,t_3*) \rightarrow (t_1,t_3*),

3. (t_1,t_3*) \rightarrow (t_2,t_3*) \rightarrow (t_1,t_2*),

4. (t_1,t_3*) \rightarrow (t_1,t_2*) \rightarrow (t_2*,t_3*),

5. (t_2,t_3*) \rightarrow (t_1,t_3*) \rightarrow (t_1,t_2*),

6. (t_2,t_3*) \rightarrow (t_1,t_2*) \rightarrow (t_1,t_3*),
```

where the "*" symbol denotes when a track is recognized as a duplicate track with respect to the other track. We see that no matter the order, both t_2 and t_3 get recognized as duplicate tracks correctly.

Training the duplicate track MVA

The training procedure is similar as before. The sample of tracks from duplicate track pairs is now used in the MVA training to distinguish duplicate tracks from good tracks. The training dataset contains

- 84339 target candidates,
- 68280 background candidates,

where the definition of a target is that the track is a duplicate track, based on the $\Delta \chi^2 > 0$ condition.

The input variables used in this MVA are

- theta angle of the track momentum,
- track quantities
 - impact parameters d_0 and z_0 and their errors,
 - CMS frame momentum p_{CMS} and momentum components p_T and p_z
 - number of hits in the SVD and CDC detectors

- track fit *p*-value,
- pair-level information
 - $-\Delta d_0, \Delta z_0, \Delta N_{CDC}, \Delta N_{SVD}, \Delta p_T, \Delta p_z, \Delta p$ -value.

The output classifier information is added to the tracks, where now each track has a certain probability of being a duplicate track. We then compare these values between both tracks in each track pair as

$$\Delta BDT_{final} = BDT_{final}^{this} - BDT_{final}^{other}, \tag{5.3}$$

which is again applied to all track pairs and overwritten for tracks which are more duplicate-like. The classifier output and the classifier output difference for each track are shown in Figure 5.12.

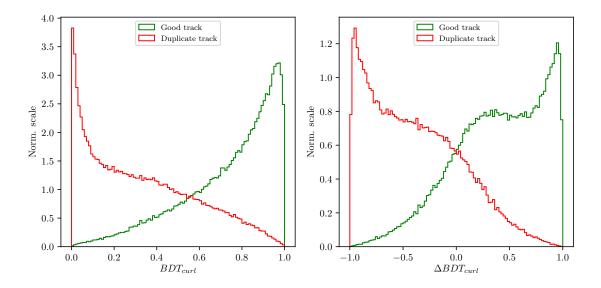


Figure 5.12.: Classifier output of the MVA training for curling track recognition (left) and difference of the classifier output, calculated for each track in a track pair (right).

Finally, we select all duplicate tracks, which survive the selection

$$\Delta BDT_{final} > 0,$$
 (5.4)

and discard them from our ROE. We can check the performance of our duplicate track classifier by applying the procedure to a validation sample of duplicate track pairs and compare the predicted result with the truth, based on Eq. (5.2). Table 5.1 shows the performance of the duplicate track recognition in the form of percentages of correctly and incorrectly identified duplicate and original tracks. The model seems to perform well and the event is now considered to be clean of duplicate tracks.

	Predicted duplicate track	Predicted good track
Duplicate track Good track	$\begin{bmatrix} 83.07 \% \\ 16.93 \% \end{bmatrix}$	22.62 % 77.38%

Table 5.1.: Ratios of correctly classified and misclassified tracks.

5.4. Belle Clean-up

For comparison, we define the Belle clean-up, used standardly at Belle, which is much simpler and relies only on a set of basic selection criteria for neutral and charged particles. This clean-up procedure is not applied in addition to our ROE clean-up, but separately, only for comparison.

In the case of photons, only a selection on the photon energy is applied, depending on the region where the photon hit the relevant part of the detector. The photon selection is summarized in Table 5.2.

Table 5.2.: Photon selection for the Belle clean-up procedure. Different selection criteria are applied on photons in different parts of the detector.

In case of tracks, pairs are selected which satisfy the following criteria:

- $p_T < 275 \text{ MeV}/c$,
- $\Delta p = |\vec{p_1} \vec{p_2}| < 100 \text{ MeV}/c$,
- $\cos \theta(\vec{p}_1, \vec{p}_2) < 15^{\circ}$ for same sign,
- $\cos \theta(\vec{p_1}, \vec{p_2}) > 165^{\circ}$ for opposite sign.

Of the two tracks, the one with a larger value of formula in Eq. 5.5 is discarded. The remaining tracks in the event then need to satisfy the conditions described in Table 5.3.

$$(\gamma |d_0|)^2 + |z_0|^2, \quad \gamma = 5.$$
 (5.5)

	$p_T < 250 \text{ MeV}/c$	$250 \text{ MeV}/c < p_T < 500 \text{ MeV}/c$	$p_T > 500 \text{ MeV}/c$
$ d_0 $	< 20 cm	< 15 cm	< 10 cm
$ z_0 $	< 100 cm	< 50 cm	< 20 cm

Table 5.3.: Track selection for the Belle clean-up procedure. Different selection criteria are applied to tracks in different p_T regions.

5.5. Clean-up Results

In this section, the results of the ROE clean-up are shown. It is obvious that cleaning up the event affects the shape of various distributions, especially ΔE and M_{BC} , which we are most interested in. Since the reconstruction procedure includes applying selection criteria on the cleaned-up variables, the clean-up also affects the efficiency of the reconstructed sample, not only the resolution.

We compare the clean-up setup, defined in this analysis, to the standard clean-up used by Belle, and to a default case, where no clean-up was applied at all. We

apply the clean-up procedure to our signal MC sample with all the applied selection criteria, defined in Section 4.6, except for the signal categorization. Figure 5.13 (left) shows signal candidate distributions of ΔE and M_{BC} for various clean-up setups. Focusing on the ROE clean-up, we see an improvement in resolution in both observed variables and an overall decrease in efficiency. The efficiency decrease is expected since the cleaned-up variables are able to better isolate the perfectly reconstructed candidates and discard the non-perfect candidates. In fact, the efficiency of the perfectly reconstructed candidates increases after the ROE clean-up, as shown in Figure 5.13 (right). The signal MC sample in case of the Belle clean-up also shows a slight improvement in the resolution after the procedure, but looking at the perfectly reconstructed candidates, we see that this clean-up procedure is not optimal. Table 5.4 shows ratios of efficiencies and FWHM's of the clean-up procedures for the perfect signal with respect to the default case, based on the ΔE distribution. While both, the Belle and ROE clean-up, improve the resolution, ROE clean-up performs significantly better and also increases the amount of the perfectly reconstructed candidates in the final sample.

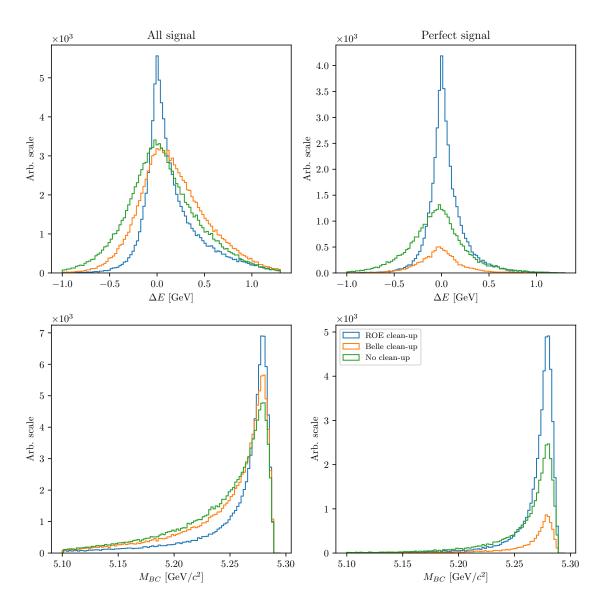


Figure 5.13.: ΔE and M_{BC} distributions for various types of cleanup procedures. The figures on the left are shown for the full signal sample after the stated selection criteria, while the figures on the right are shown for the perfectly reconstructed signal candidates. For ROE clean-up, the procedure seems to improve resolution, as well as increase the amount of perfectly reconstructed candidates, relative to the default case.

	Efficiency ratio	FWHM ratio
Belle clean-up ROE clean-up	28.5 % 140.1 %	$75.0 \% \\ 35.0 \%$

Table 5.4.: Comparison of efficiencies and FWHM's of ROE and Belle clean-up setups with respect to the default case (no clean-up) for the perfect signal.

Another variable which heavily depends on the clean-up is the charge product of

the signal and companion B meson candidate, already defined in Eq. (4.18), shown in Figure 5.14 for various clean-up procedures. The figure shows an improved resolution of the distribution of the charged product, which means that candidates migrate to the correct value of the charge product after the clean-up. Looking at the perfectly reconstructed candidates, we again see the increase in the bin corresponding to the correct charge product. As a cross-check, we can also look at ΔE and M_{BC} variables for each value of the charge product. These plots are shown for the full signal MC sample in Figure 5.15 and they show a clear resolution improvement for the correct value of the charge product in the case of the ROE clean-up. For other values of the charge product there also seems to be a small improvement for both cases of clean-up, but it is negligible compared to the plots for the optimal charge product value. This supports our choice of signal categorization, defined in Section 4.5, where we select only candidates with the correct value of the charge product.

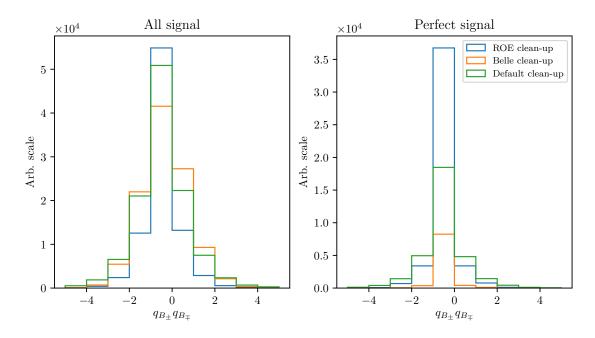


Figure 5.14.: Distribution of the charge product of both B mesons for various types of clean-up procedures, shown on the full signal MC (left) and for the perfectly reconstructed signal candidates (right). For ROE clean-up, the procedure seems to increase the number of perfectly reconstructed candidates.

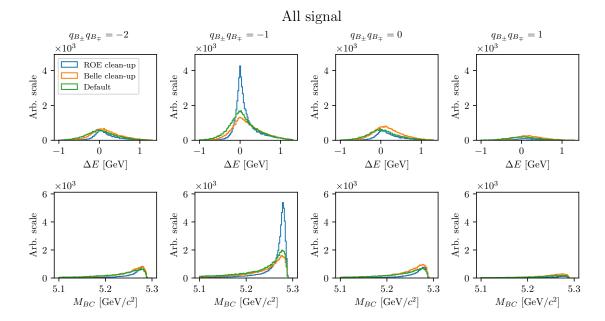


Figure 5.15.: Distributions of ΔE (top) and M_{BC} (bottom) for various types of clean-up procedures, split by specific values of the charge product, shown for the full signal MC. There is a significant improvement in resolution after ROE-cleanup, for the case of the correct value of the charge product.

5.6. ROE Clean-up Validation

The ROE clean-up seems to perform well on signal MC, based on the results in the previous section. However, it is necessary to make sure that this procedure performs as well on other simulated and measured data, which is done in this section. The clean-up procedure is validated on the control sample,

$$B^+ \to \bar{D}^0 \ell^+ \nu, \quad D^0 \to K^+ K^-,$$

which was already defined in Section 2.2. The control candidates are reconstructed in the same manner as the signal candidates. In addition to the same selection criteria applied, as in the previous section, we also apply a selection to make the control sample more significant. We keep only the candidates passing the following selection cut on the invariant mass of the two kaons

$$1.849 \text{ GeV}/c^2 < m_{KK} < 1.879 \text{ GeV}/c^2$$
 (5.6)

as shown in Figure 5.16. Further detail about this cut can be found in Section 6.1.

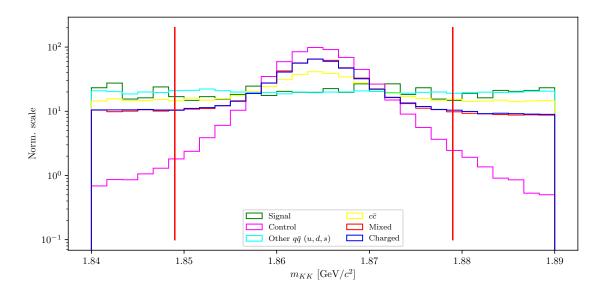


Figure 5.16.: Normalized distributions of m_{KK} for the full MC dataset. The red lines represent the edges of the selection in the m_{KK} distribution where the control sample is enhanced. The m_{KK} distribution drops quickly for the case of the control decay, while staying uniform for other contributions.

With the control sample selection determined, we now run the reconstruction with and without the ROE clean-up procedure on all available MC and data.

The effects of the ROE clean-up are shown in Figure 5.17, where we overlay the data points to a stacked histogram of MC contributions for ΔE and M_{BC} . We see that the data and MC agree well. A slight systematic trend can be seen in the ΔE variable, which is addressed in Section 7.1.1. The control sample resolution seems very poor in the case without the clean-up, but it improves significantly if the clean-up procedure is applied, as expected. The simulated background also seems to gain an improvement in the resolution, but this is likely due to the background consisting of similar candidates as the control sample. This means that the clean-up performs as expected due to the nature of the decays and does not arbitrarily shape the background to be more signal like. Additionally, it should be pointed out that, after the clean-up, the simulated background resolution is worse compared to the control decay resolution, while this is not the case if the clean-up procedure is not performed.

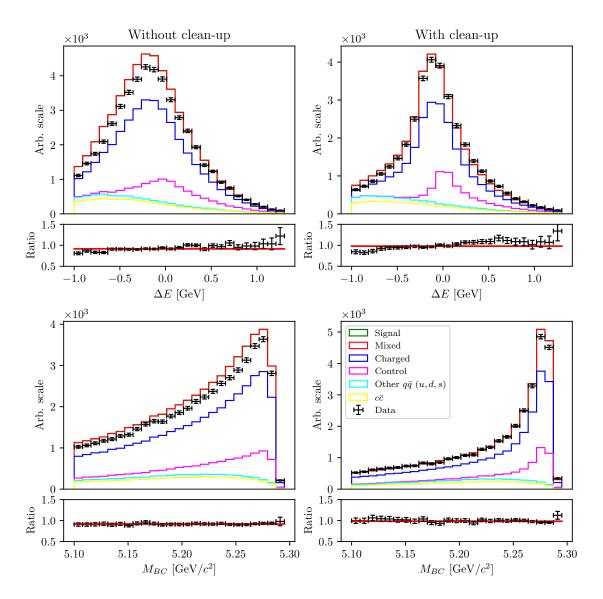


Figure 5.17.: Distributions of ΔE (top) and M_{BC} bottom for the case without (left) and with ROE clean-up (right). The resolution of the control sample is improved and the MC and data agree well in all aspects. While the simulated background resolution is also improved, it is worse compared to the resolution of the control sample.

To perform the clean-up validation in greater detail, we also compare the data and MC agreement in bins of the charge product of the two B mesons. Figure 5.18 shows the cleaned-up versions of ΔE and M_{BC} for each charge product bin in the same manner as shown in the previous section. We see that the MC and data agreement persists in all cases.

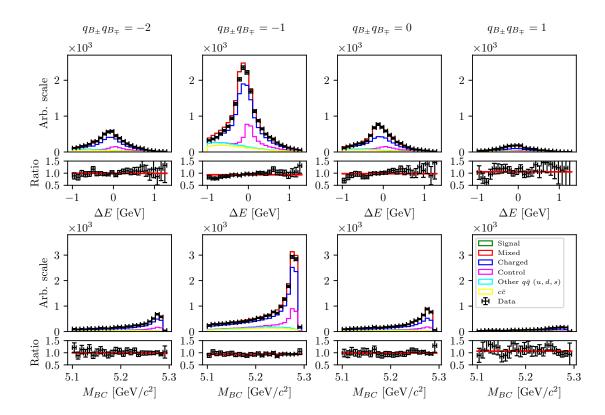


Figure 5.18.: Distributions of ΔE (top) and M_{BC} (bottom) split in bins of the charge product of the two B mesons.

The ROE clean-up procedure seems to perform well. It significantly improves the resolution of the control and, therefore, signal candidates and increases the amount of perfectly reconstructed events. The clean-up procedure was also applied to data and no disagreement with respect to the simulated MC samples was found. This means that the procedure does not differ between MC and data and does not perform on them differently. The procedure was therefore validated in great detail and is suitable to be used in this analysis.

Chapter 6.

Background Suppression

This chapter explains the procedure of suppressing various kinds of backgrounds by using MVA classifiers. More information about the MVA training, feature importance, and hyper-parameter optimization for each MVA step in this chapter can be found in Appendix B.

6.1. Resonant Background

In the analysis we study decays with kaons in the final state. This means that standard procedures in $b \to u$ analyses, in order to suppress $b \to c$ backgrounds, such as K-veto, are not possible. As a consequence, our final sample consists of combinations of K pairs coming also from $b \to c$ sources, such as $D^0 \to K^+K^-$. Such candidates usually have resonance-like properties in the two-kaon invariant mass spectrum. Figure 6.1 shows this invariant mass spectrum of two kaons, m_{KK} , where obvious resonant structures are present from sources like

- $\phi \to K^+K^-$ (sharp peak at $\sim 1.019 \text{ GeV}/c^2$),
- $D^0 \to K^+K^-$ (sharp peak at $\sim 1.864 \text{ GeV}/c^2$),
- $D^0 \to K^+\pi^-$ (wide, shifted peak, due to kaon miss-identification).

In order to suppress these resonant backgrounds, while studying signal or control decays, we define two regions

- signal region: $|m_{KK} m_{\phi}| > \Delta_{\phi}, |m_{KK} m_{D^0}| > \Delta_{D^0}, |m_{K\pi} m_{D^0}| > \Delta_{D^0},$
- control region: $|m_{KK} m_{D^0}| \le \Delta_{D^0}, |m_{K\pi} m_{D^0}| > \Delta_{D^0},$

where m_{KK} is the KK invariant mass and $m_{K\pi}$ is the invariant mass of KK candidates, where the kaon with the same charge as the B meson was given the mass of the charged π . $m_{\phi} \approx 1.019 \text{ GeV}/c^2$ and $m_{D^0} \approx 1.864 \text{ GeV}/c^2$ are nominal masses of the ϕ and D^0 mesons, and $\Delta_{\phi} \approx 8 \times 10^{-3} \text{ GeV}/c^2$ and $\Delta_{D^0} \approx 1.5 \times 10^{-2} \text{ GeV}/c^2$ are the widths around the nominal mass values for the ϕ and D^0 meson, respectively. By selecting the signal or control region, we are able to efficiently isolate the desired subset. Table 6.1 shows the subsample efficiency after selecting either of the regions.

	ϵ (Signal cand.)	ϵ (Control cand.)	$\epsilon(\phi \text{ resonance cand.})$
Signal-specific	95.4%	4.0%	$\begin{array}{ c c c c }\hline 13.6\% \\ 0.0\% \\ \end{array}$
Control-specific	1.9%	96.0%	

Table 6.1.: Efficiencies after selecting the signal or control region

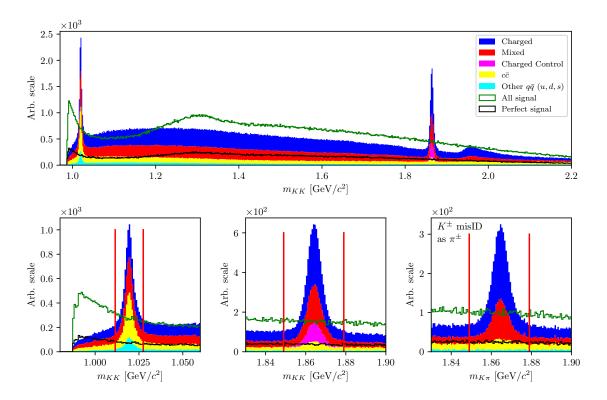


Figure 6.1.: Invariant mass of KK candidates over a wider region (top) and for a specific region around the ϕ peak (bottom left), the D^0 peak (bottom center), and again for the D^0 peak, but with the $m_{K\pi}$ mass of the KK candidates (bottom right). Signal (green) and perfect signal (black) are equally scaled up.

6.2. Continuum Suppression

Physics processes where continuum states are produced in electron and positron collisions

$$e^+e^- \to q\bar{q}$$

with q=u,d,s or c, represent an important category of backgrounds, called the continuum background. In addition to the previously used kinematic constraints, used to separate $e^+e^- \to \Upsilon(4S) \to B\bar{B}$ decays from $e^+e^- \to q\bar{q}$, properties of the event shape" are also used, since phase-space distributions of produced particles differ for these two processes. Continuum background events are produced back-to-back in the CMS frame, so hadrons produced in the quark fragmentation possess only a small transverse momentum compared to the initial momentum magnitude. This leads to a spatially confined, jet-like structure. On the other hand, B mesons

from $B\bar{B}$ events are produced almost at rest in the CMS frame. Their decay products form an isotropic distribution in the detector, which yields a spherical event shape.

6.2.1. Characteristic Variables

Information on the phase-space distribution of produced particles can be obtained in a number of different ways. In this section, different characteristic variables are presented, which are used in the MVA training. They all focus on kinematic and shape differences between the two processes, which we wish to discriminate.

Thrust and Related Variables

It is possible to define a thrust axis \vec{T} for a collection of N momenta p_i as a unit vector, along which their total projection is maximal. The thrust scalar T (or thrust) is a derived quantity, defined as

$$T = \frac{\sum_{i} |\vec{T} \cdot \vec{p_i}|}{\sum_{i} |\vec{p_i}|}.$$
(6.1)

In this case, a related variable is $|\cos\theta_T|$, where θ_T is the angle between the thrust axis of the particles from the B meson candidate and the thrust axis of all particles in the ROE. Since both B mesons in $B\bar{B}$ events are produced at rest, their decay particles and, consequentially, their thrust axes are isotropically distributed. On the other hand, particles in continuum events follow the direction of the jets in the event. As a consequence, both thrust axes are strongly directional and aligned in the opposite direction, which results in a large peak at $|\cos\theta_T| \approx 1$. Additionally, one can also use the variable $|\cos\theta_{TB}|$, which is the cosine of the angle between the thrust axis of the B candidate and the beam axis. For B candidates from $B\bar{B}$ events this distribution is again uniformly distributed, while for candidates from continuum events this distribution follows the distribution of the initially produced quark pairs, $1 + \cos^2\theta_{T,B}$. In practice, such a distribution exhibits a drop at $|\cos\theta_{TB}| \approx 1$, due to the acceptance loss of the detector in the direction of the beam pipes. Figure 6.2 shows the distributions of $|\cos\theta_T|$ (left) and $|\cos\theta_{T,B}|$ (right) for B meson candidates from various sources.

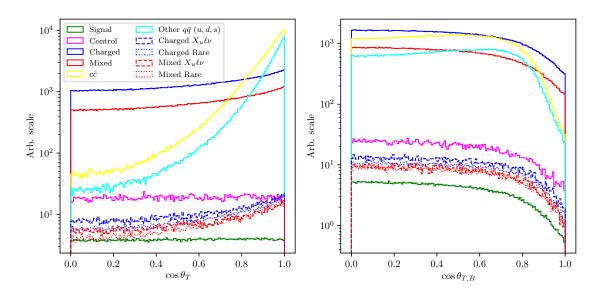


Figure 6.2.: Distributions of $|\cos \theta_T|$ (left) and $|\cos \theta_{T,B}|$ (right) for B meson candidates from various sources.

CLEO Cones

CLEO cones have been introduced by the CLEO collaboration and are an additional specific tool to provide optimal continuum background discrimination. They consist of nine variables corresponding to the momentum flow around the thrust axis of the B meson candidate, binned in nine cones of 10° around the thrust axis, as illustrated in Figure 6.3. Momentum flow is defined as the scalar sum of the momenta of all charged tracks and neutral particles, pointing in a specific angle interval. A Fisher discriminant is formed from the nine momentum flow variables and from $|\cos \theta_T|$ for B candidate and the ROE. The Fisher discriminant, \mathcal{F} , is the linear combination of the input variables, which maximizes the separation between signal and background. Additional information is provided in [20].

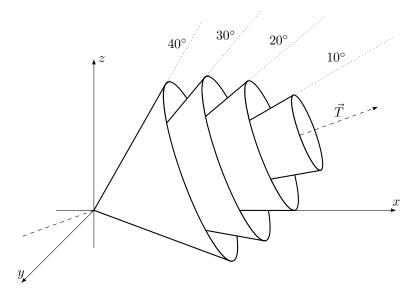


Figure 6.3.: Concept of CLEO cones. \vec{T} denotes the thrust axis of the B meson candidate in an event. Each variable corresponds to a momentum flow around the thrust axis in steps of 10° .

KSFW Moments

Fox-Wolfram moments are another useful parametrization of phase-space distribution of energy and momentum flow in an event. For a collection of N momenta p_i , the k-th order normalized Fox-Wolfram moment R_k is defined as

$$R_k = \frac{H_k}{H_0} = \frac{1}{H_0} \sum_{i,j} |p_i| |p_j| P_k(\cos \theta_{ij}), \tag{6.2}$$

where θ_{ij} is the angle between p_i and p_j , and P_k is the k-th order Legendre polynomial. For events with two strongly collimated jets, R_k takes values close to 0 (1) for odd (even) values of k, so these moments provide a convenient discrimination between $B\bar{B}$ and continuum events.

Belle developed a refined generation of Fox-Wolfram moments, called Kakuno-Super-Fox-Wolfram (KSFW) moments to further suppress the continuum background. They are described in detail in [21].

B-Flavor Tagging

While the goal of B-flavor tagging is to determine the flavor of a B meson, the variables used for flavor tagging also potentially contribute to background suppression. Flavor tagging relies on the fact that a large fraction of B mesons decay to a final state that is flavor specific, and can only be reached either through the decay of a b or a \bar{b} quark. Because of the large number of B meson decay channels, full reconstruction of a sufficiently large number of flavor-specific B candidates is not feasible. Instead, inclusive techniques are employed, that make use of different flavor-specific signatures of B decays.

The flavor tagging algorithm proceeds in two stages. In the first stage, individual flavor-specific signatures are analyzed, each of which provides a signature-specific

flavor tag. In the second stage, the results from the first stage signatures are combined into a final flavor tag. Both stages heavily rely on MVA methods in order to optimally combine all available information. The final result of each stage is qp, a product of the flavor sign, q, and the probability of a correct flavor tag, p. Further details can be found in [21].

ROE fit information

Most B tag mesons decay via $b \to c$ transitions with at least one additional vertex at a distance comparable to the decay length of a B meson. These vertices introduce a bias in the measurement of the companion B meson vertex position, which degrades the vertex resolution.

The strategy to select the optimal set of tracks for the vertex determination is to first select a subset of the tracks in the ROE, which satisfy some requirements, such as a minimum number of vertex detector hits and a maximum transverse distance to the interaction region. All tracks, which do not pass the selection criteria, are removed. In the end, all tracks are combined in a single vertex using the interaction region as a constraint. If the goodness of the vertex fit is not good enough, the worst track is removed and the vertex refitted. This procedure is repeated until the fit is satisfactory or no tracks are left. After the vertex of the ROE has been determined, Δz can be calculated as the distance of vertices between the signal B meson candidate and the ROE in the z direction. More information is available in [21].

6.2.2. MVA Training

Most of the characteristic variables, described in Section 6.2.1, were taken together in order to train a single MVA classifier for continuum suppression. All characteristic variables were checked for possible q^2 , M_{BC} or ΔE correlation. Variables with significant correlation or complex shapes in the correlation distribution were discarded from the training set, since they would have introduced unwanted dependence on the unreliable model, used for signal MC generation. Additionally, all of the characteristic variables in our set do not depend on the signal mode, they only differ in the kinematic and topological aspects of $B\bar{B}$ and continuum background events.

The training dataset consisted of 2×10^5 candidates, where 50 % of the candidates are correctly reconstructed signal events, 25 % are $u\bar{u}$, $d\bar{d}$ and $s\bar{s}$ background with expected proportions, and 25 % is $c\bar{c}$ background. Such a composition is chosen so that there are enough signal and background samples for the MVA model training, and to avoid any of the background contributions being under-represented. Since the full Belle dataset is experiment-dependent, we construct the training dataset by sampling MC events from appropriate experiments, proportionally to the size of the dataset per each experiment.

The training variable set consisted of

- B meson direction and thrust related variables
 - magnitude of thrust axes of the B and ROE candidates,
 - cosine of the angle between the thrust axis of the B candidate and thrust axis of the ROE candidate,

- cosine of the angle between the thrust axis of the B candidate and the beam direction,
- reduced Fox-Wolfram moment R_2 ,
- all 9 CLEO Cones
- KSFW Moments
 - $-R_{01}^{so}, R_{02}^{so}, R_{03}^{so}, R_{04}^{so},$
 - $-R_{10}^{so}, R_{12}^{so}, R_{14}^{so},$
 - $-R_{20}^{so}, R_{22}^{so}, R_{24}^{so},$
 - $-R_0^{oo}, R_1^{oo}, R_2^{oo}, R_3^{oo}, R_4^{oo},$
- B-flavor tagging variables
 - $-qp ext{ of } e, \mu, \ell,$
 - qp of intermediate e, μ, ℓ ,
 - qp of K, K/π , slow pion, fast hadron,
 - -qp of maximum P^* , Λ , fast-slow-correlated (FSC),
- Other
 - $-\Delta z$.

Figure 6.4 shows the classifier output for various types of background, all in expected MC proportions. B meson candidates from continuum background are dominant at lower values, while candidates from $B\bar{B}$ events populate the region with higher values.

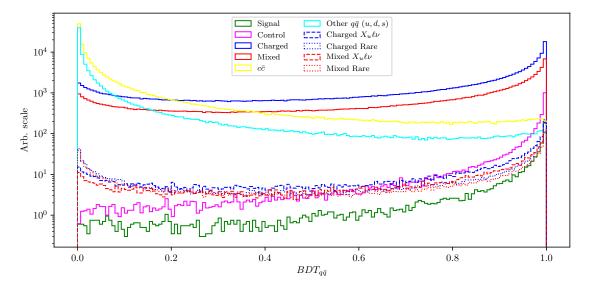


Figure 6.4.: Continuum suppression classifier output for signal and various types of background. B candidates from continuum events dominate the lower region, while candidates from $B\bar{B}$ dominate in the upper region of the classifier output.

6.3. $B\bar{B}$ Suppression

After separating continuum background from $B\bar{B}$ events, the next step is to train an MVA classifier to recognize our signal candidates among the candidates from other $B\bar{B}$ events. $B\bar{B}$ events consists of

- $b \to c\ell\nu$ background,
- $b \to u\ell\nu$ background,
- Other rare decays (radiative, penguin, rare 2- and 3-body decays, ...).

Similarly, the training dataset for this classifier consisted of 2×10^5 candidates, where 50 % of the candidates are correctly reconstructed signal events. The remaining part of the training dataset consists of all background, not including the control sample, because we are not interested in suppressing it directly. The background part of the dataset consists of 75 % generically decaying charged and neutral $B\bar{B}$ events in equal proportions, whereas the remaining 25 % is equally populated with charged and neutral $B\bar{B}$ events from $b \to u\ell\nu$ and other rare decays. The training dataset was proportionally sampled in the same manner as described in Section 6.2.2.

In order to separate this kind of background, we must be careful not to introduce correlations with the fit variables (ΔE and M_{BC}) or any kind of model dependence (correlation with q^2). This means that we can not use any information of the decay particles or the candidate, which is of kinematic nature, such as decay particles momenta, decay angles or other similar variables.

The training variable set consisted of

- vertex fit probability of $P(\chi^2, DOF)$ of the signal B meson candidate
- vertex fit probability of $P(\chi^2, DOF)$ of the ROE side,
- $\cos \theta_{BY}$ from Eq. (4.4),
- cos of the angle between the momentum vector and vector joining the IP and the production vertex of the $KK\ell$ candidate,
- B-flavor tagging variables for the two signal-side kaons,
- numbers of kaons, tracks and distant tracks in ROE,
- θ angle of the ROE momentum in CMS frame,
- ξ_Z from [22]
- $\bullet \Delta z$,
- m_{miss}^2 from Eq. (4.11),
- m_{miss}^2 for partial reconstruction of $B^0 \to D^{*-}\ell\nu$,

where distant tracks are all tracks in ROE which satisfy the condition of $|d_0| > 10.0$ cm or $|z_0| > 20.0$ cm. The last entry is a veto variable where we partially reconstruct the D^* candidate four-momentum via a linear combination of the π_s^{\pm} four-momentum in the $D^* \to D\pi_s^{\pm}$ decay. It contributes to discarding the $B^0 \to D^{*-}\ell^+\nu$ decay, where D^{*-} further decays to $D^{*-} \to \bar{D}^0\pi_s^-$. Figure 6.5 shows the veto variable with a partial reconstruction of a charged π_s^{\pm} .

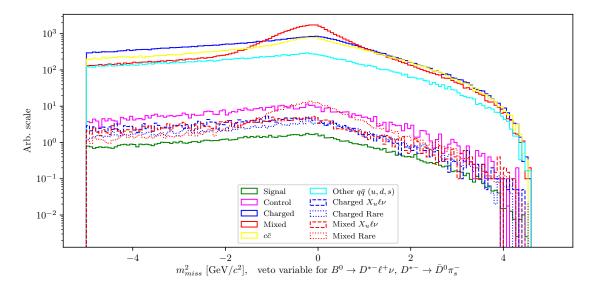


Figure 6.5.: Distribution of m_{miss}^2 for partially reconstructed $B^0 \to D^{*-}\ell^+\nu$ decays.

When the training is finished and the hyper-parameters of the classifier are optimized, the classifier output, as shown in Figure 6.6 (left), can be used for background suppression. B meson candidates from $B\bar{B}$ background are dominant at lower values, while candidates from $B\bar{B}$ events populate the region with higher values. Since the differences between signal and background $B\bar{B}$ events are smaller than $B\bar{B}$ and $q\bar{q}$ events, the resulting classifier has a smaller separation power than the one described in the previous section.

6.3.1. Boosting to Uniformity

The selection approach with standard classifiers is optimal for counting experiments, as it, by construction, produces the optimal selection for observing an excess of signal over background events. Today's BDT algorithms, which work in this way, produce non-uniform selection efficiencies and may, as a consequence, shape background distributions to look like signal. In order to minimize such behavior, it is possible to discard variables, which are correlated with the variable of interest (in our case ΔE and M_{BC}), from the training set. This, however, decreases the classifiers discriminating power. Another approach is to use a novel boosting method, uBoost, which is trained to optimize an integrated FOM under the constraint that the BDT selection efficiency for the desired class must be uniform. The uBoost algorithm balances the biases to produce the optimal uniform selection [23].

The training set used is the same as described at the beginning of this chapter, along with the same set of training variables. It will be seen later that the standard

BDT classifier shapes the background to look like signal mostly in the M_{BC} distribution, therefore we train the uBDT classifier with a uniformity constraint on the M_{BC} variable of the background candidates. The resulting classifier output is shown in Figure 6.6 (right). For this classifier, the separation power between signal and background seems worse, however, the shapes of backgrounds differ significantly, which greatly aids in the performance of signal extraction.

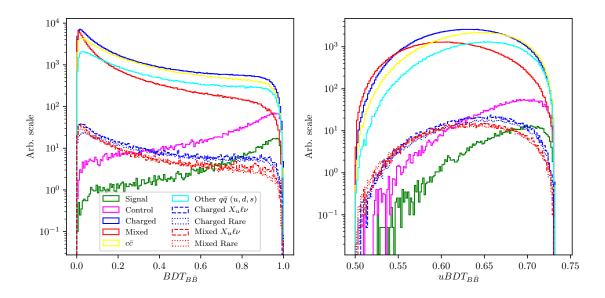


Figure 6.6.: $B\bar{B}$ suppression classifier output for signal and various types of background for the standard BDT classifier (left) and the uBDT classifier (right). B candidates from $B\bar{B}$ background events dominate the lower region, while signal and control candidates dominate in the upper region of the classifier output.

6.4. Selection Optimization

Instead of two separate $q\bar{q}$ and $B\bar{B}$ FOM optimizations, it is more efficient to do a simultaneous 2D FOM optimization, since the two classifiers are not completely uncorrelated. In the same manner as before, FOM is optimized for perfectly reconstructed signal candidates in the signal window, after the pre-selection, signal categorization, and after discarding the background resonances and the control decay. The FOM plot with the optimal point for both $B\bar{B}$ MVA classifiers is shown in Figure 6.7.

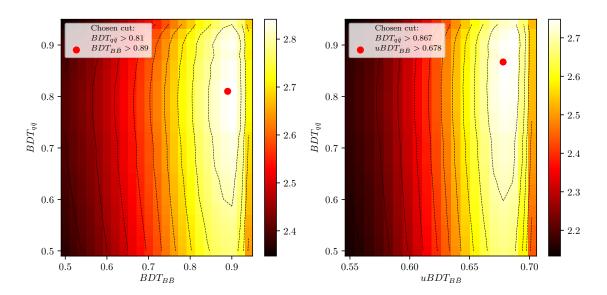


Figure 6.7.: 2D FOM optimization of continuum suppression classifier and the standard BDT (left) and uBDT (right) $B\bar{B}$ suppression classifier.

We can compare signal and major background distributions of ΔE and M_{BC} after the 2D FOM optimization for both classifiers. Figure 6.8 shows the arbitrary (left) and normalized scale (right) for ΔE (top) and M_{BC} (bottom) for the final sample optimized with the standard BDT classifier, while Figure 6.9 shows similarly for the final sample optimized with uBDT classifier. We can see that there is considerably more background in the latter case, however, also shapes of background and signal distributions differ greatly, meaning there is less room for correlation in the extraction process. The biggest change seems to be in the shape of the M_{BC} distribution, where the background component is much more signal like in the final sample optimized with the standard BDT classifier than in the other case. Additionally, the shapes are more easily constrained in the latter case, since they are present in regions where no signal is expected. The total numbers of expected signal candidates and the signal-to-noise ratios for both classifiers are:

- Standard *BDT*: $N_{sig} = 176$, $N_{sig}/N_{bkg} = 4.83 \%$,
- uBDT: $N_{sig} = 264$, $N_{sig}/N_{bkg} = 1.33 \%$.

Due to the large difference in ΔE and M_{BC} shape, we will continue the analysis with the uBDT classifier, although the comparison between both methods will be shown for the final fit result in the next chapter.

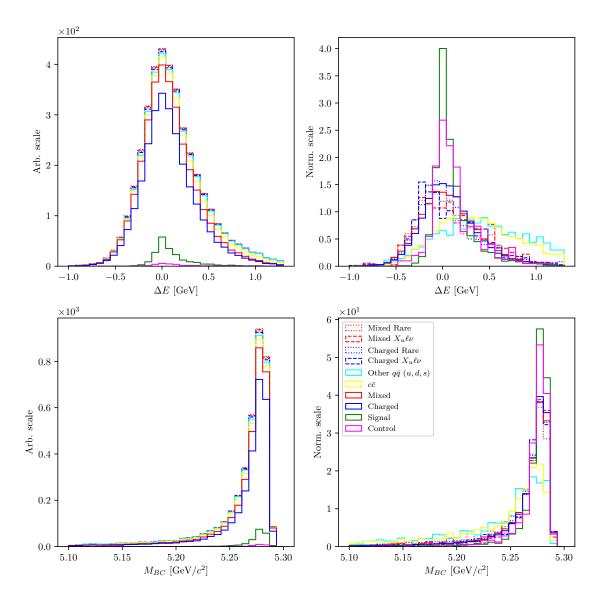


Figure 6.8.: Arbitrary (left) and normalized scale (right) for ΔE (top) and M_{BC} (bottom) for the final sample optimized with the standard BDT classifier.

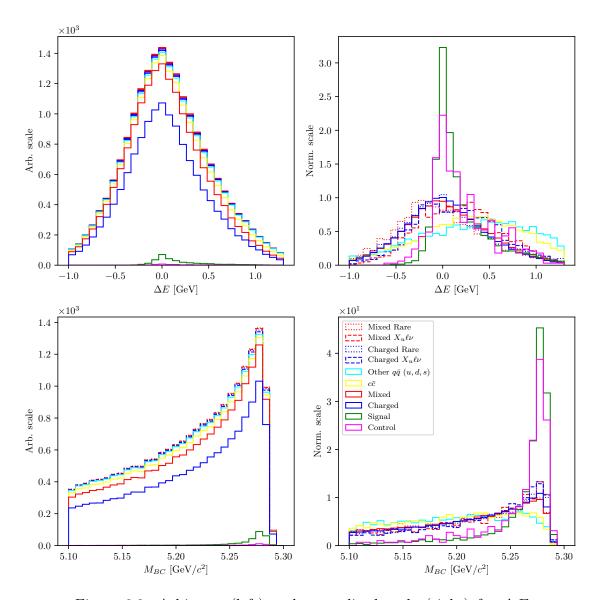


Figure 6.9.: Arbitrary (left) and normalized scale (right) for ΔE (top) and M_{BC} (bottom) for the final sample optimized with the uBDT classifier for $B\bar{B}$ suppression.

6.4.1. $B\bar{B}$ Background Composition and Lepton Veto

The majority of background candidates after the final selection is represented by candidates from $B\bar{B}$ events. In order to suppress this background even further, we need to take a look at its structure and recognize various contributions to this part of the background. Figure 6.10 shows ΔE , M_{BC} , and m_{KK} for the most significant contributions in the signal region. While most of the candidates come from events, where all reconstructed charged particles in the signal decay do not come from a single B meson, but both of them, these candidates are not so problematic. Their distribution is rather smooth and frequent in regions where we expect no signal. On the other hand, there are also contributions from some specific B meson decays, which produce more signal-like distributions. We will denote the first kind of background as $\Upsilon(4S)$ -matched and the second kind as the B-matched $B\bar{B}$ background. Fortunately, these decays are well known and well measured, so their

yields can be constrained. Especially problematic is the double semileptonic decay $B \to \bar{D}^{(*)}\ell^+\nu$, $D^+ \to \bar{K}^-\ell^+\nu$, where the secondary lepton is misidentified as a kaon. Even though the decay has two neutrinos, these events survive the m_{miss}^2 selection and produce peaks at the same positions as the signal distributions, while exhibiting only a slightly worse resolution.

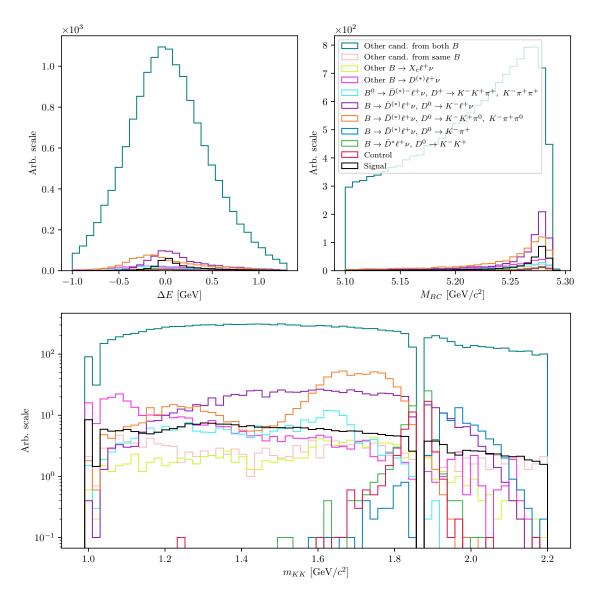


Figure 6.10.: ΔE (left), M_{BC} (right) and m_{KK} (bottom) for major contributions to the $B\bar{B}$ background in the signal region. $\Upsilon(4S)$ -matched backgrounds represents the majority, but have a smooth and wide distribution, distinguishable from signal. B-matched contributions show a peak in M_{BC} and sometimes in ΔE , but can be constrained using existing measurements.

In order to suppress the latter source of background, a lepton veto is applied to both kaons, requiring that neither of the kaons should exhibit lepton-like properties. On the candidates, which pass the final selection, we optimize the eID and μID PID cuts, where S and B in Eq. 4.1 are represented by perfect signal candidates and

by background candidates, respectively. Background in this case is represented by events in which a lepton has been misidentified as a kaon. 2D FOM plots for both kaons are shown in Figure 6.11, where K_0 denotes same-sign, and K_1 the opposite-sign kaon, with respect to the B meson which they are part of. It can be seen that in the majority of cases an electron is misidentified as K_1 . With the optimal selection

- K_0 : eID < 0.8,
- K_1 : eID < 0.1, $\mu ID < 0.8$,

we reject 77.5% of candidates from the double semileptonic decays, while the efficiency loss of the signal candidates and other types of $B\bar{B}$ background is about 5-6%. The $B\bar{B}$ background for the signal region after the lepton veto is shown in Figure 6.12, and in Figure 6.13 for the control region.

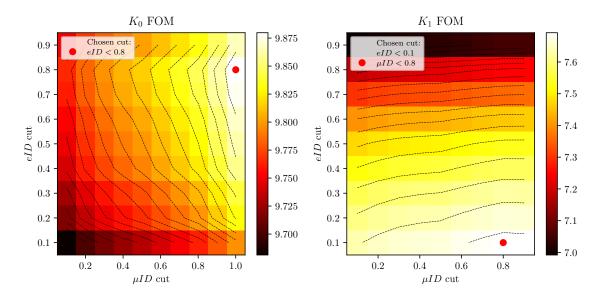


Figure 6.11.: 2D FOM for optimal eID and μID selection on samesign (left) and opposite-sign (right) kaons with respect to the B meson charge. For double semileptonic background component, in most cases an electron is missidentified as the opposite-sign kaon in the reconstruction chain.

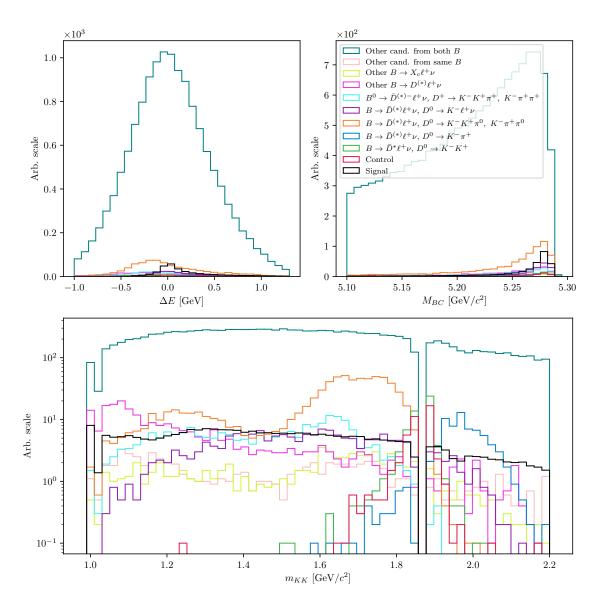


Figure 6.12.: ΔE (left), M_{BC} (right) and m_{KK} (bottom) for major contributions to the $B\bar{B}$ background in the signal region after the lepton veto. The double semileptonic background component is suppressed by a factor of 4-5.

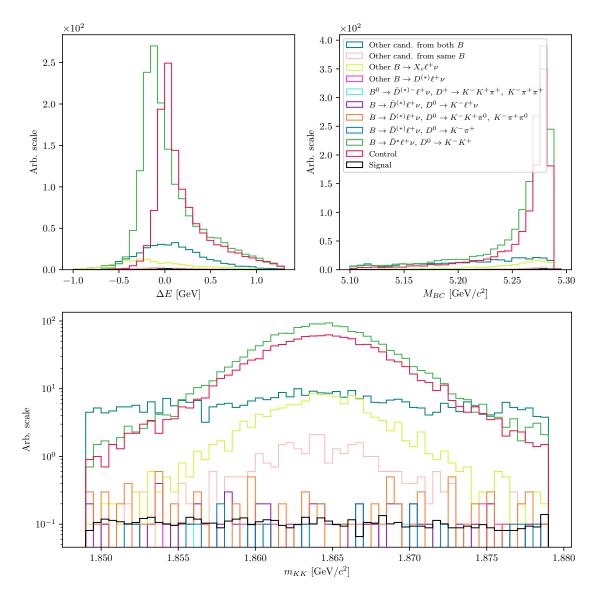


Figure 6.13.: ΔE (left), M_{BC} (right) and m_{KK} (bottom) for major contributions to the $B\bar{B}$ background in the control region after the lepton veto. The major component in this case are other $B \to D^* \ell + \nu$, $D \to K^+ K^-$ decays, besides the control decay.

6.5. Data and MC Agreement

With the final selection in place, we can check the data and MC agreement by checking the control decay region in on- and off- resonance data. Off-resonance samples provide the ability to check the agreement of the $q\bar{q}$ background component, while on-resonance samples can be used to check the validity of the control MC sample and, consequently, the signal MC sample.

6.5.1. Off-resonance Data

The off-resonance data were collected at 60 MeV below the $\Upsilon(4S)$ resonance peak energy, in order to determine the continuum background. It, therefore, offers a direct view of the $q\bar{q}$ background data sample, which we can compare to the off-resonance MC sample. Figure 6.14 shows ΔE , M_{BC} , and the $q\bar{q}$ classifier output, $BDT_{q\bar{q}}$, for off-resonance data and stacked MC in the control region, before the MVA selection, where the MC sample was scaled down by a factor of 6, due to having 6 streams of MC. These figures do not show a fit to the data, but merely an overlay of the data and stacked MC distributions. They show good data and MC agreement for the offresonance sample already before the fit. More importantly than the normalization, the shape of data and MC also seems to match, so further corrections of ΔE and M_{BC} on MC are not necessary. This is also demonstrated by the flatness of the ratio of ΔE and M_{BC} distributions, shown in the same Figure. There seems to be a difference in the classifier performance for the continuum background suppression in data and on MC in the lower region of $BDT_{q\bar{q}}$, where classifier efficiency on MC is overestimated. However, after the selection on classifier output, these differences are negligible, since a relatively small amount of continuum background passes the selection.

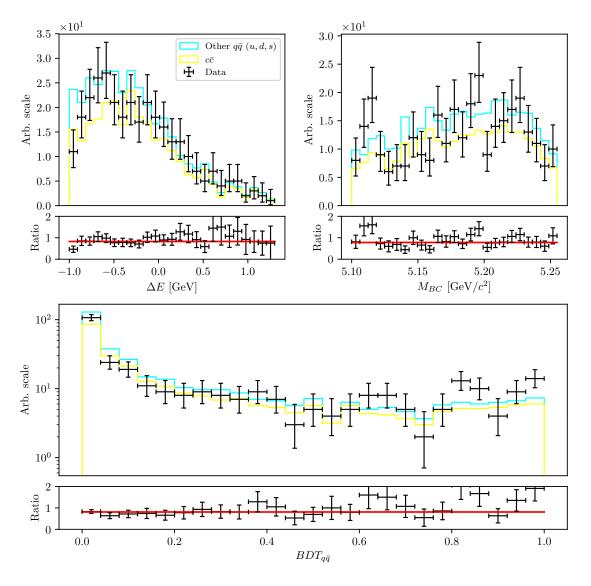


Figure 6.14.: ΔE (left), M_{BC} (right), and the $q\bar{q}$ classifier output (bottom) for off-resonance data and MC in the control region prior to any MVA selection.

6.5.2. On-resonance Data

We repeat the check on on-resonance data. Figure 6.15 shows ΔE , M_{BC} , $BDT_{q\bar{q}}$, and $uBDT_{B\bar{B}}$, where one can see inconsistencies between data and stacked MC on the lower parts of all BDT spectra. These figures again do not show the fit, but merely an overlay of the data to the stacked MC distributions. On the other hand, the data and MC seem to agree well in the upper parts of the spectra. Overall, data and MC seem to agree well after the pre-selection and without any corrections. This means that the modeling of the MC sample is precise and that the MVA selection does not introduce any additional differences between the data and MC for the control and, therefore, the signal sample.

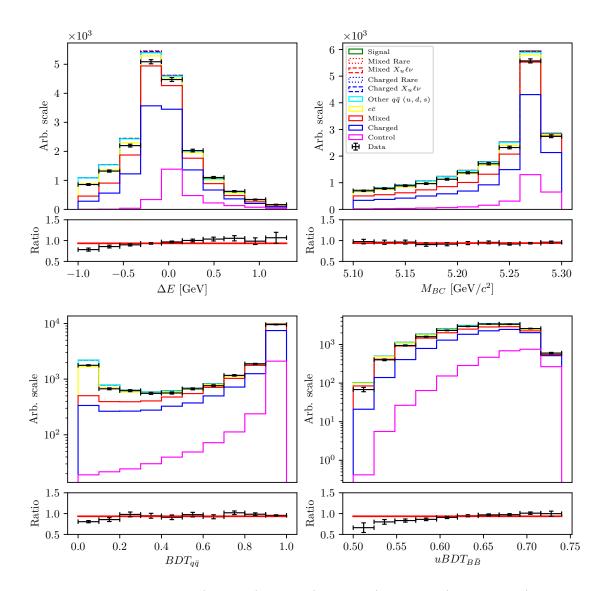


Figure 6.15.: ΔE (top left), M_{BC} (top right), the $q\bar{q}$ - (bottom left) and the $B\bar{B}$ classifier output (bottom right), for onresonance data and MC in the control region prior to the MVA selection.

Chapter 7.

Extraction of Physical Parameters

In this chapter, the procedure for signal yield extraction is presented. We use the framework of RooFit [24], where we define 2D histogram templates in ΔE and M_{BC} , for signal and several types of background, based on MC. Using these templates, the independent full sample is fitted with the binned extended maximum likelihood (ML) fit, so that the individual template ratios and their sum describe the fitted sample as best as possible. In particle physics we are often dealing with low numbers of events and need to account for the Poissonian nature of the data, therefore we use the likelihood fit, since it takes the Poisson errors into account, unlike the χ^2 fit, where the errors are assumed to be Gaussian. In this procedure, we attempt to find the parameter values that maximize the likelihood function, given the observations.

If $P(n|\vec{\alpha})$ is the probability of measuring n candidates, where $\vec{\alpha}$ is a set of parameters on which P depends, we can define the likelihood function L for a series of such measurements (i.e., bins in histogram) n_i based on Poisson statistics as

$$L(\vec{\alpha}) = \prod_{i=1} P(n_i | \vec{\alpha}) = \prod_{i=1} \frac{\mu_i^{n_i} e^{-\mu_i}}{n_i!},$$
(7.1)

where μ_i is the expected value for each measurement. It is also common to search for the minimum of the negative value of $\ln L$, or negative log-likelihood (NLL), as

$$\mathcal{L}(\vec{\alpha}) = -\ln L(\vec{\alpha}) = -\sum_{i} \ln \left(\frac{\mu_i^{n_i} e^{-\mu_i}}{n_i!} \right) = \sum_{i} \ln(n_i!) + \mu_i - n_i \ln(\mu_i). \tag{7.2}$$

Maximizing L or minimizing \mathcal{L} gives us a maximum likelihood estimate of the set of parameters $\vec{\alpha}_{ML}$ which best describe the observed data.

The ML method provides a method to estimate the fit uncertainty. This is especially useful if the log-likelihood has a non-parabolic shape, which leads to asymmetric errors. We calculate the errors using the MINOS algorithm from the MINUIT package [25], which is implemented in RooFit. The algorithm follows the log-likelihood function out of the minimum to find the appropriate intervals of confidence for each parameter, taking the parameter correlations into account.

To estimate the goodness of the likelihood fit, one option is to generate toy MC experiments and obtain the expected log-likelihood distribution. Likelihood fits, however, also offer another way to test the goodness of the fit via the likelihood ratio (LR), where we compare the likelihood obtained under the ML parameters $\vec{\alpha}_{ML}$, to the likelihood obtained under the null hypothesis parameters $\vec{\alpha}_{H_0}$. This determines how likely the data is under one model than the other. We define the LR test as

$$\lambda = -2\ln\left(\frac{L(\vec{\alpha}_{ML})}{L(\vec{\alpha}_{H_0})}\right) = -2\left[\ln L(\vec{\alpha}_{ML}) - L(\vec{\alpha}_{H_0})\right] \sim \chi_q^2,\tag{7.3}$$

which asymptotically behaves as the χ_q^2 distribution with q=m-n degrees of freedom, where m and n are degrees of freedom of $L(\vec{\alpha}_{ML})$ and $L(\vec{\alpha}_{H_0})$, respectively. In particle physics we usually study a specific decay and try to perform measurements of the signal yield, so the null hypothesis in this case is that we expect to observe no signal. This means that, for the null hypothesis, we fix the expected signal yield parameter to zero, while leaving the other parameters of $\vec{\alpha}_{H_0}$ the same as in $\vec{\alpha}_{ML}$, which results in n=m-1 degrees of freedom and in their difference q=m-n=1. For such a simple LR test of a single parameter, the LR test then follows the χ^2 distribution with 1 degree of freedom. In this case we can define the fit significance from the χ^2 value in units of σ as

Significance =
$$\sqrt{\lambda} = \sqrt{\chi^2}$$
. (7.4)

7.1. Fit Setup

We perform 10 fits to each stream of MC, where 9 streams were used for the creation of the templates and the remaining stream was used as the fitted data. When fitting the measured data, all available MC was used for creating the templates. The full signal MC sample was used for the signal template definition in case of MC, as well as the data fit. The signal part of the ulnu sample was not used in template construction, it was only used as a part of the fitted sample.

The same MC samples are used for template construction as described in Chapter 2,

- signal MC,
- 10 streams of charged and mixed $B\bar{B}$ background,
- 6 streams of $c\bar{c}$ (charm) and other $q\bar{q}$ (uds) background,
- ulnu sample, corresponding to 20× integrated luminosity of the full Belle dataset,
- rare sample, corresponding to $50\times$ integrated luminosity of the full Belle dataset.

7.1.1. Control Fit

 $B\bar{B}$ background composition in control region is shown in Figure 6.13. Due to the strict selection of the m_{KK} around the D^0 mass window, most of the decays with a D^0 proceed via $D^0 \to K^+K^-$ decay. In this case, the following fit templates are chosen

- signal template,
- $q\bar{q}$ template,
- $C_0: B^+ \to \bar{D}^0 \ell^+ \nu, D^0 \to K^- K^+$ (control decay),
- $C_1: B \to \bar{D}^*\ell^+\nu, D^0 \to K^-K^+,$
- other $B\bar{B}$ BKG template.

In control fits, all template shapes are fixed and the yields of all templates are floated, except for the $q\bar{q}$ template in cases after the $BDT_{q\bar{q}}$ cut, and the signal template in all cases, since they both have expected yields close to zero and are instead fixed to the expected MC values. Additionally, since the C_0 and C_1 decays are well known and measured, we make use of this fact in the form of a ratio N_{C_1}/N_{C_0} , which is fixed to the MC value in case of the MC fit. In case of fits to the data, we constrain the ratio to the measured value in the form of a Gaussian function with the width corresponding to the measurement uncertainty. These constraints are implemented in the fit, so the systematic effects are taken into account in the fit error. The ratio is implemented based on the decay channels shown in Table 7.1 and is defined as

$$r_1 = \frac{\left(\sum_j N_{1j} \times \rho_{1j}\right)}{N_{00} \times \rho_{00}},\tag{7.5}$$

where j runs over all channels in the category C_1 and where ρ_{ij} is the branching fraction correction factor for the specific channel N_{ij} , which incorporates information from world measurements. It is defined as

$$\rho = \frac{\mathcal{B}^{PDG}}{\mathcal{B}^{GEN}},\tag{7.6}$$

where \mathcal{B}^{PDG} is the measured branching fraction and \mathcal{B}^{GEN} is the branching fraction value used in MC generation. The branching fraction correction factor has been implemented due to differences between measured and MC branching fraction values. Each branching fraction measurement serves as a constraint used in the fit. All branching fraction constraints in the control fit are shown in Table 7.2. The measured values are cited only for the B^0 decay mode, where isospin symmetry has been assumed. The corresponding B^+ branching fractions were calculated as

$$\mathcal{B}(B^+) = \mathcal{B}(B^0) \times \tau_{B^+/B^0},$$
 (7.7)

where τ_{B^+/B^0} is the ratio of B-meson decay times, which is measured to be [8]

$$\tau_{B^+/B^0} = 1.076 \pm 0.004.$$
 (7.8)

Category	Channel	B Decay Mode	D Decay Mode	N_{MC}
C_0	N_{00}	$\mid B^+ \to \bar{D}^0 \ell^+ \nu$	$D^0 \to K^- K^+$	1184 ± 34
C_1	$\begin{array}{ c c } N_{10} \\ N_{11} \end{array}$	$\begin{vmatrix} B^+ \to \bar{D}^{*0} \ell^+ \nu \\ B^0 \to D^{*-} \ell^+ \nu \end{vmatrix}$	$\begin{array}{c} D^0 \to K^- K^+ \\ D^0 \to K^- K^+ \end{array}$	$\begin{vmatrix} 1458 \pm 38 \\ 186 \pm 16 \end{vmatrix}$

Table 7.1.: Well defined decay channels used for constraining the control fits.

In case of the MC fits, the fitted sample is also generated with MC, so $\mathcal{B}_{i}^{PDG} = \mathcal{B}_{i}^{GEN}$, therefore Eq. (7.5) simplifies to a simple MC yield ratio. On fits to real data, expected MC yields and branching fraction measurements are implemented as independent Gaussian constraints in order to properly account for correlations in Eq. (7.5).

ID	Decay	$\mid \mathcal{B}_{GEN} \mid$	$\mid \mathcal{B}_{PDG} \mid$	ρ	Ref.
0	$B^0 \to D^- \ell^+ \nu$	2.13×10^{-2}	$(2.13 \pm 0.09) \times 10^{-2}$	1.00 ± 0.04	[8]
1	$B^0 \to D^{*-}\ell^+\nu$	5.33×10^{-2}	$(4.88 \pm 0.11) \times 10^{-2}$	0.92 ± 0.02	[0]
2	$B^+ o \bar{D}^0 \ell^+ \nu$	2.31×10^{-2}	$(2.29 \pm 0.10) \times 10^{-2}$ $(5.25 \pm 0.12) \times 10^{-2}$	0.99 ± 0.04	
3	$B^+ \to \bar{D}^{*0} \ell^+ \nu$	5.79×10^{-2}	$(5.25 \pm 0.12) \times 10^{-2}$	0.91 ± 0.02	[care.]
4	$D^0 \to K^-K^+$	3.90×10^{-3}	$(3.97 \pm 0.07) \times 10^{-3}$	1.02 ± 0.02	[26]

Table 7.2.: MC and measured values of branching fractions along with the calculated correction factors used for constraining the control fit.

Smearing and Offset Parameters

With simulated data, we are able to perform detailed studies prior to looking at the measured data. However, simulated data often does not describe real data perfectly. Out of variables ΔE and M_{BC} , ΔE is especially prone to a lack of precision in energy measurements. This can introduce either overestimation of resolution on MC, as well as a possible shift in the measured energy in either direction. Due to this fact, we introduce a modification of the ΔE variable by applying a smearing factor and an offset. These are applied by simple transformations of

$$f_{\text{offset}}: x \mapsto x + a,$$
 (7.9)

$$f_{\text{smearing}}: x \mapsto \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma^2}},$$
 (7.10)

where x goes over all entries in the ΔE distribution, a is the energy offset and f_{smearing} corresponds to the normal distribution with mean μ and standard deviation σ . Since this operation is applied on the candidate level, the smearing introduces some randomness to the fit procedure, which is the reason why we repeat the fit procedure several times in order to reliably extract the parameters of interest. In the case of the M_{BC} variable the mentioned effects are not as prominent, so the smearing and offset for the latter variable are omitted.

The following parameter phase-space is scanned in order to determine the best parameter values

- smearing factor in range [0.0, 0.08] GeV in steps of 8×10^{-3} ,
- offset in range [0.0, 0.003] GeV in steps of 1.5×10^{-4} ,

where, for each parameter pair, the likelihood ratio test is performed to estimate the goodness of the fit. Figure 7.1 shows the contour plot of the likelihood ratio λ , as defined in Eq. (7.3), for 2 degrees of freedom, for MC (left) and data (right). The scan over MC serves the purpose of a consistency check, where we expect the best fit to occur in the phase-space where neither smearing nor offset are applied. In the case of data, we see that we obtain a better fit by introducing some level of smearing and offset. In both cases, the two parameters have shown no sign of significant correlation, so we treat them independently. The likelihood ratio test allows us to estimate the parameter values in the 1σ confidence interval, where we obtain the optimal parameter set

• Smearing: $40^{+15}_{-17} \text{ MeV}$,

• Offset: $6^{+4.6}_{-6}$ MeV.

We apply this transformation to our MC samples in all cases when fitting to the real data.

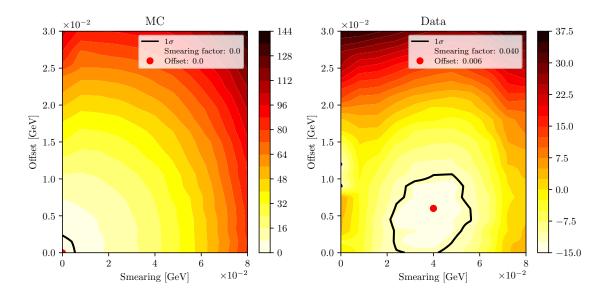


Figure 7.1.: Likelihood ratio test of an additional smearing and offset parameter to MC (left) and data (right).

7.1.2. Signal Fit

The motivation for the choice of the signal fit templates comes from Figure 6.12. The following histogram templates were defined

- signal template,
- $q\bar{q}$ template,
- a series of well defined templates from $B\bar{B}$ background:

$$-C_0: B^+ \to \bar{D}^0 \ell^+ \nu, D^0 \to K^- K^+ \text{ (control decay)},$$

$$-C_1: B \to \bar{D}^* \ell^+ \nu, D^0 \to K^- K^+,$$

$$-C_2: B \to \bar{D}^{(*)} \ell^+ \nu, D^0 \to K^- \pi^+,$$

$$-C_3: B \to \bar{D}^{(*)} \ell^+ \nu, D^0 \to K^- K^+ \pi^0, K^- \pi^+ \pi^0,$$

$$-C_4: B \to \bar{D}^{(*)} \ell^+ \nu, D^0 \to K^- \ell^+ \nu,$$

$$-C_5: B^0 \to D^{(*)} \ell^+ \nu, D^+ \to K^- K^+ \pi^+, K^- \pi^+ \pi^+,$$

$$-C_6: \text{ other } B \to \bar{D}^{(*)} \ell^+ \nu \text{ decays},$$

• remaining $B\bar{B}$ background template.

As mentioned in Chapter 6, the majority of the background comes from $B\bar{B}$ events. Various processes (C_0 to C_6) contribute to this background, which are well known and measured, so we make use of these measurements by fixing their yields in MC fits and appropriately constraining them in the data fits. Similarly, as described in the control fits, these constraints are implemented in the fit process. The remaining

 $B\bar{B}$ background is merged into a single template. The shape of all templates is fixed, while the yields are floated for all templates except for the constrained background templates. The yield constraints are based on the channels shown in Table 7.3 and defined for each template category as

$$Y_i = \eta_{\text{norm.}} \times \frac{\left(\sum_j N_{ij} \times \rho_{ij}\right)}{\rho_{00}},\tag{7.11}$$

where j runs over all channels in the category C_i and where ρ_{ij} is defined in Eq. (7.6). The first factor, $\eta_{\text{norm.}}$, serves as a normalization factor in order to scale the number of generated $B\bar{B}$ events to the number of $B\bar{B}$ events in measured data. We define it as

$$\eta_{\text{norm.}} = \frac{N_{\text{control}}^D}{N_{\text{control}}^{MC}},\tag{7.12}$$

where N_{control}^D and N_{control}^{MC} are control yields in the control fit for the data and MC, respectively.

In addition to the branching fraction constraints in Table 7.2, further constraints are defined in Table 7.4. In case of the category C_6 , we have no firm handle on the D meson decay, therefore no correction for this branching fraction can be introduced, so we set a correction factor of 1 with a 100% error for the D meson decay branching fraction. As most of the correction factors used for constraints have deviations (including the errors) from nominal values well below 100%, this value is very conservative.

Category	Channel	B Decay Mode	D Decay Mode	Expected MC Yield
C_0	N_{00}	$\mid B^+ \to \bar{D}^0 \ell^+ \nu$	$D^0 \to K^- K^+$	44 ± 7
C_1	$\begin{array}{ c c } N_{10} \\ N_{11} \end{array}$	$\begin{vmatrix} B^+ \to \bar{D}^{*0} \ell^+ \nu \\ B^0 \to D^{*-} \ell^+ \nu \end{vmatrix}$	$\begin{array}{c} D^0 \to K^- K^+ \\ D^0 \to K^- K^+ \end{array}$	$\begin{array}{ c c c }\hline 53 \pm 7 \\ 6 \pm 2 \end{array}$
C_2	$egin{array}{c c} N_{20} \\ N_{21} \\ N_{22} \end{array}$	$\begin{vmatrix} B^+ \to \bar{D}^0 \ell^+ \nu \\ B^+ \to \bar{D}^{*0} \ell^+ \nu \\ B^0 \to D^{*-} \ell^+ \nu \end{vmatrix}$	$D^{0} \rightarrow K^{-}\pi^{+}$ $D^{0} \rightarrow K^{-}\pi^{+}$ $D^{0} \rightarrow K^{-}\pi^{+}$	$ \begin{vmatrix} 23 \pm 5 \\ 41 \pm 6 \\ 6 \pm 2 \end{vmatrix} $
C_3	$ \begin{array}{c c} N_{30} \\ N_{31} \\ N_{32} \\ N_{33} \\ N_{34} \\ N_{35} \end{array} $		$D^{0} \to K^{-}K^{+}\pi^{0}$ $D^{0} \to K^{-}\pi^{+}\pi^{0}$ $D^{0} \to K^{-}K^{+}\pi^{0}$ $D^{0} \to K^{-}\pi^{+}\pi^{0}$ $D^{0} \to K^{-}K^{+}\pi^{0}$ $D^{0} \to K^{-}K^{+}\pi^{0}$ $D^{0} \to K^{-}\pi^{+}\pi^{0}$	
C_4	$ \begin{vmatrix} N_{40} \\ N_{41} \\ N_{42} \\ N_{43} \\ N_{44} \\ N_{45} \end{vmatrix} $	$\begin{vmatrix} B^{+} \rightarrow \bar{D}^{0}\ell^{+}\nu \\ B^{+} \rightarrow \bar{D}^{0}\ell^{+}\nu \\ B^{+} \rightarrow \bar{D}^{*0}\ell^{+}\nu \\ B^{+} \rightarrow \bar{D}^{*0}\ell^{+}\nu \\ B^{0} \rightarrow D^{*-}\ell^{+}\nu \\ B^{0} \rightarrow D^{*-}\ell^{+}\nu \end{vmatrix}$	$D^{0} \rightarrow K^{-}e^{+}\nu$ $D^{0} \rightarrow K^{-}\mu^{+}\nu$ $D^{0} \rightarrow K^{-}e^{+}\nu$ $D^{0} \rightarrow K^{-}\mu^{+}\nu$ $D^{0} \rightarrow K^{-}e^{+}\nu$ $D^{0} \rightarrow K^{-}\mu^{+}\nu$	$ \begin{vmatrix} 47 \pm 7 \\ 7 \pm 3 \\ 98 \pm 10 \\ 10 \pm 3 \\ 14 \pm 4 \\ 3 \pm 2 \end{vmatrix} $
C_5	$egin{array}{c} N_{50} \\ N_{51} \\ N_{52} \\ N_{53} \end{array}$	$ \begin{vmatrix} B^0 \to D^- \ell^+ \nu \\ B^0 \to D^- \ell^+ \nu \\ B^0 \to D^{*-} \ell^+ \nu \\ B^0 \to D^{*-} \ell^+ \nu \end{vmatrix} $	$D^{+} \rightarrow K^{-}K^{+}\pi^{+}$ $D^{+} \rightarrow K^{-}\pi^{+}\pi^{+}$ $D^{+} \rightarrow K^{-}K^{+}\pi^{+}$ $D^{+} \rightarrow K^{-}\pi^{+}\pi^{+}$	$ \begin{vmatrix} 102 \pm 10 \\ 63 \pm 8 \\ 31 \pm 6 \\ 21 \pm 5 \end{vmatrix} $
C_6	$egin{array}{c} N_{60} \\ N_{61} \\ N_{62} \\ N_{63} \\ \end{array}$		Other D^0 and D^+ decays	$ \begin{vmatrix} 69 \pm 8 \\ 94 \pm 10 \\ 63 \pm 8 \\ 35 \pm 6 \end{vmatrix} $

Table 7.3.: Well defined decay channels used for constraining the signal fits.

ID	Decay	$\mid \mathcal{B}_{GEN} \mid$	$\mid \mathcal{B}_{PDG} \mid$	ρ	Ref.
5	$D^0 \to K^- \pi^+$	3.82×10^{-2}	$(3.89 \pm 0.04) \times 10^{-2}$	1.02 ± 0.01	
6	$D^0 o K^-K^+\pi^0$	2.36×10^{-3}	$(3.37 \pm 0.15) \times 10^{-3}$	1.43 ± 0.06	
7	$D^0 \to K^- \pi^+ \pi^0$	13.08×10^{-2}	$(14.2 \pm 0.5) \times 10^{-2}$	1.09 ± 0.04	
8	$D^0 \to K^- e^+ \nu$	3.41×10^{-2}	$(3.53 \pm 0.028) \times 10^{-2}$	1.04 ± 0.01	[26]
9	$D^0 \to K^- \mu^+ \nu$	3.41×10^{-2}	$(3.31 \pm 0.13) \times 10^{-2}$	0.97 ± 0.04	
10	$D^+ \rightarrow K^- K^+ \pi^+$	9.06×10^{-3}	$(9.51 \pm 0.34) \times 10^{-3}$	1.05 ± 0.04	
11	$D^+ \to K^- \pi^+ \pi^+$	9.51×10^{-2}	$8.98 \pm 0.28 \times 10^{-2}$	0.94 ± 0.03	

Table 7.4.: Additional MC and measured values of D meson branching fractions along with the calculated correction factors used for constraining the signal fit.

7.2. Adaptive Binning Algorithm

The fit templates contain areas of low statistics, which are populated with bins with zero content. This is a direct consequence of having a finite MC sample and represent a liability in ML fits. Due to the low statistics in the edge regions, the locations of these empty bins can vary for the templates and the fitted sample. A problem occurs if all templates have an empty bin where the fitted sample does not. In the scope of ML fits, this effectively means that there are entries in bins, where the probability of having them is 0. We will call such bins *problematic*, because in these cases the fit does not converge.

The ideal solution for this problem would be to increase the MC statistics. Since this is not an option, we pursue other solutions, such as decreasing the number of bins. While this solves the problem, the drawback of it is a decrease in the template resolution in densely populated regions, where good resolution is most needed. The compromise solution seems to be a choice of variable bins, with fine binning in the densely populated regions and larger bins in the regions with low statistics.

We have devised an algorithm, which compares the templates and the fitted sample, and defines a variable binning so that there are no more problematic bins in the end. Figure 7.2 shows an example of how the procedure works. The algorithm does the following

- 1. define uniform binning in both dimensions,
- 2. create a 2D histogram from MC templates with expected yields,
- 3. define an *optimal* region, where most of the 2D integral is contained and where all bins have non-zero content (this region does not change throughout the process),
- 4. compare the histograms for the expected and the fitted sample, find the problematic bins,
- 5. loop until all problematic bins disappear
 - a) find the problematic bin, which is nearest to the maximum bin,
 - b) change the binning from N to N-1 from that bin and in the direction away from the maximum bin.

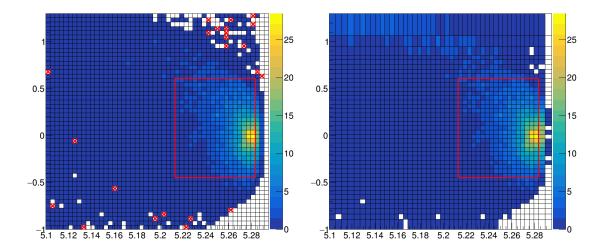


Figure 7.2.: Steps taken in the adaptive binning algorithm. Left image shows the initial 2D histogram with the defined optimal region and the problematic bins, the right image shows the final binning with the unchanged optimal region, while the problematic bins are gone due to the new binning choice.

An additional problem occurs in the plotting of the fitted templates with variable binning. It would seem that RooFit does not take the bin widths into account when plotting, while everything works as expected for the fit itself. This was bypassed by extracting the fitted yields and applying them to the templates, defined with uniform binning.

7.3. Toy MC Experiments

For the chosen final selection and fit procedure, toy MC pseudo-experiments were performed in order to confirm the behavior of the fit setup. The fit behavior is also checked as a function of the signal yield in the form of a linearity test. A detailed description of toy MC experiments is written in this section.

With toy MC experiments we study the yields, errors and the pulls of the signal fit by generating our own pseudo-datasets, according to the MC. This significantly reduces the time we would need to produce the datasets in the standard way, while still reliably describing the underlying physics behind the pseudo-dataset. All available MC was used for pseudo-dataset generation, as well as creating the templates.

7.3.1. Pseudo-Experiment: Expected Signal Yield

We constructed 3×10^3 pseudo-datasets, where each dataset was generated with the expected amount of each template category, distributed according to the Poisson distribution. All fits were performed with the optimal initial uniform binning of 19×19 bins in ΔE and M_{BC} . The optimal choice of binning is based on the binning test in Section 8.1.

Figure 7.3 shows distributions of the fit yields, errors and the pull distribution of all pseudo-fits. The pull distribution g of a normally distributed variable x with a mean μ and a width σ is defined as

$$g = \frac{x - \mu}{\sigma},\tag{7.13}$$

and is also distributed normally, with mean zero and unit width. The fits in Figure 7.3 seem to be under control, although there is a slight bias present in the negative direction, which can also be seen in the pull distribution plot. The latter follows a normal distribution with a mean of (-0.11 ± 0.02) and a standard deviation of (1.01 ± 0.01) . The mean (\bar{X}) and the standard deviation (S) were calculated in the usual way, while their errors $\sigma_{\bar{X}}$ and σ_{S} were calculated as [27]

$$\sigma_{\bar{X}} = \frac{S}{\sqrt{N}}, \quad \sigma_S = \frac{S}{\sqrt{2(N-1)}}, \tag{7.14}$$

where N is the number of performed measurements. A decision has been made to put this yield difference as one of the contributions to the overall systematic uncertainty.

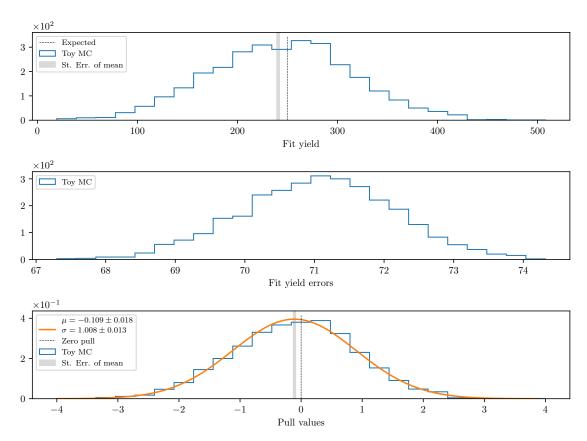


Figure 7.3.: Toy MC fits of pseudo-data showing the fit yield (top), the fit errors (center) and the pull distribution of the fits (bottom).

7.3.2. Pseudo-Experiment: Linearity Test

Linearity test is used for determining the sensitivity of the fit to the amount of signal in the fitted sample. Since this is the first measurement of this decay channel,

MC modeling is not reliable and could be very different from reality, so we need to perform this test in order to determine our sensitivity to smaller, as well as larger amounts of expected signal.

The pseudo-datasets are generated in the same way as in the previous section, with the exception of signal, which is generated in various amounts. 50 steps from [0.1, 10] in the logarithmic scale are taken for fractions of signal amount, where, for each fraction, we generate 500 pseudo-datasets according to Poisson statistics.

Figure 7.4 shows the difference between the mean fit yield and the expected yield, the mean pull and the mean significance at each signal fraction value. The expected MC result lies at the fraction value $10^0 = 1$. The plots show a small bias with respect to the expected signal yield, while the pulls seem to be described by the normal distributions throughout the fraction range with a slight offset, which will be incorporated as a contribution to the overall systematic uncertainty. At expected value, we are at about 3.6σ statistical significance.

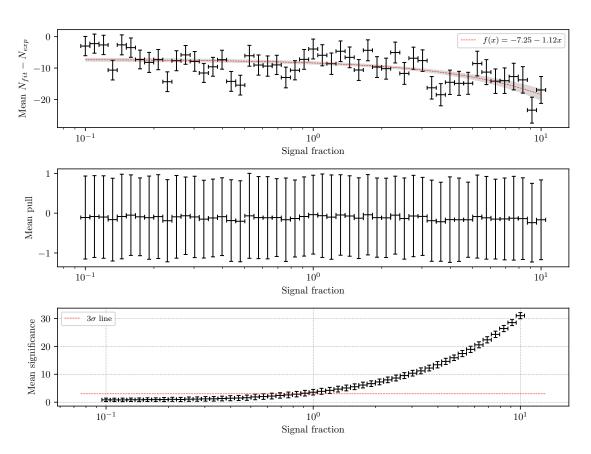


Figure 7.4.: The mean fit yield and expected yield difference (top), the mean pull (center) and the mean significance (bottom) as a function of the signal fraction.

Chapter 8.

Fit Results

In this chapter, we present the results of signal and control fits on MC, as well as in data, along with the control decay branching fraction measurement.

8.1. Signal MC Fit Results

With the signal fit setup described in Section 7.1.2, we proceed to fit the 10 streams of MC. To compare both methods of $B\bar{B}$ suppression, two different samples were prepared and used in the fit. Since the choice of initial uniform binning is not obvious, we perform fits to all streams of MC, for each binning choice in the range $N \times N, N \in [4,30]$. Figure 8.1 shows the fitted and the true MC matched yield differences, the pulls, and fit significances for both final samples for each binning case. The difference between fitted an expected signal yield should be equal to 0 to ensure no bias is present in the fit, while the average pull distribution for each bin case should have a central value of 0, and a width of 1. From the top plot in Figure 8.1, we see that the yield difference is consistent with 0, while exhibiting a slight, but a consistent overestimation of the fitted yield. This difference is accounted for as a separate contribution to the overall systematic uncertainty. The pull distribution seems to be closer to the normal distribution for the case of the uBDT classifier. Also the fit significance is larger by approximately 1σ , so we conclude that the uBDT classifier outperforms the standard BDT setup. The binning in ΔE and M_{BC} is chosen at the plateau of the significance, where no significant bias is present and is somewhere in the region of 20 bins in each dimension.

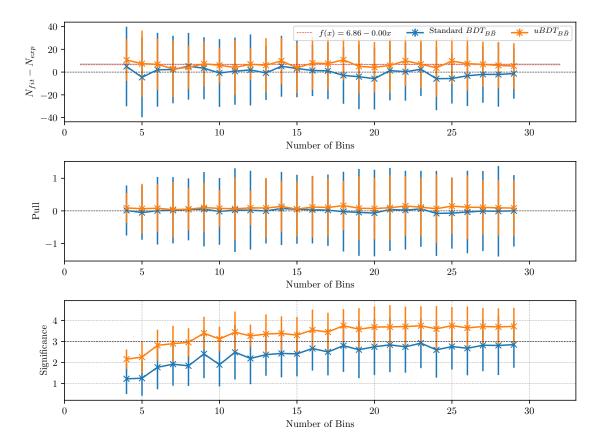


Figure 8.1.: Expected yield difference (top), pulls (center) and fit significance (bottom) as a function of binning in ΔE and M_{BC} for the final sample, optimized with the standard BDT classifier (blue) and the uBDT classifier (orange).

For the chosen binning of 19×19 in ΔE and M_{BC} we perform the 10 stream MC fits, where an average stream fit is shown in Figure 8.2, while all fit results are shown in Figure 8.3. All stream fit results were fitted with a 0th degree polynomial. The global result seems to describe the expected value precisely, with the residual bias much smaller than the average statistical error. The normalized χ^2 value with 10-1=9 degrees of freedom for the global fit is $\chi_9^2=1.23$, while the average statistical significance of the fits is around 3.5σ .

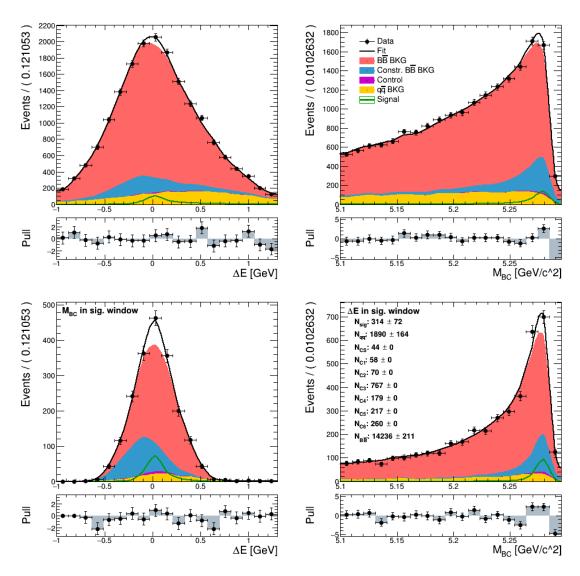


Figure 8.2.: An example fit to one stream of MC. Left column shows the ΔE and the right column shows the M_{BC} distribution of the full fitted sample in the full fit region (top) and the in signal enhanced region (bottom).

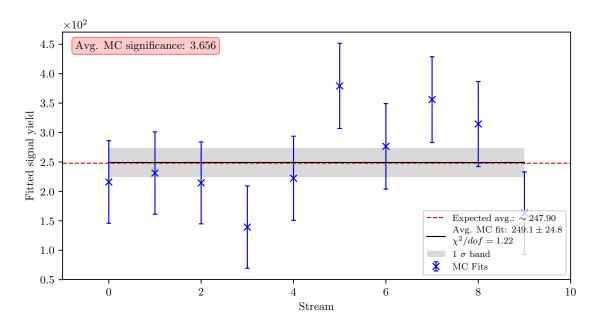


Figure 8.3.: Fits to all 10 streams of MC and the global fit with a 0^{th} degree polynomial. The red line shows the mean value of the global fit and the gray band shows the 1σ confidence interval.

8.2. Control Fit Result

With the control fit setup described in Section 7.1.1, we proceed to fit the control sample for the 10 streams of MC and 1 stream of data. A typical MC stream fit is shown in Figure 8.4 for MC and in Figure 8.5 for the data, while all fit results are shown in Figure 8.6, where all streams of MC are fitted with a 0th degree polynomial. The control fit results for the split and joined lepton modes are shown in Table 8.1.

	N^{MC}	$N^{ m data}$
$\ell = e \text{ or } \mu$	1182 ± 11	1187 ± 44
$\ell = e$	591 ± 8	583 ± 28
$\ell = \mu$	592 ± 7	613 ± 30

Table 8.1.: Control sample fit results for MC and data for various lepton final state modes.

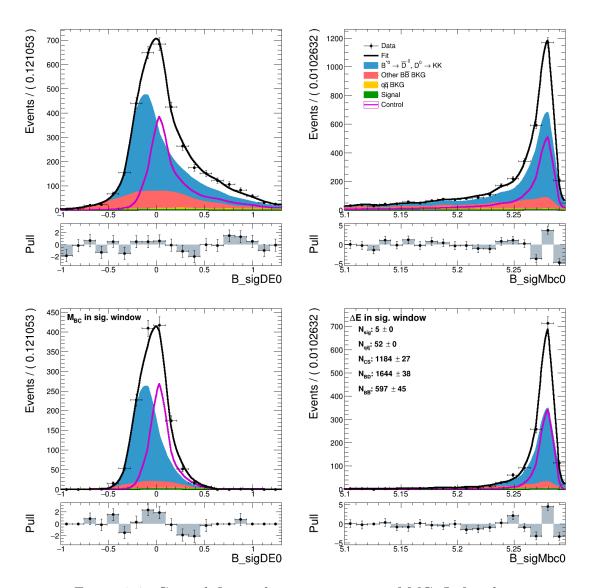


Figure 8.4.: Control fit result on one stream of MC. Left column shows the ΔE and the right column shows the M_{BC} distribution in the full fit window (top) and the in signal enhanced window (bottom).

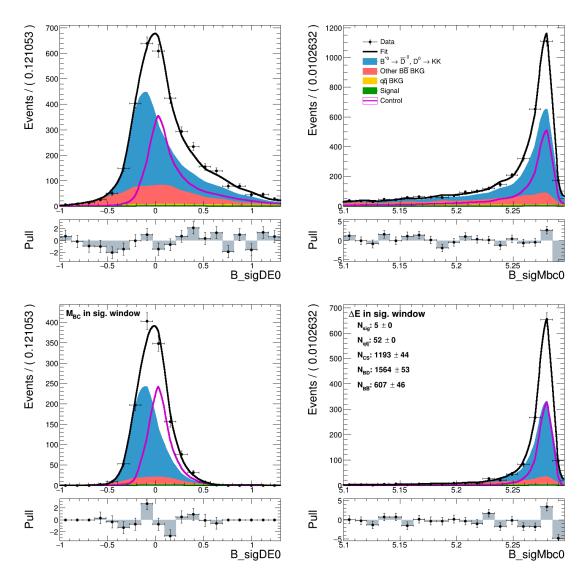


Figure 8.5.: Control fit result on real data. Left column shows the ΔE and the right column shows the M_{BC} distribution in the full fit window (top) and the signal enhanced window (bottom).

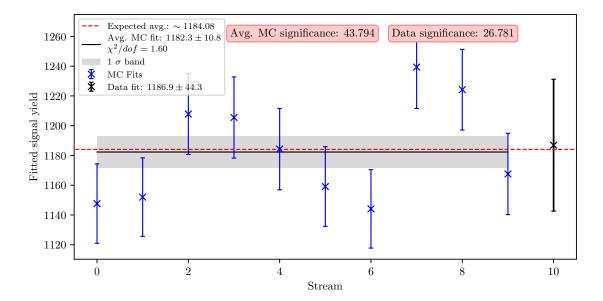


Figure 8.6.: Control fit to the data and all 10 streams of MC. The red line shows the mean value of the global MC fit with a 0th degree polynomial. The gray band shows the 1σ confidence interval around the global MC fit.

8.2.1. Branching Ratio Measurement for Control Decay

After acquiring the fit results on MC and data, we are able to determine the branching fraction of the control decay, which is defined as

$$\mathcal{B}_{\text{control}}^{MC} = \frac{N_{\text{control}}^{\text{MC}} \times \epsilon_{MC}}{2N_{B\bar{B}}^{MC}},$$

$$\mathcal{B}_{\text{control}} = \frac{N_{\text{control}} \times \epsilon_{MC} \times \rho_{PID}}{2N_{B\bar{B}}},$$
(8.1)

$$\mathcal{B}_{\text{control}} = \frac{N_{\text{control}} \times \epsilon_{MC} \times \rho_{PID}}{2N_{B\bar{B}}},$$
(8.2)

where $N_{\rm control}^{\rm MC}$ and $N_{\rm control}$ are yields of the control fit on MC and data, ϵ_{MC} is the MC efficiency of the control sample, ρ_{PID} the PID correction factor, and $N_{R\bar{R}}^{MC}$ and $N_{B\bar{B}}$ the numbers of $B\bar{B}$ meson pairs on MC and data, respectively. The factor of 2 in the denominator comes from the fact that there are 2 B mesons in each B meson pair ($\times 1/2$), where only about 50% of the B meson pairs are charged B^+B^- meson pairs $(\times 2)$, and from the fact that we are interested in the branching fraction to the lepton final state of either e or μ , and not their sum ($\times 1/2$).

The control sample efficiency was determined on a separate signal MC sample of the control decay, where we generated $5 \times 10^6~B^+\bar{B}^-$ pairs, with one B always decaying via $B^+ \to \bar{D}^0 \ell^+ \nu$, $D^0 \to K^+ K^-$. After applying the final selection, the full and split efficiencies, with respect to the lepton final state, were determined to be

$$\epsilon_{MC} = (8.61 \pm 0.04) \times 10^{-3},$$

$$\epsilon_{MC}^e = (4.25 \pm 0.03) \times 10^{-3},$$

$$\epsilon_{MC}^\mu = (4.35 \pm 0.03) \times 10^{-3}.$$

The efficiency error was estimated according to the binomial distribution

$$\sigma_{\epsilon_{MC}} = \frac{1}{N} \sqrt{n(1 - \frac{n}{N})},$$

where n is a subset of the full set N.

The PID correction factor is obtained by taking into account the known PID efficiency differences between the data and MC. It is described in detail in Section 9.1.1 and is determined to be

$$\rho_{PID} = 0.99 \pm 0.02$$

for the e and μ mode, as well as both of them together.

The number of BB meson pairs can be counted on MC and has been measured for the data sample by the collaboration. The values are

$$N_{B\bar{B}}^{MC} = 765.98 \times 10^6,$$

 $N_{B\bar{B}} = (771.58 \pm 10.56) \times 10^6.$

Finally, we can determine the branching fractions based on the calculations in Eq. (8.2). The obtained values are shown in Table 8.2 and graphically presented in Figure 8.7, along with the MC generated value and the current PDG world average. Both MC and the data results for the control decay branching fraction are in agreement with the expected and the world average values. One should note that the error bars correspond to statistical uncertainty and the PID systematic uncertainty only. Other systematic uncertainties are not included, since this measurement is not the goal of our analysis.

	$\mathcal{B}_{GEN} \ [\times 10^{-5}]$	$\mathcal{B}_{PDG} \left[\times 10^{-5} \right]$	$\mid \mathcal{B}^{MC} \mid \times 10^{-5} \mid$	$\mathcal{B}^{\text{data}} \left[\times 10^{-5} \right]$
$\ell = e \text{ or } \mu$ $\ell = e$ $\ell = \mu$	9.01	9.10 ± 0.42	$8.97 \pm 0.09 9.07 \pm 0.14 8.88 \pm 0.12$	9.03 ± 0.40 9.02 ± 0.49 9.18 ± 0.50

Table 8.2.: Control sample fit results for MC and data for various lepton final state modes.

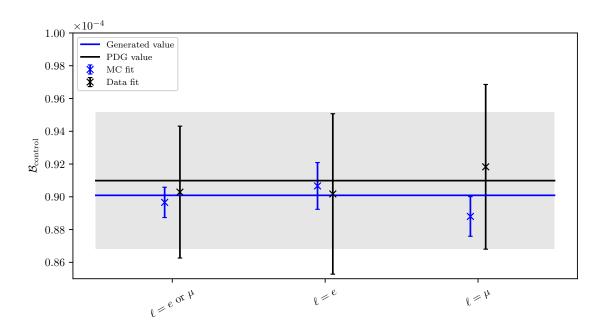


Figure 8.7.: Various branching fraction determinations for the control decay of our analysis.

8.3. Results of Signal Fit in Data

8.3.1. Signal Yield

Finally, after validating the analysis procedure using the signal MC sample, control MC sample and the control data sample, we can continue to perform the signal extraction process on the full Belle $\Upsilon(4S)$ data sample. Figure 8.8 shows the fit result in projections of ΔE and M_{BC} for both fit regions. The extracted signal yield in data, as well as the yields of the remaining contributions, are shown in Table 8.3. Values of all the constraints are shown in Table 8.4. As the main results, we obtain the signal yield of

$$N_{\text{sig}} = 491 \pm 86$$
, fit significance = 6.0σ . (8.3)

The uncertainty includes the statistical uncertainty, as well as the partial systematic uncertainty, due to the limited knowledge of various branching fractions, entering the fit in a form of Gaussian constraints. The significance of the result, taking into account only the mentioned uncertainties, is 6.0σ . The latter is obtained via the likelihood-ratio test, as previously described in Eq. (7.4).

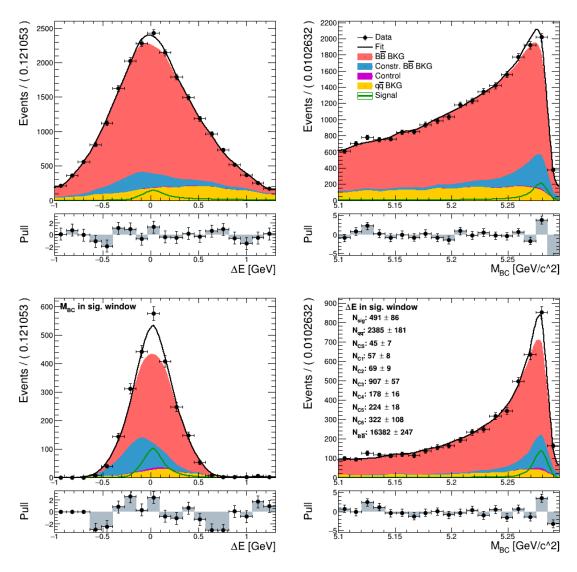


Figure 8.8.: Signal fit result on real data. Left column shows the ΔE and the right column shows the M_{BC} distribution in the full fit window (top) and in the signal enhanced window (bottom).

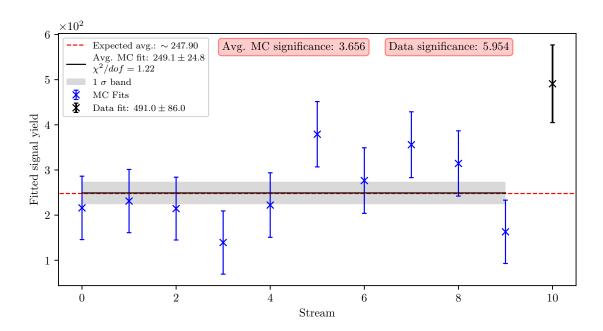


Figure 8.9.: Signal fit to the data and all 10 streams of MC. The red line shows the mean value of the global MC fit with a $0^{\rm th}$ degree polynomial. The gray band shows the 1σ confidence interval around the global MC fit.

Category	Fit Yield
Signal	491 ± 86
$q\bar{q}$ background	2385 ± 181
C_0	45 ± 7
C_1	57 ± 8
C_2	69 ± 9
C_3	907 ± 57
C_4	178 ± 16
C_5	224 ± 18
C_6	322 ± 108
Other $B\bar{B}$ background	16382 ± 247

Table 8.3.: Yields of all signal fit contributions in data.

Constraint	Value	Constraint	Value
\mathcal{B}_0	0.049 ± 0.001	N_{30}	103 ± 10
\mathcal{B}_1	1.076 ± 0.004	N_{31}	212 ± 14
\mathcal{B}_2	0.021 ± 0.001	N_{32}	136 ± 12
\mathcal{B}_3	1.014 ± 0.018	N_{33}	268 ± 16
\mathcal{B}_4	1.018 ± 0.010	N_{34}	20 ± 4
\mathcal{B}_5	1.439 ± 0.063	N_{35}	35 ± 6
\mathcal{B}_6	1.092 ± 0.038	N_{40}	48 ± 7
\mathcal{B}_7	1.035 ± 0.008	N_{41}	7 ± 3
\mathcal{B}_8	0.971 ± 0.038	N_{42}	99 ± 10
\mathcal{B}_9	1.053 ± 0.038	N_{43}	10 ± 3
\mathcal{B}_{10}	0.946 ± 0.029	N_{44}	14 ± 4
\mathcal{B}_{11}	1.264 ± 0.446	N_{45}	3 ± 2
$N_{ m control}^{MC}$	1181 ± 11	N_{50}	104 ± 10
$N_{ m control}^{Data}$	1210 ± 43	N_{51}	64 ± 8
N_{00}	44 ± 7	N_{52}	31 ± 6
N_{10}	54 ± 7	N_{53}	22 ± 5
N_{11}	6 ± 2	N_{60}	69 ± 8
N_{20}	23 ± 5	N_{61}	94 ± 10
N_{21}	41 ± 6	N_{62}	63 ± 8
N_{22}	6 ± 2	N_{63}	35 ± 6

Table 8.4.: Mean values and standard deviations of constraints after the fit.

In this fit setup, we perform a random smearing of the ΔE (see Eq. 7.9 and description there). Because of the randomness involved in the smearing, such a method results in small random fluctuations of the fitted central values. In order to check the consistency of the fit result because of the ΔE smearing, we perform 500 fit repetitions. The result in Eq. (8.3) represents the average result, corresponding to the central values of the yield, uncertainty, and the significance of the signal fit in data.

The systematic part of the fit uncertainty can be estimated by fixing the Gaussian constraints to central values, presented in Table 8.4, and repeating the fit. The resulting uncertainty and significance are purely statistical and are found to be

$$\sigma_{\text{stat}} = 81$$
, fit significance = 6.3 σ . (8.4)

Other sources of systematic uncertainties and their estimations are presented in details in Chapter 9.

8.3.2. Branching Ratio

Similarly as for the control decay, we are able to calculate the branching fraction of the signal decay via the formulas

$$\mathcal{B}_{\text{sig}}^{MC} = \frac{N_{\text{sig}}^{\text{MC}} \times \epsilon_{MC}}{2N_{B\bar{B}}^{MC}},\tag{8.5}$$

$$\mathcal{B}_{\text{sig}} = \frac{N_{\text{sig}} \times \epsilon_{MC} \times \rho_{PID}}{2N_{B\bar{B}}},\tag{8.6}$$

where $N_{\rm sig}^{\rm MC}$ and $N_{\rm sig}$ are yields of the signal fit on MC and data, ϵ_{MC} is the MC efficiency of the signal sample, ρ_{PID} the PID correction factor, and $N_{B\bar{B}}^{MC}$ and $N_{B\bar{B}}$ are the numbers of $B\bar{B}$ meson pairs on MC and in data, respectively.

The signal efficiency was determined on the same signal MC sample as was used throughout the analysis. The full signal efficiency is determined to be

$$\epsilon_{MC} = (1.052 \pm 0.003) \times 10^{-2},$$

where the efficiency error was calculated in the same manner as in Section 8.2.1. The PID correction factors for signal and the numbers of $B\bar{B}$ meson pairs on MC and data are the same as in the case of control decay branching fraction measurement.

Finally, we can determine the branching fractions based on the calculations in Eq. (8.6). The obtained values are shown in Table 8.5. The measured value is almost twice as large as the MC value. The errors in all cases are statistical only. The result shows that the MC contribution to our simulated samples is underestimated and indicates that the branching fraction of the decay may be large enough to affect results in precision physics, in cases where it is ignored.

Table 8.5.: Signal decay branching fraction results on MC and in data.

8.3.3. Signal Distribution in bins of m_{KK}

It is possible to take a deeper look in the signal distribution over the m_{KK} variable by performing the signal fit in bins of the m_{KK} distribution, instead of a single fit over the whole region. This offers a deeper insight into the decay process and provides more details about the reliability of our MC samples. Results can be used to check theoretical predictions or to update existing MC generators. Table 8.6 shows the selected regions in m_{KK} , along with the corresponding signal yields on MC and in data. The results are graphically presented in Figure 8.10. Figures for each m_{KK} window fit can be found in Appendix C.1. In the given results, the m_{KK} regions around the ϕ and the D^0 resonances have been excluded (see Section 6.1).

Region	Expected	MC fit	Data fit
$0.980 < m_{KK} < 1.232$	44	42 ± 11	113 ± 29
$1.232 < m_{KK} < 1.483$	66	64 ± 12	185 ± 40
$1.483 < m_{KK} < 1.735$	67	67 ± 15	166 ± 47
$1.735 < m_{KK} < 1.987$	38	35 ± 9	87 ± 46
$1.987 < m_{KK} < 2.238$	24	26 ± 7	35 ± 25
$2.238 < m_{KK} < 2.490$	9	8 ± 4	-10 ± 13
$2.490 < m_{KK} < 2.742$	4	7 ± 4	-8 ± 7
$2.742 < m_{KK} < 2.993$	1	0 ± 1	-2 ± 5
$2.993 < m_{KK} < 3.245$	0	-3 ± 1	-2 ± 9
$3.245 < m_{KK} < 3.497$	0	-0 ± 1	-1 ± 6
$3.497 < m_{KK} < 3.748$	0	-0 ± 1	-2 ± 8
$3.748 < m_{KK} < 4.000$	0	-1 ± 1	-1 ± 224

Table 8.6.: Various signal fit yields for each of the defined m_{KK} windows.

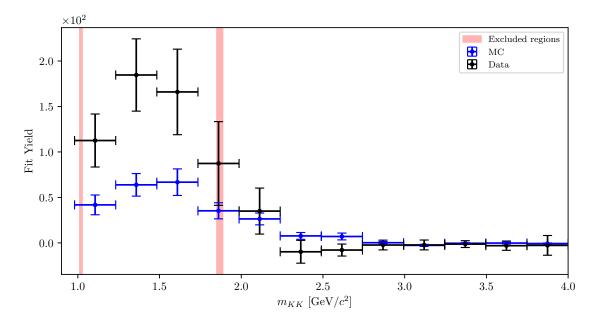


Figure 8.10.: Signal yield distribution as a function of m_{KK} .

8.3.4. Signal Distribution in bins of q^2

Similarly, we look at the signal distribution over the square momentum transfer to the lepton pair, q^2 , by performing the signal fit in bins of the q^2 distribution. In addition to the possibility of improved modeling of the MC generators, this enables the extraction of the V_{ub} parameter of the CKM matrix, provided that the appropriate form factors of the decay are available. Table 8.7 shows the selected regions in q^2 , along with the corresponding signal yields on MC and data. The results are graphically presented in Figure 8.11. Figures for each q^2 window fit can be found in Appendix C.2.

Region	Expected	MC fit	Data fit
$0.000 < q^2 < 1.500$	27	22 ± 11	-19 ± 31
$1.500 < q^2 < 3.000$	31	30 ± 4	25 ± 29
$3.000 < q^2 < 4.500$	30	30 ± 5	83 ± 32
$4.500 < q^2 < 6.000$	28	28 ± 6	124 ± 31
$6.000 < q^2 < 7.500$	31	29 ± 8	96 ± 31
$7.500 < q^2 < 9.000$	25	24 ± 8	83 ± 30
$9.000 < q^2 < 10.500$	24	22 ± 8	37 ± 28
$10.500 < q^2 < 12.000$	19	19 ± 5	57 ± 24
$12.000 < q^2 < 13.500$	16	15 ± 5	17 ± 18
$13.500 < q^2 < 15.000$	9	5 ± 3	15 ± 14
$15.000 < q^2 < 16.500$	5	3 ± 2	17 ± 3
$16.500 < q^2 < 18.000$	1	-3 ± 1	2 ± 4

Table 8.7.: Various signal fit yields for each of the defined q^2 windows.

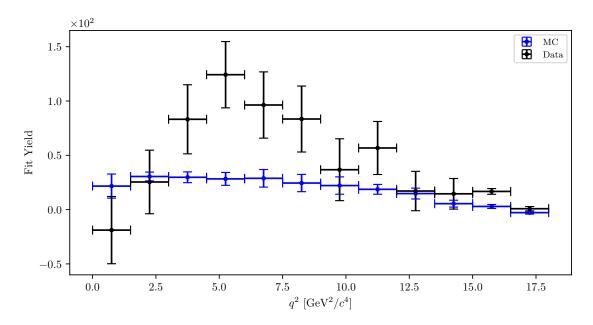


Figure 8.11.: Signal yield distribution as a function of q^2 .

The measured q^2 and m_{KK} distributions exhibit differences with respect to the MC models used. Differences are larger in the q^2 distribution, where the measured signal appears to fall more rapidly than in simulation in the region below 2.5 ${\rm GeV}^2/c^4$, and is more concentrated in the region of 2.5 ${\rm GeV}^2/c^4 < q^2 < 10~{\rm GeV}^2/c^4$. In m_{KK} , both data and simulation are concentrated in the region below 2 ${\rm GeV}/c^2$.

Chapter 9.

Systematic Uncertainty

In this chapter, the systematic errors of the analysis are discussed. These uncertainties arise due to various reasons, some of them being the difference between real and simulated data, or due to the nature of the approaches, taken in a specific analysis. Depending on their type, some uncertainties are generic and prepared beforehand, in order to be used in all analyses, while others are analysis specific and possible sources need to be thought through thoroughly.

9.1. Contributions to the Systematic Uncertainty

9.1.1. PID Efficiency Correction

The PID selection efficiency for the three charged particles in our signal decay needs to be corrected on MC due to various differences, when comparing to data. The Belle PID group has prepared a set of correction factors and tables of systematic uncertainties for PID efficiencies for all charged particles. In case of kaon ID and lepton ID, the tables are binned in experiment numbers, particle momentum, and in $\cos \theta$ of the particle direction, where, for each bin, a ratio of efficiencies between MC and data is provided, as well as the systematic errors. Each particle's correction factor and error is shown in Table 9.1, as well as the corresponding entry for all 3 particles. The entries are shown for both the signal and the control region, where the differences are related to the kinematic variations between the two decay modes.

The central values were obtained with a weighted average over all experiments, where 100% correlation for error calculation was assumed. A full correlation was also assumed when calculating the KK PID error, as both K use the same PID information.

The final PID efficiency systematic error on the full MC sample is determined to be

$$\sigma_{\rm sys}^{\rm PID} = 10, \quad \delta_{\rm sys}^{\rm PID} = 2.0\%,$$
 (9.1)

for the signal, as well as the control decay.

PID correction and systematic uncertainties	Control decay	Signal decay
Same sign K (w.r.t the B meson)	1.005 ± 0.009	1.007 ± 0.010
Opposite sign K (w.r.t the B meson)	1.004 ± 0.009	1.006 ± 0.009
e	0.977 ± 0.011	0.976 ± 0.011
μ	0.985 ± 0.009	0.980 ± 0.009
ℓ	0.981 ± 0.007	0.980 ± 0.007
KKe	0.986 ± 0.021	0.988 ± 0.022
$KK\mu$	0.994 ± 0.020	0.993 ± 0.021
$KK\ell$	0.991 ± 0.019	0.990 ± 0.020

Table 9.1.: PID correction factors and systematic uncertainties for various charged particles and their combinations.

9.1.2. Fit Bias and Binning Effects

Signal and background templates in our analysis are not perfectly distinct from one another and may potentially cause some over- or underestimation of the fitted signal yield. In order to study this problem, we estimate the bias from the binning study performed in Section 8.1, as well as the linearity test toy MC study in Section 7.3.2. The two bias functions describe a bias in each direction and are approximated as

$$f_{\min}(x) = -7.25 - 1.12x - \sigma_{f_{\min}}(x), \tag{9.2}$$

$$\sigma_{f_{\min}}(x) = \sqrt{0.050x^2 - 0.175x + 0.410},\tag{9.3}$$

$$f_{\text{max}}(N_b) = 6.86 + \sigma_{f_{\text{max}}}(N_b),$$
 (9.4)

$$\sigma_{f_{\text{max}}}(N_b) = \sqrt{0.004N_b^2 - 0.113N_b + 1.112}$$
(9.5)

where x represents the signal yield fraction of the data fit and N_b represents the binning choice of the fit. Values of 1σ intervals have been added to the bias functions, in order to be more conservative. The extracted signal yield in data, with the fit setup of $N_b = 19$ bins, was determined to be $N_{sig} = 491$, which leads to $x = N_{sig}/N_{sig}^{MC} = 491/249 \approx 2$. The bias interval is therefore

$$\sigma_{\text{sys}}^{\text{bias}} = ^{+7}_{-10}, \quad \delta_{\text{sys}}^{\text{bias}} = ^{+1.5\%}_{-2.0\%}.$$
 (9.6)

9.1.3. Gaussian Constraints

As mentioned in Section 8.3, it is possible to estimate the size of the systematic error of the Gaussian constraints. By fixing the constraints to the central values in Table 8.4, we obtain the pure statistical error, which can then be subtracted from the average fit error in order to determine the systematic uncertainty contribution, arising form the Gaussian constraints. Due to the nature of implementing the smearing of the ΔE variable, there is some randomness involved in our fits, so we perform 500 fits for the fixed and non-fixed case, in order to determine the uncertainties. The split errors are then

$$\bar{\sigma}_{\text{fit}} = 86, \tag{9.7}$$

$$\bar{\sigma}_{\text{stat}} = 81,$$
 (9.8)

$$\sigma_{\text{sys}}^{GC} = 26, \quad \delta_{\text{sys}}^{GC} = 5.4\%,$$
(9.9)

where GC stands for the Gaussian constraints. We see that the constrained channels are very well defined and introduce a relatively small level of uncertainty.

9.1.4. Fit Template Smearing and Offset

The smearing and offset of the ΔE variable was discussed in Section 7.1.1, where we estimated the central value of the parameters, as well as their range in the 1σ confidence interval. We have performed a study of the effects of different smearing and offset parameter values in data. From Section 7.1.1, the parameter values are

- Smearing: 40^{+15}_{-17} MeV,
- Offset: $6^{+4.6}_{-6}$ MeV.

Since the two parameters are largely uncorrelated, we are able to perform the study in the form of signal fits with four different combinations of parameters in the given range. For each parameter setting, the Gaussian constraints are fixed and 500 fits are performed to obtain the following results

- Result: $\bar{N}_{sig} = 458$, • set: [smearing, offset]: [23 MeV,6 MeV],
- set: [smearing, offset]: [55 MeV,6 MeV], Result: $\bar{N}_{sig} = 532$.
- set: [smearing, offset]: [40 MeV,0 MeV], Result: $\bar{N}_{sig} = 532$,
- set: [smearing, offset]: [40 MeV,12.6 MeV], Result: $\bar{N}_{sig} = 460$,

which results in the following estimate of systematic uncertainties for smearing and offset parameters

$$\sigma_{\text{sys}}^{\text{sm.}} = {}^{+41}_{-33}, \quad \delta_{\text{sys}}^{\text{sm.}} = {}^{+8.3\%}_{-6.7\%},$$

$$\sigma_{\text{sys}}^{\text{off.}} = {}^{+41}_{-31}, \quad \delta_{\text{sys}}^{\text{off.}} = {}^{+8.4\%}_{-6.3\%}.$$
(9.10)

$$\sigma_{\text{sys}}^{\text{off.}} = {}^{+41}_{-31}, \quad \delta_{\text{sys}}^{\text{off.}} = {}^{+8.4\%}_{-6.3\%}.$$
 (9.11)

9.1.5. Effects of a Finite MC sample

The shape of signal and backgrounds templates in our analysis is fixed and only their normalization is considered as a floating parameter in the fit. Due to the finite size of the MC sample, the template shape introduces an additional source of uncertainty, as it may differ if produced in a separate, equal-sized MC sample. Since the bins in these 2D histogram templates are statistically independent, we take the content of each bin and vary the values according to the Poisson distribution. This procedure is repeated 500 times with the Gaussian constraints fixed to the central values in Table 8.4 and with a fixed random seed for the application of smearing in ΔE . To estimate the size of this uncertainty, we take the width of the fit yield distribution. The resulting finite MC sample contribution of the systematic uncertainty is

$$\sigma_{\text{sys}}^{MC} = 26, \quad \delta_{\text{sys}}^{MC} = 5.3\%.$$
 (9.12)

9.1.6. MVA Selection Efficiencies

Control sample fits allow us to check the behavior of the optimized MVA selection on MC, as well as in data, and see if any of the MVA steps introduce a possible disagreement between the two. We compare control yields, their ratios, and ratios of selection efficiencies (double ratios). The following selection criteria scenarios are studied

- (a) final selection before any MVA step,
- (b) (a) $+ BDT_{q\bar{q}}$ cut,
- (c) (a) + $uBDT_{B\bar{B}}$ cut,
- (d) (a) $+ BDT_{a\bar{a}} + uBDT_{B\bar{B}}$ cut (final selection).

The results for control fit yields, their ratios, and double ratios are shown in Figure 9.1. The plot shows that the yield ratios and selection efficiency ratios are consistent with 1. This means that data and MC are in agreement before, as well as after applying the final selection. This is an important check since the behavior of our analysis on the control sample suggests that the final selection is not over-optimized to signal MC.

We estimate the systematic error, due to the MVA selection steps, as the standard deviation of double ratio entries around the nominal values, for each step in the MVA selection, except for the results coming from measurements with specific lepton states. The systematic error estimate for this contribution is

$$\sigma_{\text{sys.}}^{\text{MVA}} = 5, \quad \sigma_{\text{sys.}}^{\text{MVA}} = 1.0\%.$$
 (9.13)

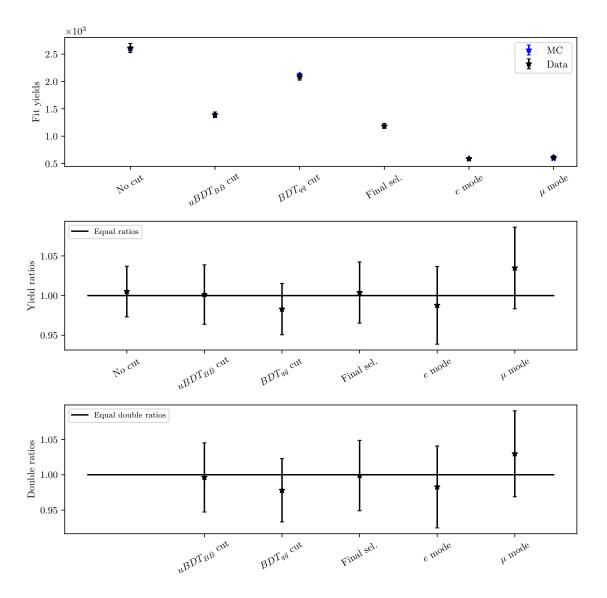


Figure 9.1.: Fit yields, their ratios and ratios of selection efficiencies (double ratios) for the control sample fits in data and on MC.

9.1.7. Model Uncertainty Effects

The signal decay model used in the MC generation is ISGW2 [18], which is known to result in unrealistic predictions and poor agreement with data, so it is not the most precise model for our signal MC sample. Due to this model unreliability, our analysis has been set up as model-independent as possible via means of not using variables, which exhibit model dependence. Such variables are, for example, the squared momentum transfer to the lepton pair (q^2) , the invariant mass of the two kaon daughters (m_{KK}) , or decay angle between any two charged particles in the final state.

In order to test the effects of model dependency on our final result, we prepare three additional signal MC samples, produced with three extreme scenarios of decay model choice. In the first scenario, we generate the signal MC sample with a generic phase-space decay mode, PHSP [10], which results in a continuum-like distributions

of q^2 and m_{KK} . In the other scenario, only resonant-like contributions of m_{KK} are used. The third scenario is obtained by sampling the first model in a non-uniform way, so that the final distribution is more dominant in the higher region of the q^2 distribution. These scenarios act as extreme cases of decay model choice and present a reasonable, if not a conservative measure of the model uncertainty.

Shape Effects of the Model

The shape effect is evaluated by using signal templates, constructed from different signal MC samples. We perform 500 fits for each case, with the Gaussian constraints fixed to the central values, presented in Table 8.4. The differences between the mean values of fit yields serve as an estimate of the model uncertainty. Figure 9.2 shows the generated m_{KK} and q^2 distributions of the three mentioned decay models, as well as distributions of ΔE and M_{BC} after the final selection. Figures show a good agreement of ΔE and M_{BC} for different cases of the generator model.

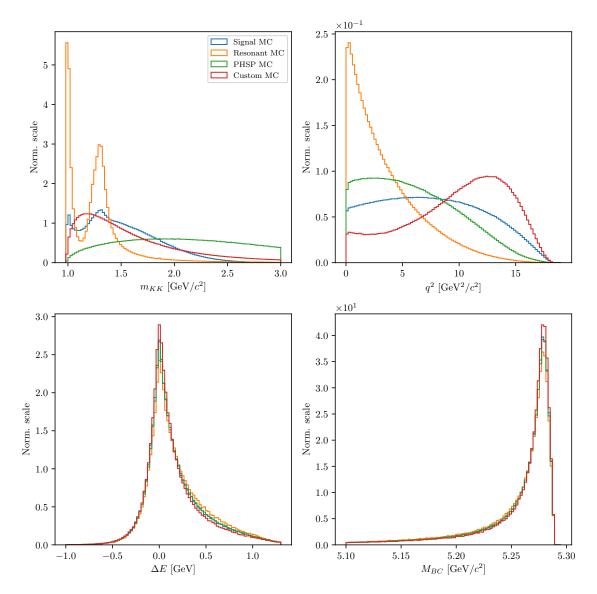


Figure 9.2.: m_{KK} (top left), q^2 (top right), ΔE (bottom left), and M_{BC} (bottom right) for the main and the three extreme cases of signal MC generator choice.

The resulting average signal yields for the three model choices are

$$\bar{N}_{sig} = 491,$$
 (9.14)

$$\bar{N}_{siq}^{\text{res.}} = 494,$$
 (9.15)

$$\bar{N}_{sig}^{\text{PHSP}} = 536, \tag{9.16}$$

We see that, in general, the model with a worse resolution in ΔE and M_{BC} will likely result in a larger yield, and vice versa for a model with a better resolution. Overall, these models give a conservative estimation of the systematic uncertainty with a value of

$$\sigma_{\text{sys}}^{\text{mod.}} = ^{+45}_{-39}, \quad \delta_{\text{sys}}^{\text{mod.}} = ^{+9.3\%}_{-8.0\%}.$$
 (9.18)

Model Efficiency

The second way that a model can affect our analysis is the efficiency of the model. Due to the different properties of the model, we can expect different efficiencies in model dependent variables like m_{KK} and q^2 . The efficiency as a function of these variables is shown in Figure 9.3 for all discussed generator models. In a perfect scenario, a model-independent analysis should produce a flat efficiency with respect to the model dependent variables. While this is rarely achieved in practice, the overall discrepancy of the efficiency functions from very different models is relatively small.

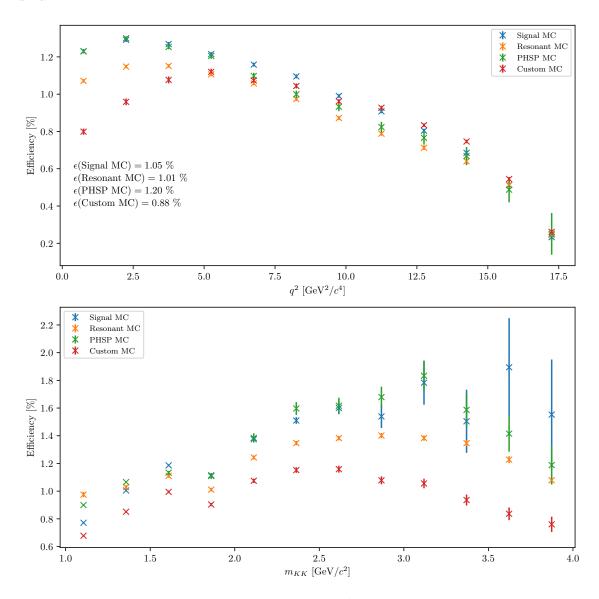


Figure 9.3.: Efficiency as a function of q^2 (top) and m_{KK} (bottom) for the main and three extreme cases of signal MC generator choice.

To estimate the systematic uncertainty due to the efficiency effects of the model, we compute the overall efficiencies and take the extreme cases of relative differences with regard to the main signal MC model. This results in

$$\sigma_{\text{sys}}^{\text{mod.}} = {}^{+70}_{-79}, \quad \delta_{\text{sys}}^{\text{mod.}} = {}^{+14.3\%}_{-16.2\%}.$$
 (9.19)

As mentioned in Section 8.3.4, the measured q^2 and m_{KK} distributions differ to some extent from the simulated ones. Since the reconstruction efficiency is not constant as a function of these variables, this could present an additional source of the systematic uncertainty. However, in m_{KK} , the measured signal is concentrated in the region of $m_{KK} < 2 \text{ GeV}/c^2$, where the efficiency is relatively constant and within the range of the already estimated systematic uncertainty of around $\pm 15\%$. In q^2 , the data distribution is wider and encompasses the region of a more pronounced efficiency dependence. We estimated the branching fraction using the signal yield as obtained in bins of q^2 (see Table 8.7) and the efficiency as a function of q^2 . The result, although with a larger statistical uncertainty, due to low statistics in q^2 bins, is within the assigned systematic uncertainty, and therefore consistent with the branching fraction, found using the average reconstruction efficiency.

9.2. Summary of Systematic Uncertainties

The summary of all systematic uncertainties is shown in Table 9.2. The full estimate of the systematic uncertainty is summed up in quadrature and applied to the result in Section 8.3.2.

Source	σ	δ [%]
PID	10	2.0
Fit Bias	$+7 \\ -10$	$\begin{array}{c c} +1.5 \\ -2.0 \end{array}$
Gaussian Constraints	26	5.4
Template Smearing	$+41 \\ -33$	$\begin{array}{ c c c c c } +8.3 \\ -6.7 \end{array}$
Template Offset	$\begin{array}{ c c c c c } +41 \\ -31 \end{array}$	$\begin{vmatrix} +8.4 \\ -6.3 \end{vmatrix}$
Finite MC Effects	26	5.3
MVA Selection	5	1.0
Model Shape	$+45 \\ -39$	$\begin{vmatrix} +9.3 \\ -8.0 \end{vmatrix}$
Model Efficiency	$\begin{vmatrix} +70 \\ -79 \end{vmatrix}$	$\begin{vmatrix} +14.3 \\ -16.2 \end{vmatrix}$
Total	$\begin{vmatrix} +109 \\ -107 \end{vmatrix}$	$\begin{vmatrix} +22.2 \\ -21.9 \end{vmatrix}$

Table 9.2.: Summary of systematic uncertainties in this analysis.

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Appendix A.

ROE MVA Control Plots

A.1. ROE Clean-up π^0 Training

A.1.1. Variable Importance

	Name	Alias	Importance
0	chiProb	$ v_0 $	0.280
1	useCMSFrame(daughterAngleInBetween(0,1))	v_1	0.203
2	<pre>daughter(0,useCMSFrame(p))</pre>	v_2	0.073
3	InvM	v_3	0.072
4	<pre>daughter(1,clusterHighestE)</pre>	v_4	0.061
5	daughter(1,clusterTheta)	v_5	0.049
6	daughter(1,p)	v_6	0.047
7	<pre>daughter(0,clusterHighestE)</pre>	v_7	0.029
8	daughter(0,clusterTheta)	v_8	0.024
9	daughter(0,clusterE9E25)	v_9	0.018
10	daughter(0,minC2HDist)	v_{10}	0.018
11	daughter(1,minC2HDist)	v_{11}	0.017
12	daughter(1,clusterE9E25)	v_{12}	0.016
13	useRestFrame(daughterAngleInBetween(0,1))	v_{13}	0.014
14	daughter(1,clusterNHits)	v_{14}	0.013
15	daughter(0,clusterNHits)	v_{15}	0.011
16	daughter(0,clusterErrorE)	v_{16}	0.009
17	daughter(1,clusterErrorE)	v_{17}	0.009
18	SigMBF	v_{18}	0.007
19	useCMSFrame(p)	v_{19}	0.006
20	daughter(0,p)	v_{20}	0.005
21	SigM	v_{21}	0.005
22	<pre>daughter(1,useCMSFrame(p))</pre>	v_{22}	0.005
23	useLabFrame(daughterAngleInBetween(0,1))	v_{23}	0.005
24	p	v_{24}	0.003

Table A.1.: Variable names, aliases and importance in the scope of π^0 MVA training for ROE clean-up.

A.1.2. Variable Distributions

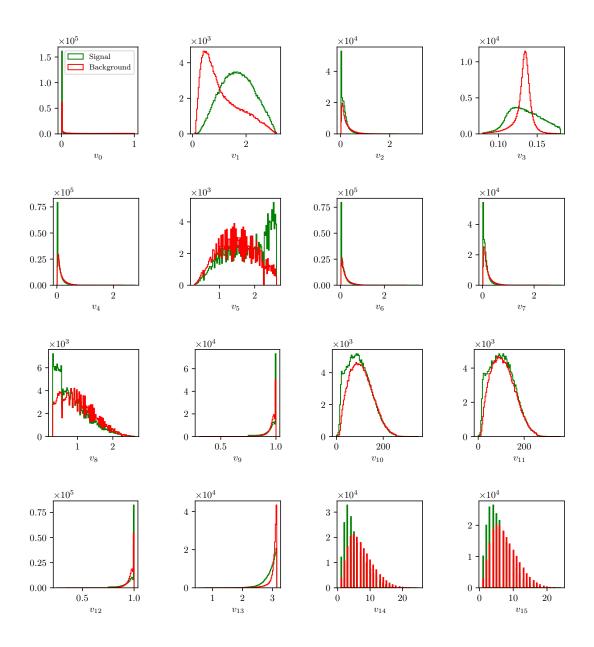


Figure A.1.: Feature distributions for MVA training of π^0 candidates in the scope of ROE clean-up.

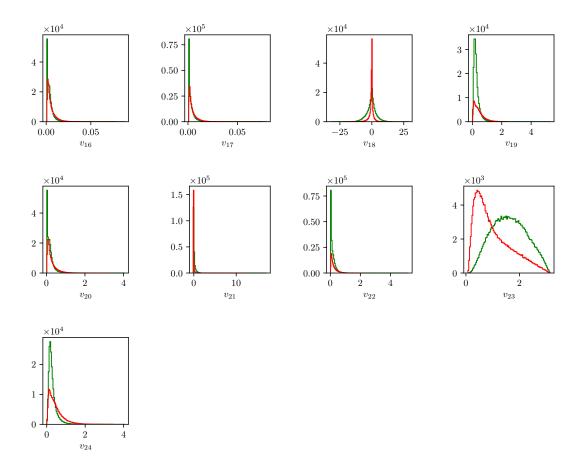


Figure A.1.: Feature distributions for MVA training of π^0 candidates in the scope of ROE clean-up.

A.1.3. Hyper-parameter Optimization

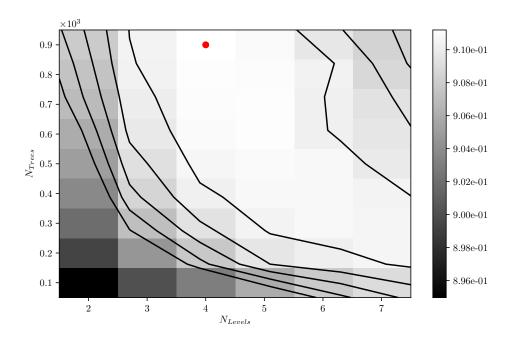


Figure A.2.: Hyper-parameter optimization of nTrees and nLevels in the BDT forest training of π^0 candidates in the scope of the ROE clean-up.

A.1.4. Results

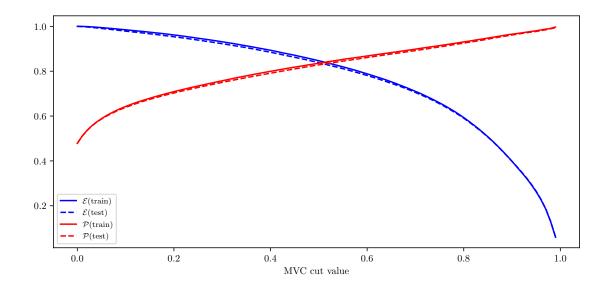


Figure A.3.: Efficiency (\mathcal{E}) and purity (\mathcal{P}) of the MVA classifier output for π^0 candidates training on the train (solid) and test (dashed) samples.

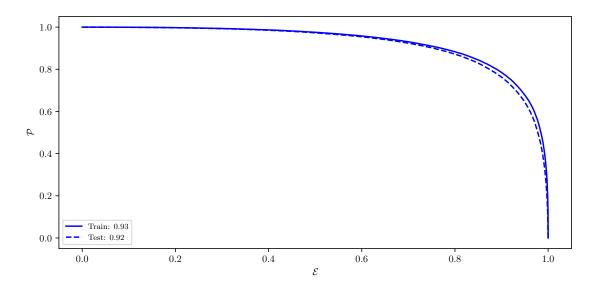


Figure A.4.: ROC curves of the MVA classifier output for π^0 candidates training on the train (solid) and test (dashed) samples.

A.2. ROE Clean-up γ Training

A.2.1. Variable Importance

	Name	Alias	Importance
0	p	v_0	0.327
1	pi0p	v_1	0.243
2	clusterHighestE	v_2	0.226
3	minC2HDist	v_3	0.052
4	cosTheta	v_4	0.036
5	clusterE9E25	v_5	0.031
6	clusterNHits	v_6	0.025
7	clusterUncorrE	v_7	0.022
8	clusterR	v_8	0.015
9	useCMSFrame(p)	v_9	0.013
10	clusterErrorE	v_{10}	0.010
11	clusterReg	v_{11}	0.000

Table A.2.: Variable names, aliases and importance in the scope of γ MVA training for ROE clean-up.

A.2.2. Variable Distributions

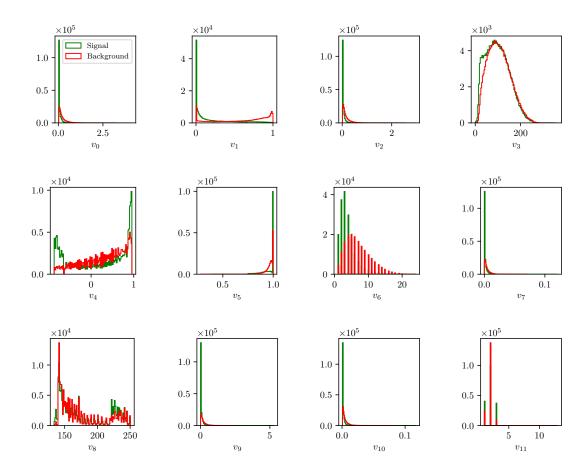


Figure A.5.: Feature distributions for MVA training of γ candidates in the scope of ROE clean-up.

A.2.3. Hyper-parameter Optimization

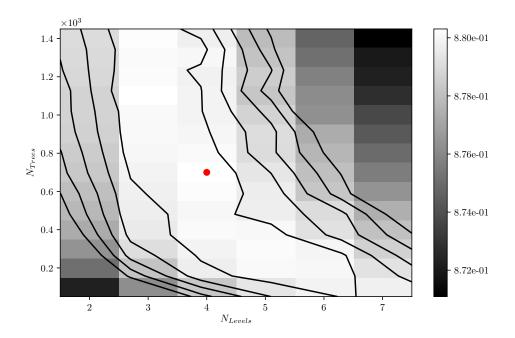


Figure A.6.: Hyper-parameter optimization of nTrees and nLevels in the BDT forest training of γ candidates in the scope of the ROE clean-up.

A.2.4. Results

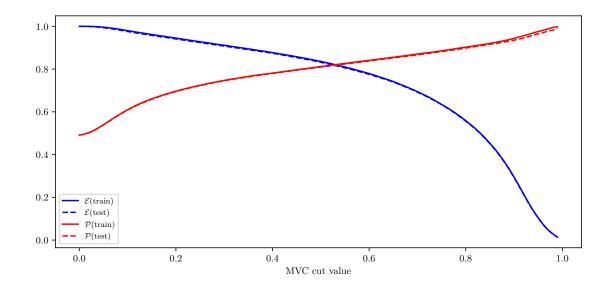


Figure A.7.: Efficiency (\mathcal{E}) and purity (\mathcal{P}) of the MVA classifier output for γ candidates training on the train (solid) and test (dashed) samples.

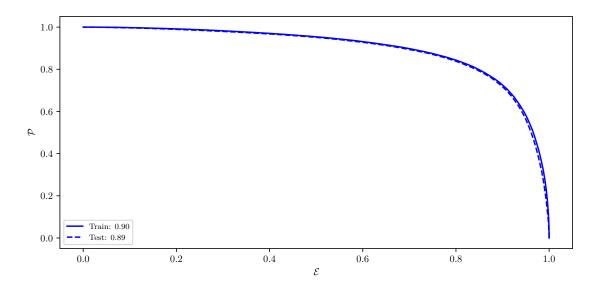


Figure A.8.: ROC curves of the MVA classifier output for γ candidates training on the train (solid) and test (dashed) samples.

A.3. ROE Clean-up Duplicate Pair Training

A.3.1. Variable Importance

	Name	Alias	Importance
0	useCMSFrame(daughterAngleInBetween(0,1))	$ v_0 $	0.132
1	daughter(0,phi0Err)	v_1	0.082
2	useLabFrame(daughterAngleInBetween(0,1))	v_2	0.055
3	daughter(1,d0)	v_3	0.051
4	daughter(1,phi0Err)	v_4	0.051
5	daughter(0,d0)	v_5	0.050
6	daughter(1,nCDCHits)	v_6	0.040
7	daughter(1,d0Err)	v_7	0.037
8	daughter(0,nCDCHits)	v_8	0.034
9	daughter(1,z0)	v_9	0.032
10	daughter(0,z0)	v_{10}	0.030
11	daughter(0,d0Err)	v_{11}	0.028
12	daughter(0,nSVDHits)	v_{12}	0.028
13	daughter(1,pz)	v_{13}	0.027
14	<pre>daughter(1,useCMSFrame(p))</pre>	v_{14}	0.024
15	extraInfo(decayModeID)	v_{15}	0.023
16	daughter(0,pz)	v_{16}	0.020
17	daughter(1,nSVDHits)	v_{17}	0.020
18	daughter(0,pValue)	v_{18}	0.020
19	daughter(1,tanlambda)	v_{19}	0.018
20	daughter(1,pValue)	v_{20}	0.018
21	daughter(0,tanlambda)	v_{21}	0.017
22	daughter(0,phi0)	v_{22}	0.016

23	daughter(1,phi0)	v_{23}	0.016
24	<pre>daughter(0,useCMSFrame(p))</pre>	v_{24}	0.015
25	daughter(0,z0Err)	v_{25}	0.014
26	daughter(1,omega)	v_{26}	0.013
27	daughter(0,omega)	v_{27}	0.013
28	daughter(1,z0Err)	v_{28}	0.012
29	daughter(0,pt)	v_{29}	0.011
30	<pre>daughter(0,omegaErr)</pre>	v_{30}	0.011
31	<pre>daughter(1,omegaErr)</pre>	v_{31}	0.010
32	daughter(1,pt)	v_{32}	0.009
33	<pre>daughter(0,tanlambdaErr)</pre>	v_{33}	0.009
34	<pre>daughter(1,tanlambdaErr)</pre>	v_{34}	0.009
35	<pre>useRestFrame(daughterAngleInBetween(0,1))</pre>	v_{35}	0.003
36	daughter(1,charge)	v_{36}	0.000
37	daughter(0,charge)	v_{37}	0.000

Table A.3.: Variable names, aliases and importance in the scope of duplicate track pair MVA training for ROE clean-up.

A.3.2. Variable Distributions

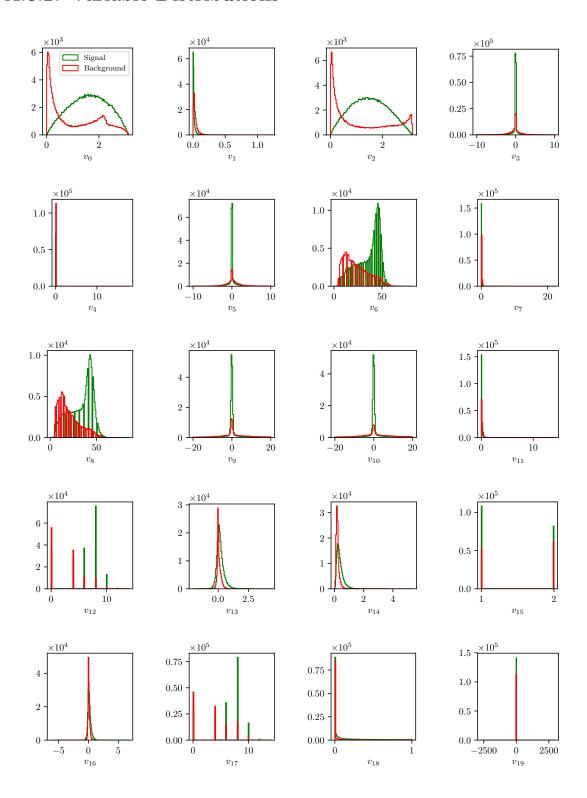


Figure A.9.: Feature distributions for MVA training of duplicate track pair candidates in the scope of ROE clean-up.

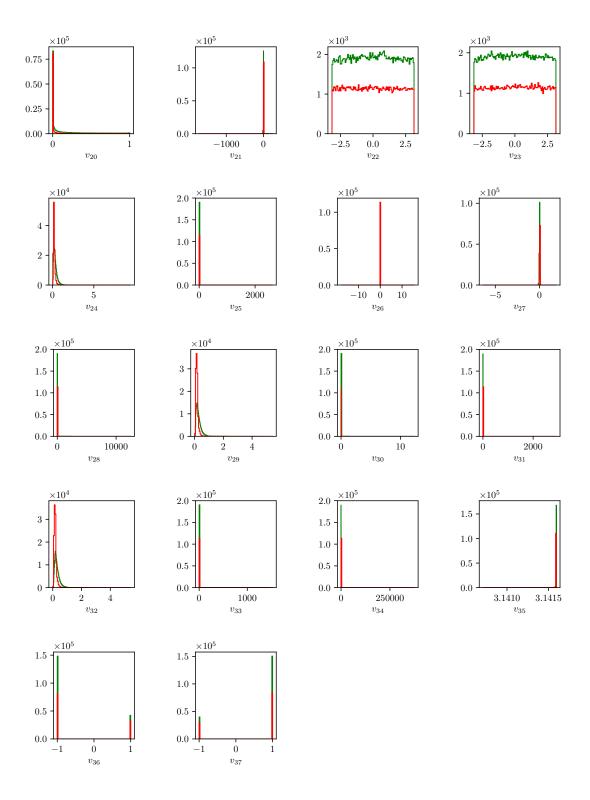


Figure A.9.: Feature distributions for MVA training of duplicate track pair candidates in the scope of ROE clean-up.

A.3.3. Hyper-parameter Optimization

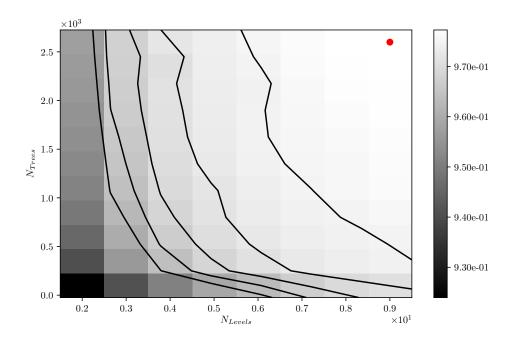


Figure A.10.: Hyper-parameter optimization of nTrees and nLevels in the BDT forest training of duplicate track pair candidates in the scope of the ROE clean-up.

A.3.4. Results

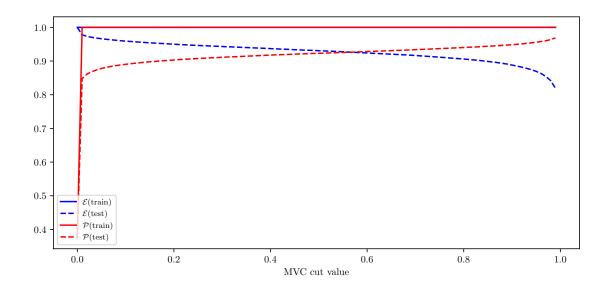


Figure A.11.: Efficiency (\mathcal{E}) and purity (\mathcal{P}) of the MVA classifier output for duplicate track pair candidates training on the train (solid) and test (dashed) samples.

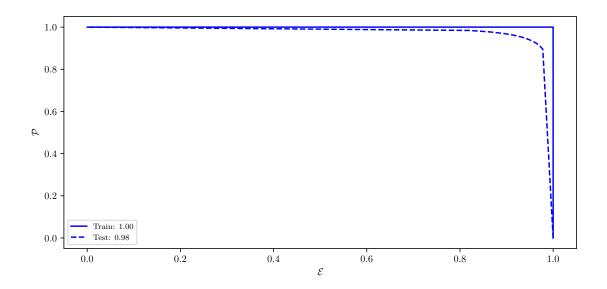


Figure A.12.: ROC curves of the MVA classifier output for duplicate track pair candidates training on the train (solid) and test (dashed) samples.

A.4. ROE Clean-up Duplicate Track Training

A.4.1. Variable Importance

	Name	Alias	Importance
0	extraInfo(d0Diff)	v_0	0.214
1	extraInfo(z0Diff)	v_1	0.087
2	d0	v_2	0.069
3	<pre>extraInfo(pValueDiff)</pre>	v_3	0.060
4	z0	v_4	0.058
5	phi0Err	v_5	0.056
6	<pre>extraInfo(pzDiff)</pre>	v_6	0.055
7	<pre>extraInfo(ptDiff)</pre>	v_7	0.045
8	z0Err	v_8	0.043
9	<pre>extraInfo(nCDCHitsDiff)</pre>	v_9	0.037
10	<pre>extraInfo(nSVDHitsDiff)</pre>	v_{10}	0.034
11	pt	v_{11}	0.032
12	d0Err	v_{12}	0.030
13	pValue	v_{13}	0.029
14	nCDCHits	v_{14}	0.028
15	nSVDHits	v_{15}	0.028
16	pz	v_{16}	0.025
17	cosTheta	v_{17}	0.024
18	phi0	v_{18}	0.023
19	useCMSFrame(p)	v_{19}	0.021

Table A.4.: Variable names, aliases and importance in the scope of duplicate track MVA training for ROE clean-up.

A.4.2. Variable Distributions

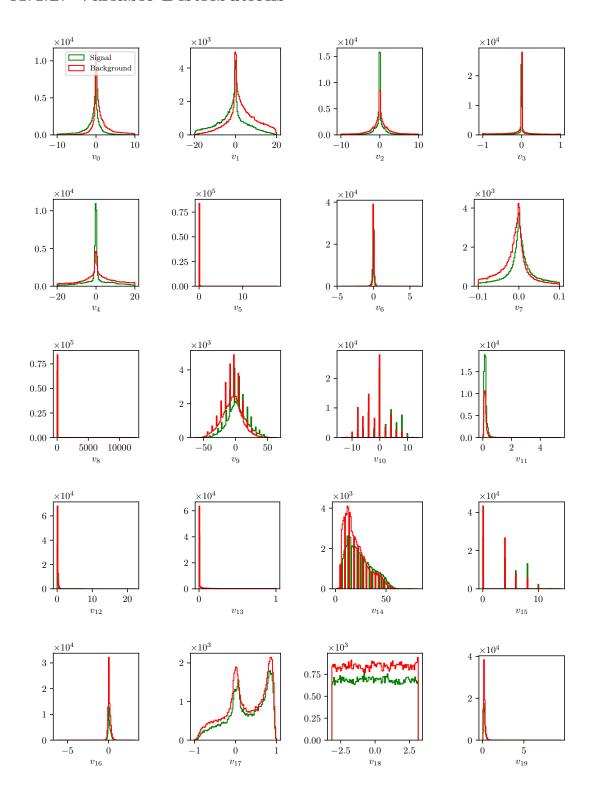


Figure A.13.: Feature distributions for MVA training of duplicate track candidates in the scope of ROE clean-up.

A.4.3. Hyper-parameter Optimization

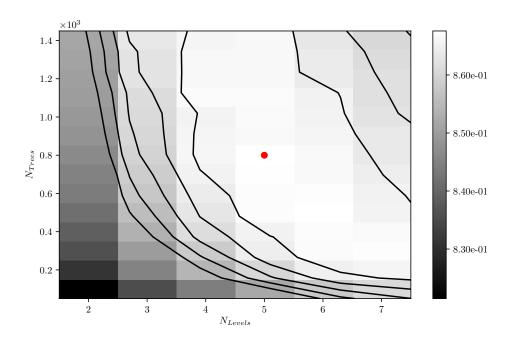


Figure A.14.: Hyper-parameter optimization of nTrees and nLevels in the BDT forest training of duplicate track candidates in the scope of the ROE clean-up.

A.4.4. Results

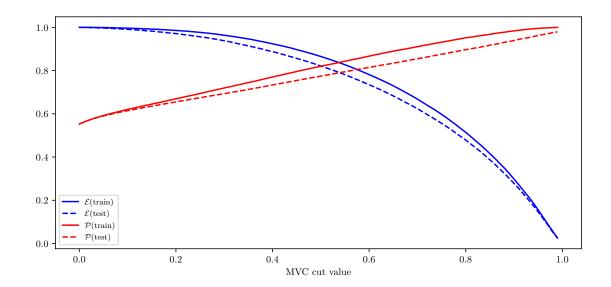


Figure A.15.: Efficiency (\mathcal{E}) and purity (\mathcal{P}) of the MVA classifier output for duplicate track candidates training on the train (solid) and test (dashed) samples.

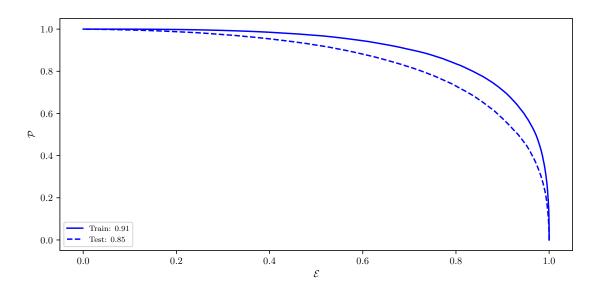


Figure A.16.: ROC curves of the MVA classifier output for duplicate track candidates training on the train (solid) and test (dashed) samples.

Appendix B.

MVA Control Plots

B.1. $q\bar{q}$ Suppression Training

B.1.1. Variable Importance

	Name	Alias	Importance
0	B_CosTBTO	v_0	0.291
1	B_hso02	v_1	0.096
2	B_ThrustB	v_2	0.096
3	B_roeFit_dz	v_3	0.075
4	B_R2	v_4	0.054
5	B_hso12	v_5	0.044
6	B_hoo2	v_6	0.032
7	B_ThrustO	v_7	0.027
8	B_qpKaon	v_8	0.024
9	B_cc2_CcROE	v_9	0.023
10	B_hoo0	v_{10}	0.019
11	B_cc3_CcROE	v_{11}	0.019
12	B_cc4_CcROE	v_{12}	0.016
13	B_CosTBz	v_{13}	0.015
14	B_hso01	v_{14}	0.015
15	B_cc1_CcROE	v_{15}	0.015
16	B_cc5_CcROE	v_{16}	0.013
17	B_cc6_CcROE	v_{17}	0.012
18	$B_{-}qpFastHadron$	v_{18}	0.012
19	B_cc7_CcROE	v_{19}	0.010
20	B_cc9_CcROE	v_{20}	0.010
21	B_cc8_CcROE	v_{21}	0.010
22	$B_{-}qpMaximumPstar$	v_{22}	0.008
23	B_hso10	v_{23}	0.008
24	B_hso04	v_{24}	0.007
25	$B_{-}qpLambda$	v_{25}	0.006
26	B_hoo1	v_{26}	0.006
27	$B_{-}qpKaonPion$	v_{27}	0.006
28	B_hoo4	v_{28}	0.006
29	$B_qpSlowPion$	v_{29}	0.006
30	B_hso03	v_{30}	0.005
31	B_hso14	v_{31}	0.004

32	B_qpFSC	v_{32}	0.004
33	B_hoo3	v_{33}	0.004

Table B.1.: Variable names, aliases and importance in the scope of $q\bar{q}$ suppression MVA training.

B.1.2. Variable Distributions

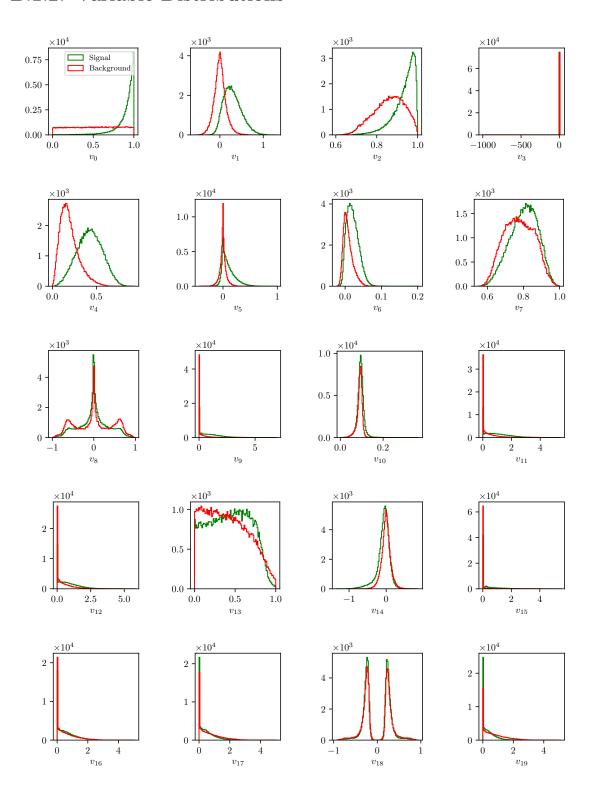


Figure B.1.: Feature distributions for MVA training of $q\bar{q}$ background suppression.

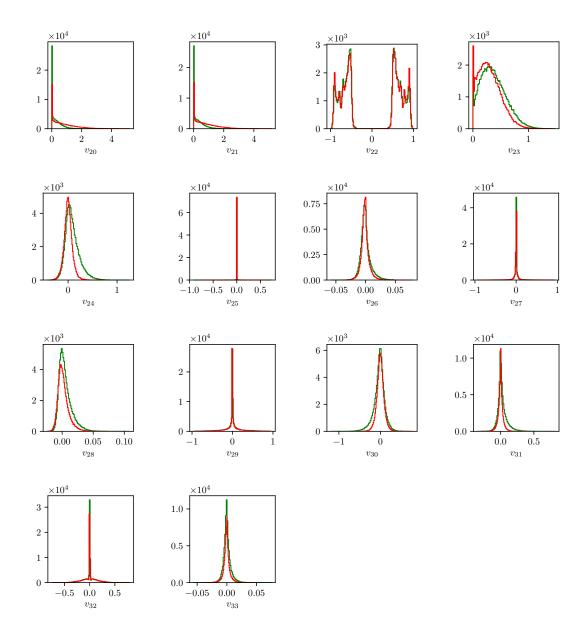


Figure B.1.: Feature distributions for MVA training of $q\bar{q}$ background suppression.

B.1.3. Hyper-parameter Optimization

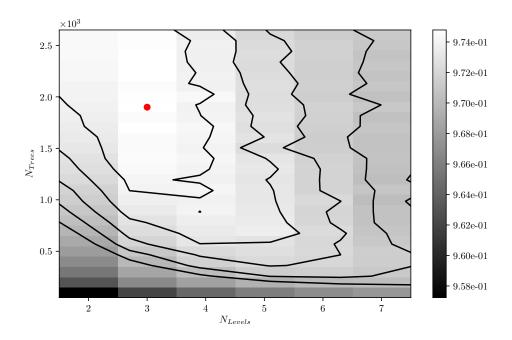


Figure B.2.: Hyper-parameter optimization of nTrees and nLevels in the BDT forest training of $q\bar{q}$ background suppression.

B.1.4. Results

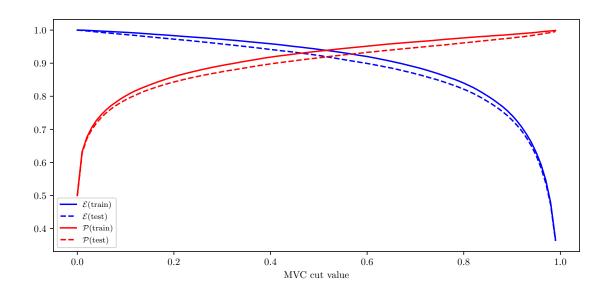


Figure B.3.: Efficiency (\mathcal{E}) and purity (\mathcal{P}) of the MVA classifier output for $q\bar{q}$ background suppression training on the train (solid) and test (dashed) samples.

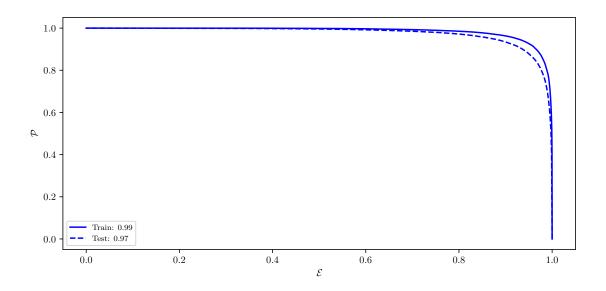


Figure B.4.: ROC curves of the MVA classifier output for $q\bar{q}$ background suppression training on the train (solid) and test (dashed) samples.

B.2. Standard $B\bar{B}$ Suppression Training

B.2.1. Variable Importance

	Name	Alias	Importance
0	B_cosMomVtxKKlnu	v_0	0.372
1	B_ROE_PThetacms0	v_1	0.096
2	B_nROETrkO	v_2	0.079
3	B_K1FT	v_3	0.063
4	$B_{-}cosBY$	v_4	0.051
5	B_roeFit_dz	v_5	0.047
6	B_xiZ0	v_6	0.043
7	$B_cosMomVtx$	v_7	0.038
8	B_{-} chi P rob	v_8	0.031
9	$B_nKaonInROE$	v_9	0.028
10	$B_{missM2Veto1}$	v_{10}	0.026
11	$B_{missM2Veto2}$	v_{11}	0.021
12	$B_nROEDistTrk$	v_{12}	0.018
13	$B_cosMomVtxKK$	v_{13}	0.018
14	B_KOFT	v_{14}	0.017
15	B_{QVeto1}	v_{15}	0.016
16	$B_{missM20}$	v_{16}	0.015
17	B_{-} Tag $VPvalue$	v_{17}	0.012
18	B_QVeto2	v_{18}	0.010

Table B.2.: Variable names, aliases and importance in the scope of $B\bar{B}$ background suppression.

B.2.2. Variable Distributions

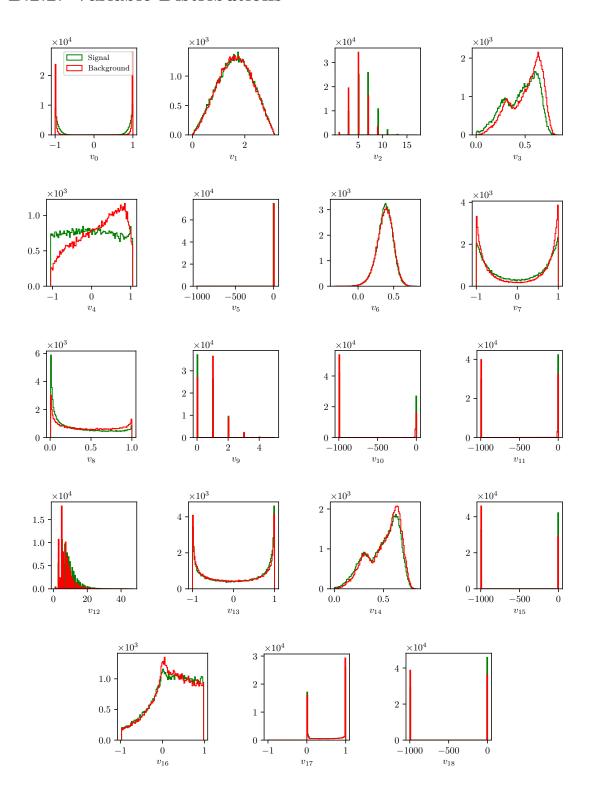


Figure B.5.: Feature distributions for MVA training of $B\bar{B}$ background suppression.

B.2.3. Hyper-parameter Optimization

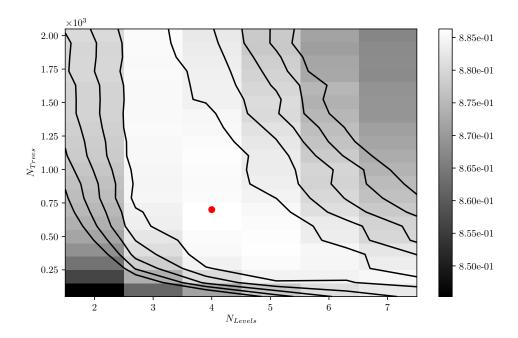


Figure B.6.: Hyper-parameter optimization of nTrees and nLevels in the BDT forest training of $B\bar{B}$ background suppression.

B.2.4. Results

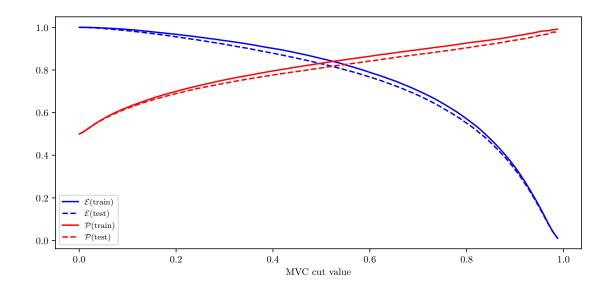


Figure B.7.: Efficiency (\mathcal{E}) and purity (\mathcal{P}) of the MVA classifier output for $B\bar{B}$ background suppression training on the train (solid) and test (dashed) samples.

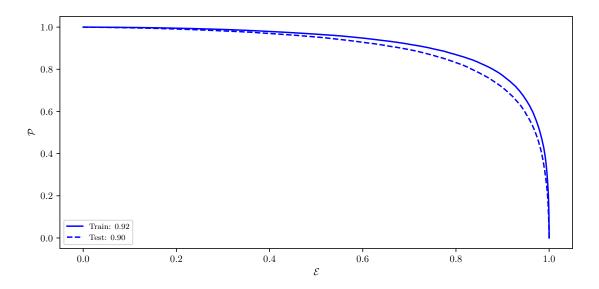


Figure B.8.: ROC curves of the MVA classifier output for $B\bar{B}$ background suppression training on the train (solid) and test (dashed) samples.

B.3. Uniformity Boosted $B\bar{B}$ Suppression Training

B.3.1. Hyper-parameter Optimization

Hyper-parameters were not optimized due to the large CPU time consumption of the algorithm. The following set up of the hyper-parameters was chosen

• nTrees: 300

• nLevels: 4

B.3.2. Results

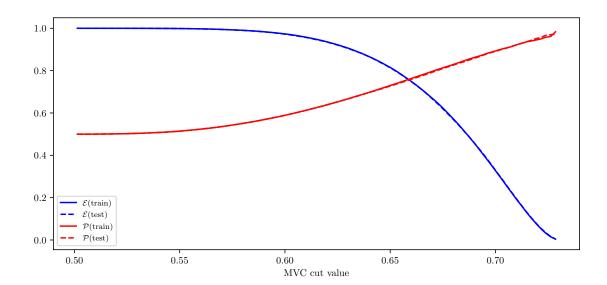


Figure B.9.: Efficiency (\mathcal{E}) and purity (\mathcal{P}) of the uniformity boosted MVA classifier output for $B\bar{B}$ background suppression training on the train (solid) and test (dashed) samples.

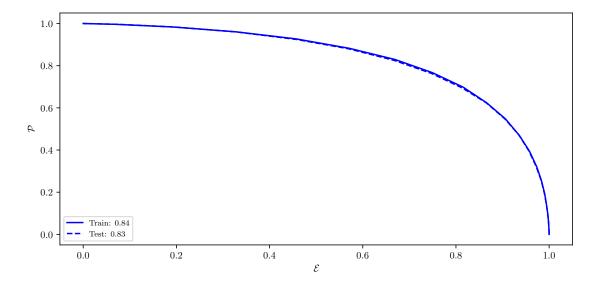


Figure B.10.: ROC curves of the uniformity boosted MVA classifier output for $B\bar{B}$ background suppression training on the train (solid) and test (dashed) samples.

Appendix C.

Other Plots

C.1. Signal Fits in m_{KK}

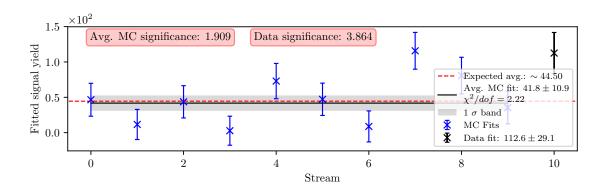


Figure C.1.: Signal fit result for the 1st m_{KK} window for MC and data in the range $0.980 < m_{KK} < 1.232$.

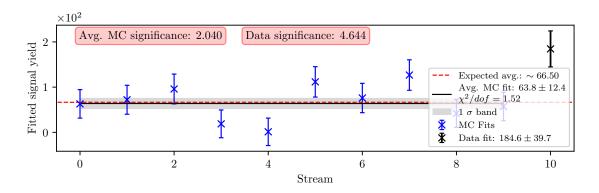


Figure C.2.: Signal fit result for the $2^{\rm nd}$ m_{KK} window for MC and data in the range $1.232 < m_{KK} < 1.483$.

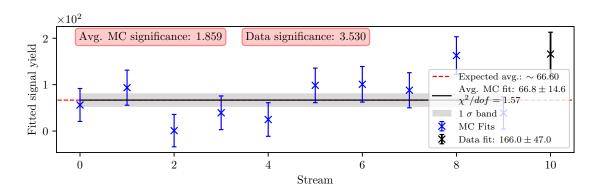


Figure C.3.: Signal fit result for the $3^{\rm rd}$ m_{KK} window for MC and data in the range $1.483 < m_{KK} < 1.735$.

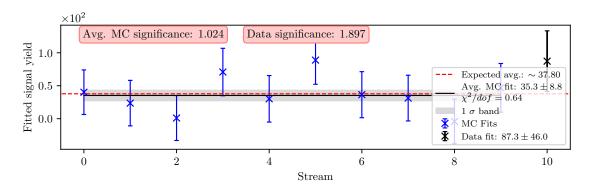


Figure C.4.: Signal fit result for the 4th m_{KK} window for MC and data in the range 1.735 $< m_{KK} < 1.987$.

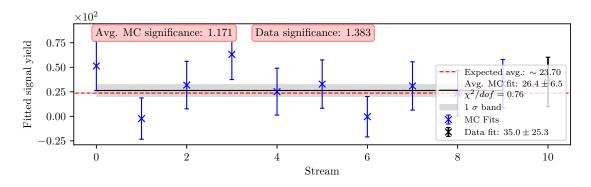


Figure C.5.: Signal fit result for the 5th m_{KK} window for MC and data in the range $1.987 < m_{KK} < 2.238$.

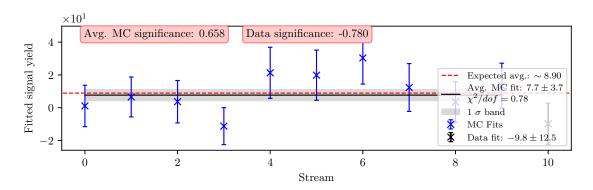


Figure C.6.: Signal fit result for the 6th m_{KK} window for MC and data in the range $2.238 < m_{KK} < 2.490$.

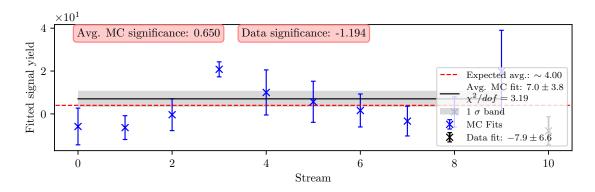


Figure C.7.: Signal fit result for the $7^{\rm th}$ m_{KK} window for MC and data in the range $2.490 < m_{KK} < 2.742$.

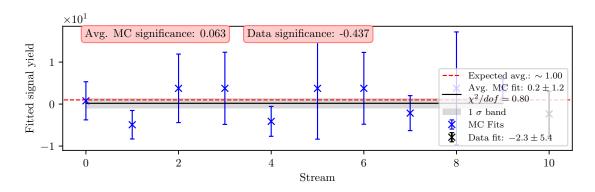


Figure C.8.: Signal fit result for the $8^{\rm th}$ m_{KK} window for MC and data in the range $2.742 < m_{KK} < 2.993$.

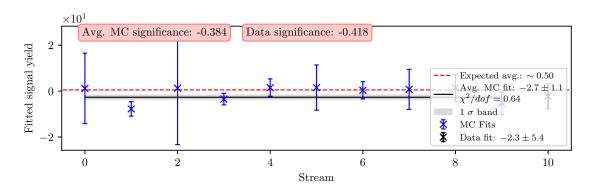


Figure C.9.: Signal fit result for the 9th m_{KK} window for MC and data in the range $2.993 < m_{KK} < 3.245$.

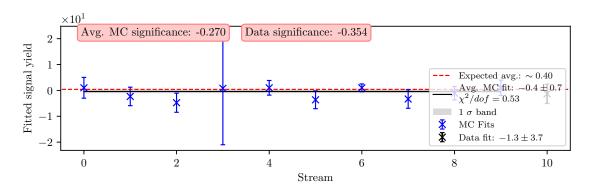


Figure C.10.: Signal fit result for the $10^{\rm th}$ m_{KK} window for MC and data in the range $3.245 < m_{KK} < 3.497$.

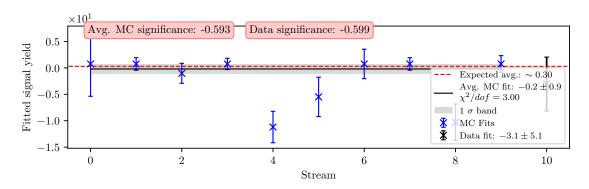


Figure C.11.: Signal fit result for the 11th m_{KK} window for MC and data in the range $3.497 < m_{KK} < 3.748$.

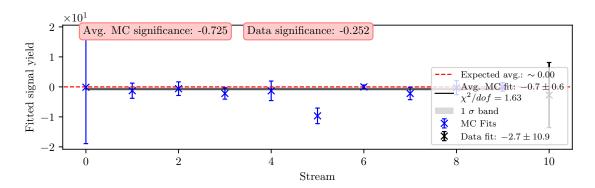


Figure C.12.: Signal fit result for the 12th m_{KK} window for MC and data in the range $3.748 < m_{KK} < 4.000$.

C.2. Signal Fits in q^2

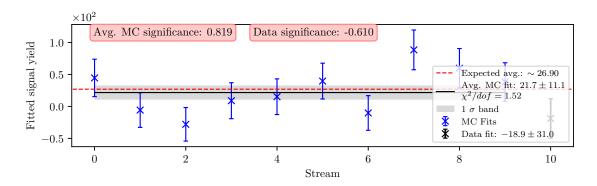


Figure C.13.: Signal fit result for the 1st q^2 window for MC and data in the range $0.000 < q^2 < 1.500$.

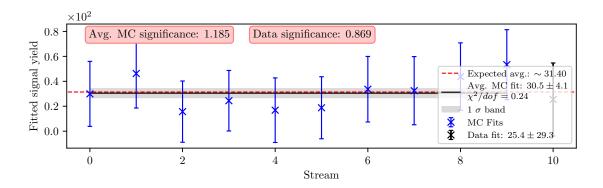


Figure C.14.: Signal fit result for the $2^{\rm nd}$ q^2 window for MC and data in the range $1.500 < q^2 < 3.000$.

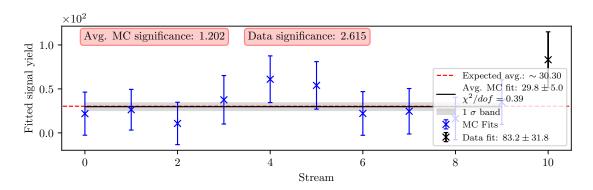


Figure C.15.: Signal fit result for the $3^{\rm rd}$ q^2 window for MC and data in the range $3.000 < q^2 < 4.500$.

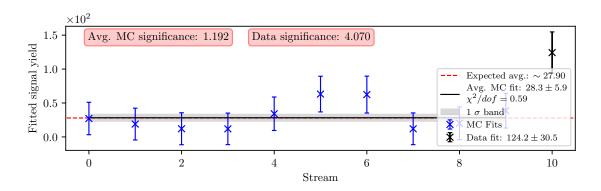


Figure C.16.: Signal fit result for the 4th q^2 window for MC and data in the range $4.500 < q^2 < 6.000$.

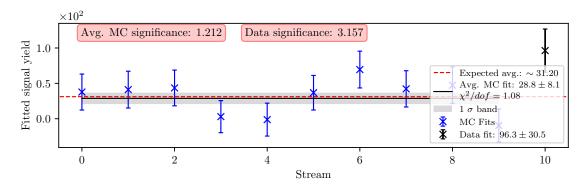


Figure C.17.: Signal fit result for the 5th q^2 window for MC and data in the range $6.000 < q^2 < 7.500$.

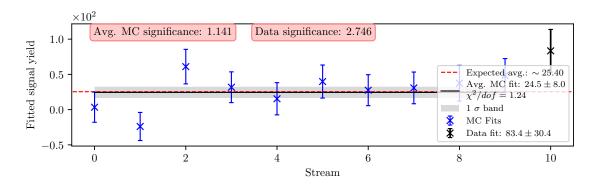


Figure C.18.: Signal fit result for the 6th q^2 window for MC and data in the range $7.500 < q^2 < 9.000$.

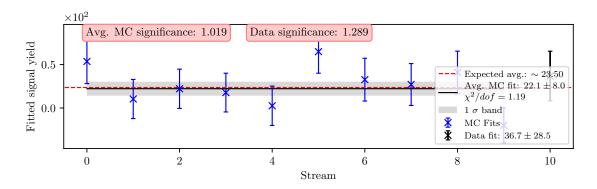


Figure C.19.: Signal fit result for the 7th q^2 window for MC and data in the range $9.000 < q^2 < 10.500$.

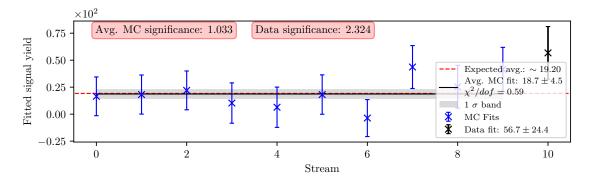


Figure C.20.: Signal fit result for the 8th q^2 window for MC and data in the range $10.500 < q^2 < 12.000$.

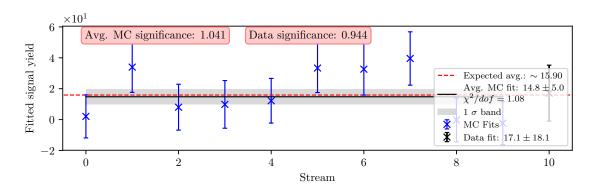


Figure C.21.: Signal fit result for the 9th q^2 window for MC and data in the range $12.000 < q^2 < 13.500$.

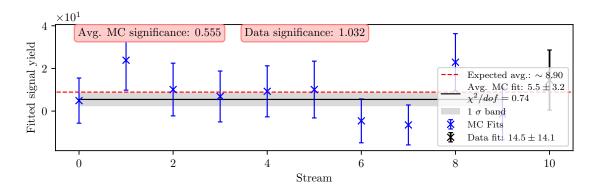


Figure C.22.: Signal fit result for the $10^{\rm th}$ q^2 window for MC and data in the range $13.500 < q^2 < 15.000$.

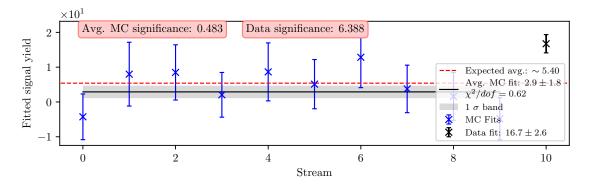


Figure C.23.: Signal fit result for the $11^{\rm th}$ q^2 window for MC and data in the range $15.000 < q^2 < 16.500$.

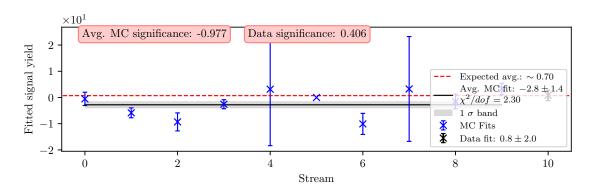


Figure C.24.: Signal fit result for the $12^{\rm th}$ q^2 window for MC and data in the range $16.500 < q^2 < 18.000$.