MACHINE LEARNING AND PATTERN RECOGNITION REPORT

Fingerprint Spoofing Detection

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Contents

Introduction The project task consists of a binary classification problem. The goal is to perform fingerprint spoofing detection, i.e. to identify genuine vs counterfeit fingerprint images. The dataset consists of labeled samples corresponding to the genuine (True, label 1) class and the fake (False, label 0) class. The samples are computed by a feature extractor that summarizes high-level characteristics of a fingerprint image. The data is 6-dimensional.

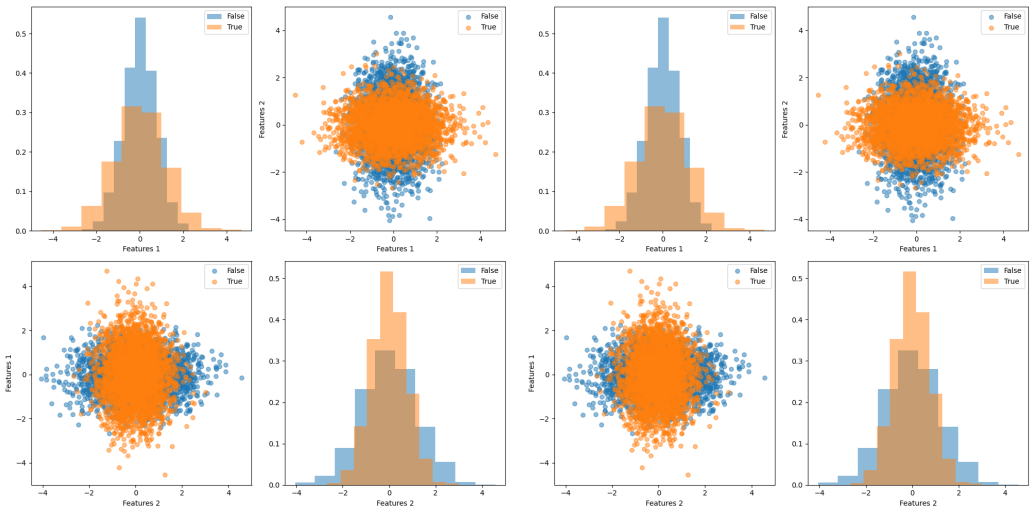
1 Dataset Analysis

In our analysis process, one first begins to represent what are the data related to the various features and how among the various features the data are distributed by making a visual representation in pairs of features.

1. Starting with the analysis of the first two features and creating a histogram and a scatter Figure 1, one can see:

   - Both features overlap
   - Follow a Gaussian distribution
   - Feature 1 has a peak at [-0.213, 0.276] and it is worth 0.541 for the false class, instead for Feature 2 the peak at [-0.402, 0.165] and it is worth 0.516 for the true class.

   Seeing Figure 1 again, it would appear that Figure 1a and Figure 1b appear to be visually the same but in b they are represented centred which as can be seen is quite similar because Feature 1 has $\mu = 0.00170711$ and $\sigma^2 = 1.00134304$, instead Feature 2 has $\mu = 0.00503903$ and $\sigma^2 = 0.9983527$



(a) Without mean (b) With mean

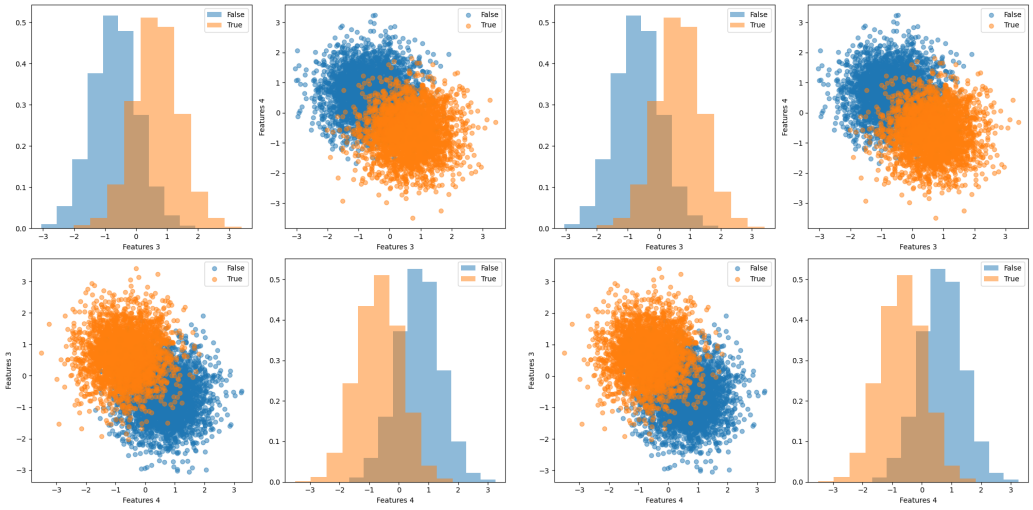
Figure 1: Feature 1 vs Feature 2 - Without centering the data relative to the average (a) and with centering the data relative to the average (b)

2. For features 3 and 4, observed in Figure 2, on the other hand, they have:

   - do not overlap like the previous two
   - Follow a Gaussian distribution but the true and false labels are centred at different points

- Feature 3 has a peak at [-1.063, -0.568] and it is worth 0.517 for the false class, instead for Feature 4 the peak at [0.290, 0.783] and it is worth 0.525 for the false class.

Seeing Figure 2a and Figure 2b, data are already similar because, the mean calculated with reference to the two classes is close to 0, in fact: Feature 3 has $\mu = -0.00560753$ and $\sigma^2 = 1.0024818$, instead Feature 4 has $\mu = 0.00109537$ and $\sigma^2 = 0.99029389$



(a) Without mean (b) With mean

Figure 2: Feature 3 vs Feature 4 - Without centering the data relative to the average (a) and with centering the data relative to the average (b)

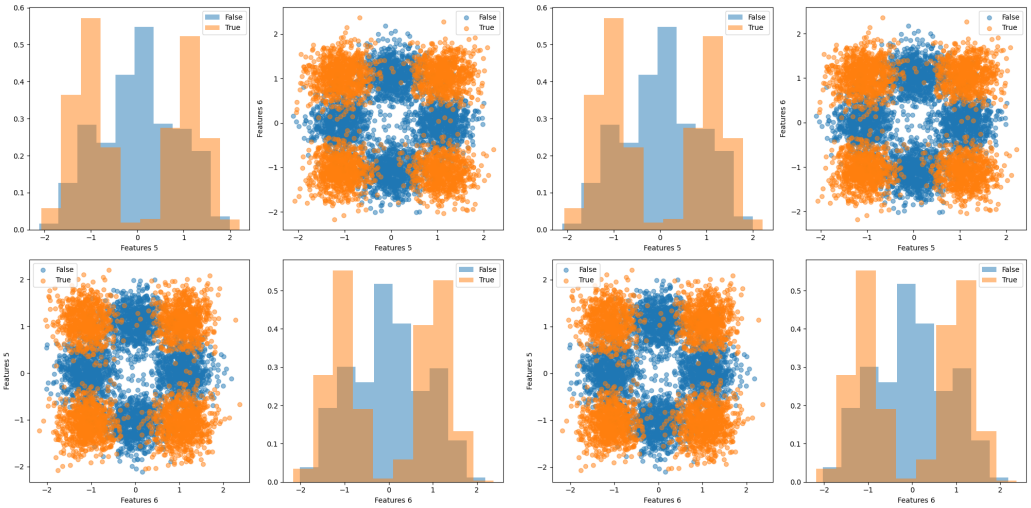
3. For features 5 and 6, observed in Figure 3, one can see:

   - Do not totally overlap
   - For both features, the true labels don't follow a Gaussian distribution as opposed to the false ones, which could be more approximate
   - Feature 5 has a peak at [-1.211, -0.783] and it is worth 0.572 for the true class, instead for Feature 6 has a peak at [-1.273, -0.817] and it is worth 0.553 for the true class.

   Also in this other case, Figure 3a and Figure 3b are similar because again the average is close to 0. Feature 5 has $\mu = -0.00700025$ and Feature 6 has $\mu = 0.00910515$

2 Dimensionality Reduction

Before proceeding with classification, two techniques of dimensionality reduction PCA and LDA can be analysed. The goal is to find a subspace of the feature space that preserves most of the useful information, that is, mapping from the *n–dimensional* feature space to *m–dimensional* space, with $m \ll n$



(a) Without mean (b) With mean

Figure 3: Feature 5 vs Feature 6 - Without centering the data relative to the average (a) and with centering the data relative to the average (b)

2.1 PCA

is an unsupervised technique. Where starting from a dataset $X = \{x_1, \ldots, x_k\}$ and calculated average. It starts with the empirical covariance matrix:

$$C = \frac{1}{K} \sum (x_i - \bar{x})(x_i - \bar{x})^T \tag{1}$$

We compute the eigen-decomposition of $C = U \Sigma U^T$ and project the data in the subspace spanned by the m columns of U corresponding to the m largest eigenvalues.

$$y_i = P^T (x_i - \bar{x}) \tag{2}$$

where P is the matrix of the m columns of U associated to the m highest eigenvalues of C. A cross-validation approach can be used to figure out the optimal value of m to be selected. To evaluate each eigenvalue, one would have to calculate the variance corresponding to the axis. The percentage can be calculated as the rate between the sum of the m eigenvalues and the sum of all of them. In Figure 4 we can see how it changes in the project. A good m, corresponds to that value which allows a percentage greater than 95%, so in our case we would need all 6 features.

2.2 LDA

is a supervised technique. To find a direction that has the best separation between classes, we measure spread between classes in terms of class covariance. The objective is to maximize the $between - class$ variability over $within - class$ variability ratio for the transformed samples:

$$\max_w \frac{w^T S_B w}{w^T S_W w} \tag{3}$$

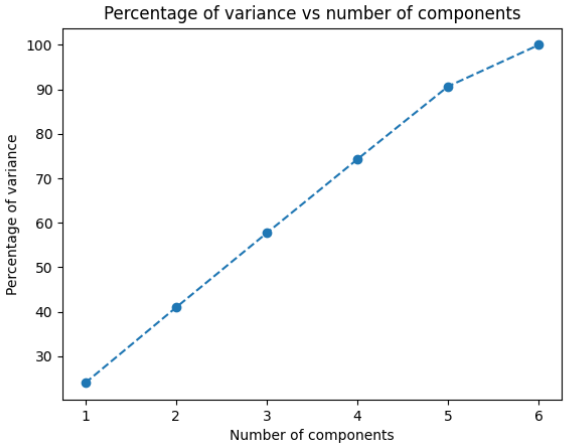


Figure 4: Cross validation for PCA impact evaluation

Method	**Num Samples**	**Error**	**Error Rate** (%)
LDA - First threshold	2000	186	9.30
LDA - Second threshold	2000	186	9.30
PCA (m=5) + LDA - First threshold	2000	186	9.30
PCA (m=5) + LDA - Second threshold	2000	185	9.25
PCA (m=6) + LDA - First threshold	2000	186	9.30
PCA (m=6) + LDA - Second threshold	2000	184	9.20

Table 1: Table showing the results of the LDA and PCA + LDA method for classification.

where:

$$S_B \triangleq \frac{1}{N}\sum_{c=1}^{K} n_c(\mu_c - \mu)(\mu_c - \mu)^T \tag{4}$$

$$S_W \triangleq \frac{1}{N}\sum_{c=1}^{K}\sum_{i=1}^{n_c}(x_{c,i} - \mu_c)(x_{c,i} - \mu_c)^T \tag{5}$$

μ is dataset mean, μ_c is class mean, n_c is the number of samples in class c, and N is the total number of samples.

The directions of LDA can be calculated by solving the generalised eigenvalue problem, as one wants to find the associated eigenvectors $S_w^{-1}S_b$. This method allows us to find at most C-1 discriminant directions, where C is the number of classes, since the objective of LDA is to maximize the separation between classes. In our case there are 2 classes so there is only one direction

2.3 Our project

PCA and LDA are applied to the dataset, in particular, m = 6 is used, and in Figure 5 we can observe what are the outcomes for the indicated directions.

At a later stage, they were used to carry out a classification. The available dataset was divided into two sub-portions one for training and the other for validation. In Table 1 we can see the error of the classification, errors made in the classification when varying m (only for some values) and the threshold were reported

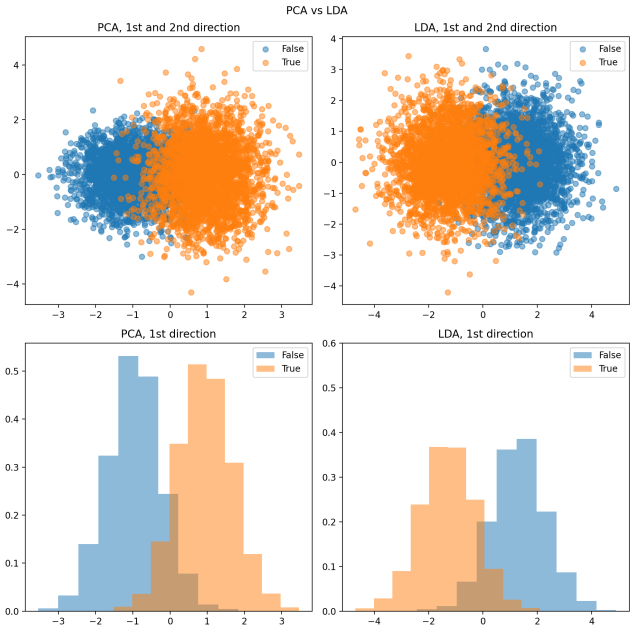


Figure 5: Comparing resulta between PCA and LDA

3 Multivariate Gaussian Density

Multivariate Gaussian Density is an extension of the Gaussian Density to multiple dimensions. It is used to describe the distribution of a vector of random variables in a $multi-dimensional$ space, and it could be defined as:

$$N(x \mid \mu, \Sigma) = \frac{1}{(2\pi)^{\frac{M}{2}} |\Sigma|^{\frac{1}{2}}} \exp^{-\frac{1}{2}(x-\mu)^T \Sigma^{-1}(x-\mu)} \tag{6}$$

where M is the size of the feature vector x, and $|\Sigma|$ is the determinant of Σ. Since the computation of the exponential could cause problems, the logarithm is applied, so from Equation 6 we get Equation 7:

$$\log N(x \mid \mu, \Sigma) = -\frac{M}{2} \log(2\pi) - \frac{1}{2} \log |\Sigma| - \frac{1}{2}(x-\mu)^T \Sigma^{-1}(x-\mu) \tag{7}$$

Thus, applying Gaussian probability density to the 6 features in our dataset, the following results in Figure 6, are obteinded.

It can observe that **features 1 - 2 - 3 - 4** fit the Gaussian distribution, the histogram of the data is well approximated by the Gaussian curve. For **features 5 - 6** this does not happen and therefore could not be a good model.

4 Model evaluation for classification

To assess whether one model classifies correctly or it is better than another, it is important to introduce what may be a unit of measurement. In particular, the evaluation of a model is done using the **DCF (Detection Cost Function)** or also called **empirical Bayes Risk**. But before delving into this method of valuation, we must consider what is called the working point of a binary classification application. This point is characterised by a triplet of values:

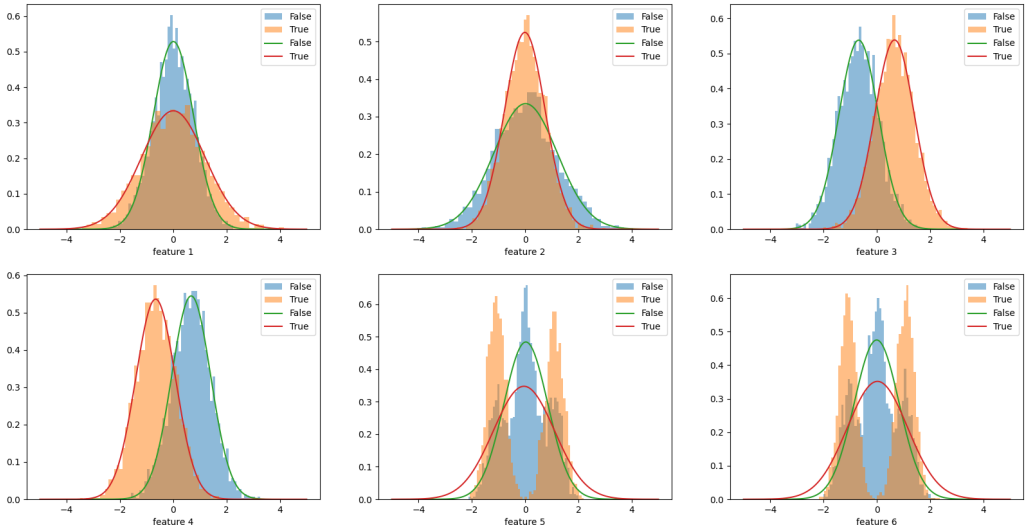


Figure 6: Gaussian Density

$$(\pi_T, C_{fn}, C_{fp}) \tag{8}$$

where π_T is the prior probability of the target class, C_{fn} is the cost of a false negative and C_{fp} is the cost of a false positive.

But from π_T, one can have attention on a particular triplet that is defined by:

$$(\tilde{\pi}, C_{fn}, C_{fp}) = (\tilde{\pi}, 1, 1) \tag{9}$$

where $\tilde{\pi}$ in Equation 9 is called **effective prior** probability of the target class, defined as:

$$\tilde{\pi} = \frac{\pi_T C_{fn}}{\pi_T C_{fn} + (1 - \pi_T) C_{fp}} \tag{10}$$

So we can define DCF_u as:

$$B_{emp} = DCF_u = \sum_{c=1}^{K} \frac{\pi_c}{N_c} \sum_{i|c_i=c} C(a(x_i, R) \mid c) \tag{11}$$

From Equation 11, we can obtain:

$$DCF_u(\pi_T, C_{fn}, C_{fp}) = \pi_T C_{fn} P_{fn} + (1 - \pi_T) C_{fp} P_{fp} \tag{12}$$

where:

$$P_{fn} = \frac{FN}{FN + TP}\ , \quad P_{fp} = \frac{FP}{FP + TN} \tag{13}$$

From the Equation 11, it is possible to find **minDCF** and **actDCF**. When calculating minDCF, the threshold used is the one that minimize DCF, and there are various methods for finding it. Whereas for actDCF, the

threshold used is:

$$t' = -\log \frac{\tilde{\pi}}{1-\tilde{\pi}} \tag{14}$$

5 Classification Models Analysis

To perform the classification, the dataset must first be divided into two sub-portions, the training and validation sub-portions.

5.1 Gaussian models

Since, it is dealing with a binary classification task, it will assign a probabilistic score to each sample in terms of the class-posterior log-ratio:

$$\log r(x_t) = \log \frac{P(C = h_1 \mid x_t)}{P(C = h_0 \mid x_t)} \tag{15}$$

Analysing Equation 15 in more detail, it becomes:

$$\log r(x_t) = \log \frac{f_{X|C}(x_t \mid h_1)}{f_{X|C}(x_t \mid h_0)} + \log \frac{P(C = h_1)}{P(C = h_0)} \tag{16}$$

The first addend of the equation is called the *llr* or *log-likelihood ratio* and an optimal decision is given by Equation 17.

$$\log r(x_t) \gtrless 0 \tag{17}$$

Considering $P(C = h_1) = \pi$ and $P(C = h_0) = 1 - \pi$, from Equation 16 and Equation 17, it is possible to write that the class assignment is based on Equation 16 and Equation 17, to obtain Equation 18.

$$llr(x_t) = \log \frac{f_{X|C}(x_t \mid h_1)}{f_{X|C}(x_t \mid h_0)} \gtrless -\log \frac{\pi}{1-\pi} \tag{18}$$

The optimal class decision is based on a comparison between the and a threshold , if the llr is greater than th the sample is assigned to class h_1, otherwise to class h_0. It is necessary to find the parameters θ, μ_c, Σ_c; this can be done by maximising the log-likelihood. Parameter estimation is part of the training phase and this therefore performed on the training part of the dataset, then an estimation of the error rate can be performed on the validation part.

5.1.1 Multivariate Gaussian Classifier

The first classifier is MVG and it is given by the empirical mean and covariance matrix for each class,

$$\mu_c^* = \frac{1}{N_c} \sum_{i|c_i=c} x_i \ , \quad \Sigma_c^* = \frac{1}{N_c} \sum_{i|c_i=c} (x_i - \mu_c^*)(x_i - \mu_c^*)^T \tag{19}$$

5.1.2 Naive Bayes Gaussian Classifier

This model makes an important assumption that simplifies the number of parameters to be estimated, it assumes that the features are independent given their class. This causes the covariance matrix to be a diagonal matrix,

consequently, matching MVG with a diagonal covariance matrix. However, the assumption of independence may be too restrictive and lead to inferior performance if the features are indeed correlated.

$$\mu^*_{c,[j]} = \frac{1}{N_c} \sum_{i|c_i=c} x_{i,[j]} \ , \quad \sigma^2_{c,[j]} = \frac{1}{N_c} \sum_{i|c_i=c} (x_{i,[j]} - \mu^*_{c,[j]})^2 \tag{20}$$

5.1.3 Tied Covariance Gaussian Classifier

The assumption of the latter model consists of its own average for each class, but an equal covariance matrix for all classes.

$$\mu^*_c = \frac{1}{N_c} \sum_{i|c_i=c} x_i \ , \quad \Sigma^* = \frac{1}{N} \sum_c \sum_{i|c_i=c} (x_i - \mu_c)(x_i - \mu_c)^T \tag{21}$$

The characteristic of this model is that it is strongly correlated to LDA.

5.1.4 Gaussian Models Comparison

A threshold of 0 was used to perform our results, which means that $P(C = 1) = P(C = 0) = 1/2$. This model was applied and the outcomes can be seen in the Table 2.

Features	**Model**	**Error Rate** (%)
	no PCA	
1 to 6	MVG	7.00
1 to 6	Naive Bayes	7.20
1 to 6	Tied Covariance	9.30
1 to 4	MVG	7.95
1 to 4	Naive Bayes	7.65
1 to 4	Tied Covariance	9.50
1 - 2	MVG	36.50
1 - 2	Naive Bayes	36.30
1 - 2	Tied Covariance	49.45
3 - 4	MVG	9.45
3 - 4	Naive Bayes	9.45
3 - 4	Tied Covariance	9.40
	PCA m = 5	
1 to 6	MVG	7.10
1 to 6	Naive Bayes	8.75
1 to 6	Tied Covariance	9.30
	PCA m = 6	
1 to 6	MVG	7.00
1 to 6	Naive Bayes	8.90
1 to 6	Tied Covariance	9.30

Table 2: Table showing the results of the Error Rate for different Models and Features.

Comparing the results with the Table 1, we can see that for some configurations there were improvements in terms of error rate. This means that Gaussian models are better able to classify the data. If we go into the details of how the error rate changes as a function of the observed features, we can see that:

- **1 to 6**: in the case we consider all 6 features, the error rate is quite low and its range goes from 7.00% to 9.30%. This means that they all provide useful information.
- **1 to 4**: if we consider features from 1 to 4, we can see that the error rate increases slightly. This allow us to say that features 5 and 6 have useful but not fundamental information to change the outcome.
- **1 - 2**: features 1 and 2 have a rather high error rate, meaning that they don't contain relevant information.
- **3 - 4**: on the other hand, the latter two features considered have a rather low error rate, a value close to the case where all features are considered. This means that the information contained in these two features is relevant for making the classification.

Starting with the previous performance, it may be interesting to analyse how performance changes as three main parameters vary:

- $\tilde{\pi}$: represents prior probability of the positive class
- C_{fn}: misclassification cost of a sample predicted as negative but it is positive
- C_{fp}: misclassification cost of a sample predicted as positive but it is negative

	MVG	**Naive Bayes**	**Tied Covariance**
Application $(\tilde{\pi}, C_{fn}, C_{fp}) = (0.5, 1, 1)$			
actDCF	0.1399	0.1439	0.1860
minDCF	0.1302	0.1311	0.1812
Application $(\tilde{\pi}, C_{fn}, C_{fp}) = (0.9, 1, 1)$			
actDCF	0.4001	0.3893	0.4626
minDCF	0.3423	0.3509	0.4421
Application $(\tilde{\pi}, C_{fn}, C_{fp}) = (0.1, 1, 1)$			
actDCF	0.3051	0.3022	0.4061
minDCF	0.2629	0.2569	0.3628
Application $(\tilde{\pi}, C_{fn}, C_{fp}) = (0.5, 1, 9)$			
actDCF	0.3051	0.3022	0.4061
minDCF	0.2629	0.2569	0.3628
Application $(\tilde{\pi}, C_{fn}, C_{fp}) = (0.5, 9, 1)$			
actDCF	0.4001	0.3893	0.4626
minDCF	0.3423	0.3509	0.4421

Table 3: Table showing minDCF and actDCF for different models and applications.

Analysing the results of the Table 3, it is possible to observe:

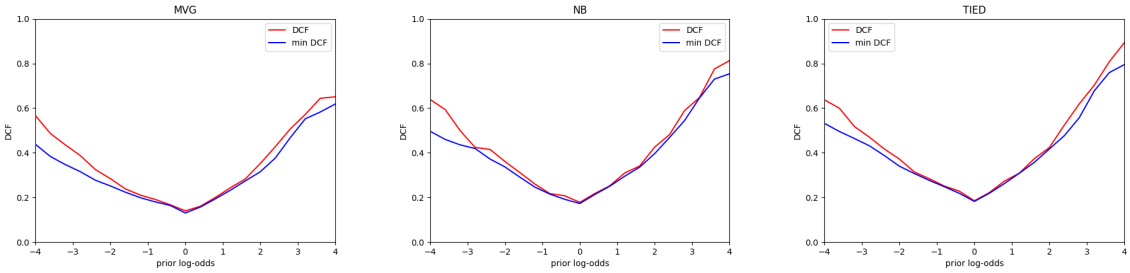
- Observing how the $\tilde{\pi}$ varies, it can be seen that the best outcome is obtained when it takes the value 0.5. On the other hand, when it takes value 0.1 and 0.9, the outcome gets worse because it penalises false negatives and false positives respectively
- Observing how the values of C_{fn} and C_{fp} change when they assume value 9. The outcomes worsen, in particular there is a greater impact on the model when the cost of false negatives increases.

Starting from the result obtained in Table 3, it is possible to consider the application of PCA as pre-processing technique focusing on the cases of $\tilde{\pi}$ equal to 0.1, 0.5 and 0.9 and $C_{fn} = C_{fp} = 1$, obtaining the outcomes shown in Table 4

	MVG	**Naive Bayes**	**Tied Covariance**
Application $(\tilde{\pi}, C_{fn}, C_{fp}) = (0.5, 1, 1)$			
no PCA			
actDCF	0.1399	0.1439	0.1860
minDCF	0.1302	0.1311	0.1812
PCA			
$m = 5$			
actDCF	0.1419	0.1749	0.1860
minDCF	0.1331	0.1737	0.1812
$m = 6$			
actDCF	0.1399	0.1780	0.1860
minDCF	0.1302	0.1727	0.1812
Application $(\tilde{\pi}, C_{fn}, C_{fp}) = (0.9, 1, 1)$			
no PCA			
actDCF	0.4001	0.3893	0.4626
minDCF	0.3423	0.3509	0.4421
PCA			
$m = 5$			
actDCF	0.3980	0.4660	0.4626
minDCF	0.3512	0.4340	0.4451
$m = 6$			
actDCF	0.4001	0.4512	0.4626
minDCF	0.3423	0.4359	0.4421
Application $(\tilde{\pi}, C_{fn}, C_{fp}) = (0.1, 1, 1)$			
no PCA			
actDCF	0.3051	0.3022	0.4061
minDCF	0.2629	0.2569	0.3628
PCA			
$m = 5$			
actDCF	0.3042	0.3930	0.4051
minDCF	0.2738	0.3545	0.3648
$m = 6$			
actDCF	0.3051	0.3920	0.4061
minDCF	0.2629	0.3535	0.3628

Table 4: Show minDCF and actDCF for different models and applications before and after applying PCA.

From the results obtained in the Table 4, it can be seen that the application of PCA was not very helpful because in no case did the outcomes improve, instead they remained the same or even worsened. Analysing overall, it can be said that the model that tends to perform worse is the Tied Covariance, whereas MVG and



(a) MVG (b) Naive Bayes (c) Tied Covariance

Figure 7: Error Bayes plots

Naive Bayes are rather similar. In particular for MVG and Naive Bayes, the best result is obtained for the $0.5, 1, 1$ configuration whether applying PCA or not. In addition, it can be said that there is a good calibration for this configuration because applying or not applying PCA, minDCF and actDCF doesn't change what is not the case for the other configurations.

In Figure 7 the Bayes error was calculated for a prior log odds in the range $(-4, +4)$, for the three models with a configuration having a $\tilde{\pi} = 0.1$ and applying the PCA as pre-processing.

5.2 Logistic Regression Classifier

Logistic Regression is a discriminative classification model, directly evaluating the posterior probability $C \mid X$. In particular by determining that hyperplane which maximises the posterior probability. Starting from the results obtained from the Tied Gaussian that provides log-likelihood ratios that are linear functions of our data, where log-posterior probability ratio is:

$$\log \frac{P(C = h_1 \mid X)}{P(C = h_0 \mid X)} = \log \frac{f_{X|C}(x \mid h_1)}{f_{X|C}(x \mid h_0)} + \log \frac{\pi}{1 - \pi} = \omega^T x + b \tag{22}$$

where prior information has been absorbed in the bias term b of the Equation 22. So from this point we can define the score function as:

$$s(x) = \omega^T x + b = 0 \tag{23}$$

where it is positive for samples of class h_1 and negative for samples of class h_0. Given ω and b we can compute the posterior class probability as:

$$P(C = h_1 \mid x, \omega, b) = \sigma(\omega^T x + b) = \sigma(s(x)) \tag{24}$$

where σ is sigmoid function defined as:

$$\sigma(x) = \frac{1}{1 + e^{-x}} \tag{25}$$

This approach assumes that the decision rules will be hyperplanes orthogonal to vector w.

5.2.1 Binary Logistic Regression

Binary Logistic Regression Not Prior-Weighted

The objective is to minimise the loss function $J(\omega, b)$, but to this is introduced what is a penalty term, so the new function becomes:

$$J(\omega, b) = \frac{\lambda}{2} \|w\|^2 + \frac{1}{n} \sum_{i=1}^{n} \log(1 + e^{-z_i(\omega^T x_i + b)}), \quad z_i = \begin{cases} 1 & \text{if } c_i = 1 \\ -1 & \text{if } c_i = 0 \end{cases} \tag{26}$$

where λ of Equation 26 is the regularization term, this term has been introduced to make problem solvable in case of linearly separable classes.

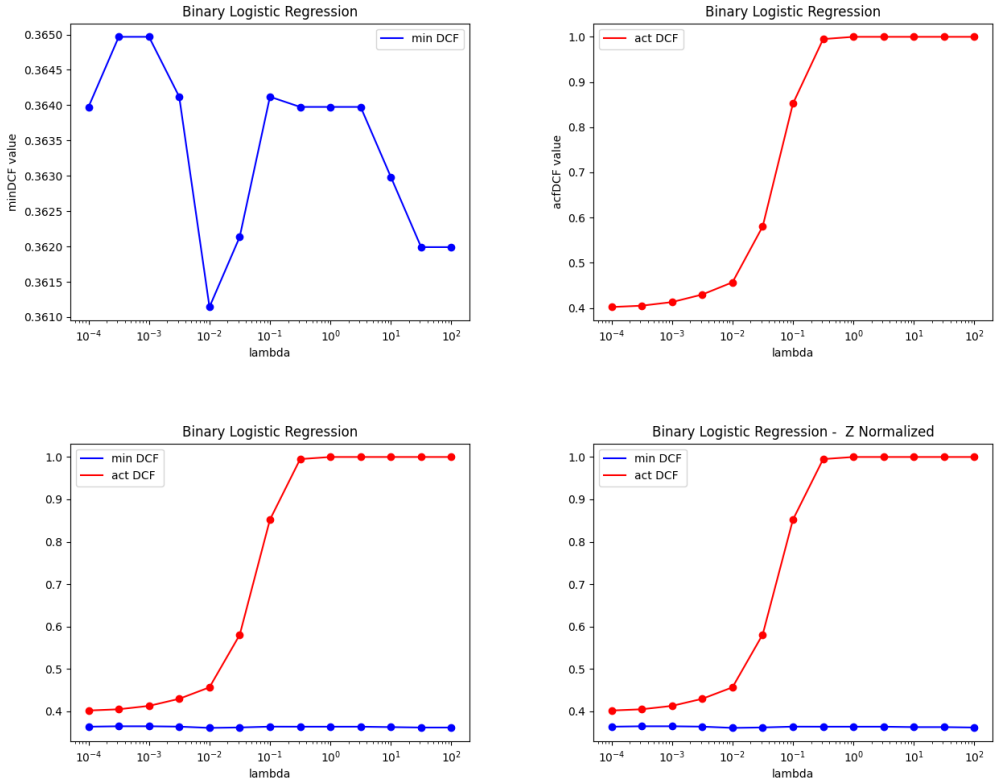


Figure 8: Binary Logistic Regression not Prior-Weighted

In this model $\pi_T = 0.1$ is used. In Table 5, it can be seen how the values of minDCF and actDCF vary when λ changes, Z-normalization is applied or not and if the whole training set or a portion was used. It can be deduced from the values obtained that the application of z-normalisation brings no advantage. On the other hand, by using only 50 samples, it can see that using a limited number of samples can significantly influence the model and could lead to misleading results that are not representative of the entire sample. Consequently, as many samples as possible should be used for training to obtain a more accurate model. Figure 8 and Figure 9 give a graphic representation of how minDCF and actDCF vary with λ

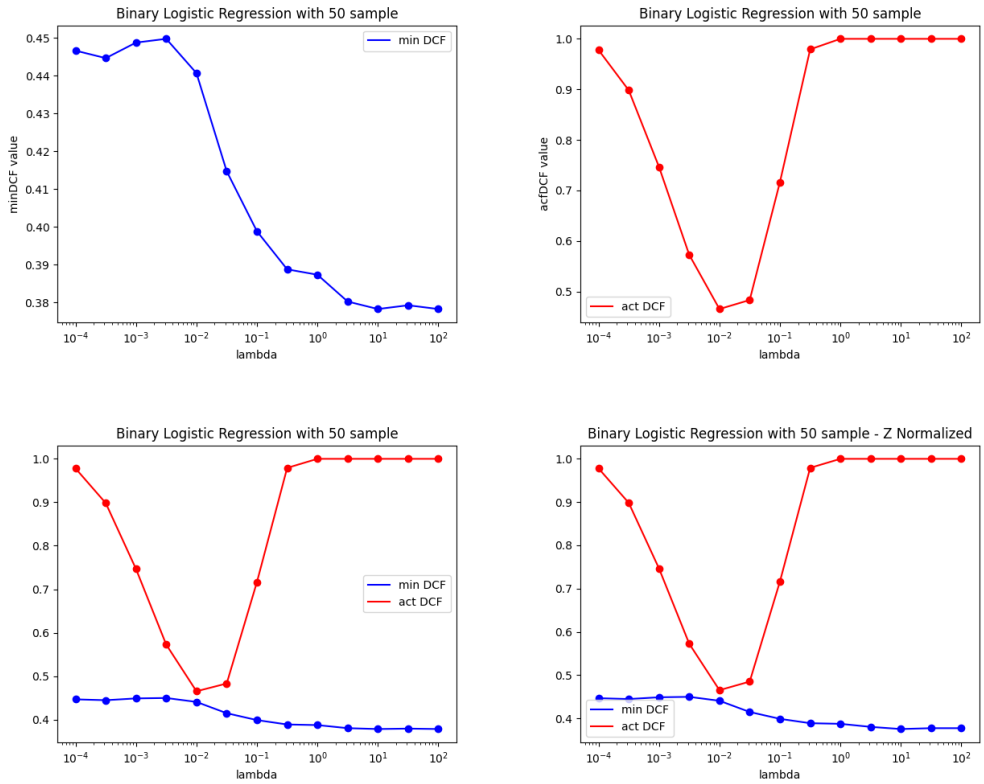


Figure 9: Binary Logistic Regression not Prior-Weighted with 50 Samples

Binary Logistic Regression Not Prior-Weighted				
λ	**minDCF**		**actDCF**	
	no z-norm	**z-norm**	**no z-norm**	**z-norm**
10^{-4}	0.3640	0.3640	0.4021	0.4021
10^{-3}	0.3650	0.3650	0.4130	0.4130
10^{-2}	0.3611	0.3611	0.4568	0.4568
10^{-1}	0.3641	0.3641	0.8522	0.8522
Binary Logistic Regression Not Prior-Weighted (50 Samples)				
λ	**minDCF**		**actDCF**	
	no z-norm	**z-norm**	**no z-norm**	**z-norm**
10^{-4}	0.4466	0.4466	0.9780	0.9780
10^{-3}	0.4487	0.4487	0.7466	0.7466
10^{-2}	0.4407	0.4407	0.4652	0.4652
10^{-1}	0.3988	0.3988	0.7164	0.7164

Table 5: Show minDCF and actDCF for Binary Logistic Regression Not Prior-Weighted model

Binary Logistic Regression Prior-Weighted

Another possible Logistic Regression approach is that Prior-Weighted; it allows to simulate different priors for class 1. Therefore, the objective function becomes:

$$J(\omega, b) = \frac{\lambda}{2} \|w\|^2 + \sum_{i=1}^{n} \xi_i \log(1 + e^{-z_i(\omega^T x_i + b)}), \quad \xi_i = \begin{cases} \frac{\pi_t}{n_T} & \text{if } z_i = +1 (c_i = 1) \\ \frac{1-\pi_T}{n_F} & \text{if } z_i = -1 (c_i = 0) \end{cases} \tag{27}$$

Now it is possible analyze the results obtained from the Prior-Weighted model with $\pi_T = 0.1$.

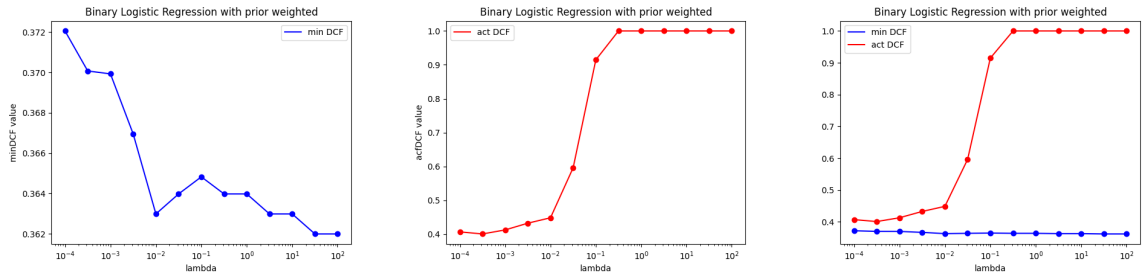


Figure 10: Binary Logistic Regression Prior-Weighted

Binary Logistic Regression Prior-Weighted		
$\pi_T = 0.1$		
λ	**minDCF**	**actDCF**
10^{-4}	0.3721	0.4071
10^{-3}	0.3699	0.4129
10^{-2}	0.3630	0.4487
10^{-1}	0.3648	0.9147

Table 6: Show minDCF and actDCF for Binary Logistic Regression Prior-Weighted

The role of the prior is to weight the samples during model training. In particular, samples in the higher priority class receive a higher weight than those in the lower priority class. This can be useful if the dataset is unbalanced, the choice of the prior must be made at the beginning and this choice can affect the model a lot, in fact we may even have a worsening of the model.
Comparing the results obtained in Table 5 and Table 6, there isn't noticeable change in the outcomes on minDCF and actDCF this means that our dataset is not unbalanced. The interesting value to observe is the value of actDCF when $\lambda = 10^-1$.

Binary Logistic Regression with Pre-processing (PCA)

In this case, PCA can be applied as pre-processing, and by looking at the values in Table 7, it can be seen that it does not result much improvement of the system.

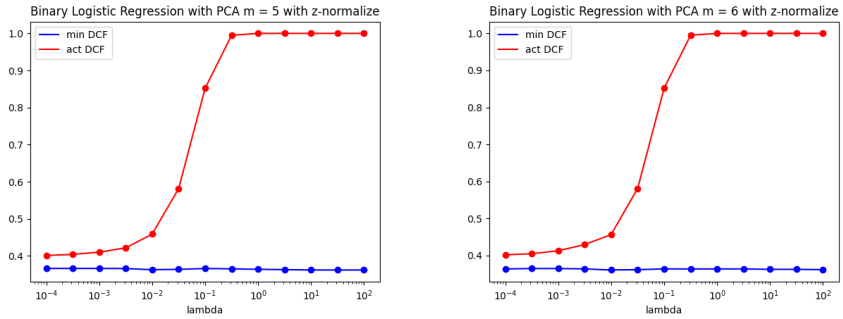


Figure 11: Binary Logistic Regression applying PCA and z-normalization

Binary Logistic Regression with PCA				
λ	**minDCF**		**actDCF**	
	no z-norm	**z-norm**	**no z-norm**	**z-norm**
		$m = 5$		
10^{-4}	0.3661	0.3661	0.4011	0.4011
10^{-3}	0.3661	0.3661	0.4100	0.4100
10^{-2}	0.3618	0.3628	0.4578	0.4588
10^{-1}	0.3660	0.3660	0.8502	0.8522
		$m = 6$		
10^{-4}	0.3640	0.3640	0.4021	0.4021
10^{-3}	0.3650	0.3650	0.4130	0.4130
10^{-2}	0.3611	0.3611	0.4568	0.4568
10^{-1}	0.3641	0.3641	0.8522	0.8522

Table 7: Show minDCF and actDCF for Binary Logistic Regression applying PCA and with adn withoud z-normalization

5.2.2 Quadratic Logistic Regression

In this step we can analyze training on a Quadratic Logistic Regression model by performing features expansion, so it's possible write log-likelihood ratio as:

$$\log \frac{P(C = h_1|x)}{P(C = h_0|x)} = x^T Ax + b^T x + c = s(\mathbf{x}, \mathbf{A}, \mathbf{b}, \mathbf{c}) \tag{28}$$

The Equation 28 is quadratic in x but it's linear in A and b. It can be rewritten to obtain a decision function that is linear for the expanded features space but quadratic in original features space.
So we can write features expansion as:

$$\Phi(x) = \begin{bmatrix} vec(xx^T) \\ x \end{bmatrix}, \quad w = \begin{bmatrix} vec(A) \\ b \end{bmatrix} \tag{29}$$

where $vec(X)$ in Equation 29 is the operator that stacks the columns of X into a single column vector. In this way we can write the posterior log-likelihood as:

$$s(x, w, c) = s^T \phi(x) + c \tag{30}$$

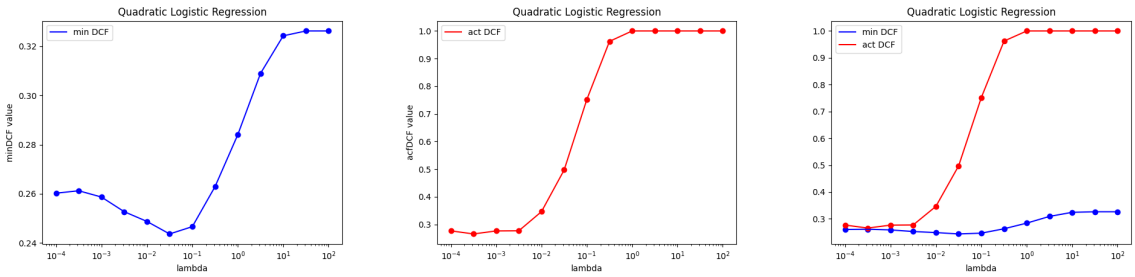


Figure 12: Quadratic Logistic Regression minDCF and actDCF

Quadratic Logistic Regression		
λ	**minDCF**	**actDCF**
10^{-4}	0.2602	0.2768
10^{-3}	0.2587	0.2765
10^{-2}	0.2487	0.3464
10^{-1}	0.2466	0.7520
Best Result		
$3.162 * 10^{-2}$	0.2436	0.4972

Table 8: Show minDCF and actDCF for Quadratic Logistic Regression and best result in logistic regression

Looking the value in Table 8, we can see that this method gives better results than the methods seen before. This is possible because the quadratic method allows us to extract characteristics that are not possible by a linear method.

Summarize

From the results obtained, one can observe:

- The application of z-normalization does not bring any improvements, which means that being a linear transformation didn' t change the ability of logistic regression to separate classes, if labels and characteristics are linearly separable.
- Looking at the graphs depicted in Figure 8, Figure 9, Figure 10, Figure 11 and Figure 12, it can be seen that λ significantly affects the performance of the model in term of minDCF and actDCF. In particular, it can be observed that for values above 10^{-1} there is significant degradation, which is why the tables have shown values up to this λ value. This is because, remember that a larger value of λ can lead to overgeneralise resulting, so underfitting. Whereas too small λ can lead to low generalisation and thus to overfitting.
- It can be seen that there is a difference between minDCF and actDCF, which means that the models don't have a good calibration. In our case, Quadratic Logistic Regression seems to give the best results, so we can deduce that the model must be able to capture non-linear relationships between the variables and thus find non-linear separation rules in the feature space.

5.3 Support Vector Machine Classifier

5.3.1 Linear Support Vector Machines

Support Vector Machines are linear classifiers that look for maximum margin separation hyperplanes. The primal formulation of the soft-margin SVM problem consists in minimizing the function:

$$\mathbf{J}(w,b) = \frac{1}{2}||w||^2 + C\sum_{i=1}^{N} max(0, 1 - z_i(w^T x_i + b)) \tag{31}$$

where in Equation 31 N is the number of training samples, C is the regularization parameter, and z_i is the margin of the i-th sample.
The dual formulation of the problem is:

$$\mathbf{J}(\alpha) = -\frac{1}{2}\alpha^T\ \mathbf{H}\ \alpha + \alpha^T\mathbf{1} \qquad 1 \leq \alpha_i \leq C,\ \ \forall i \in \{1, ..., N\},\ \ \sum_{i=1}^{n} \alpha_i z_i = 0 \tag{32}$$

where H in Equation 32 is $H_{ij} = z_i z_j x_i^T x_j$ and the dual solution is the maximizer of $J^D(\alpha)$.
Primal and dual solutions are related through:

$$w^* = \sum_{i=1}^{N} \alpha_i^* z_i x_i \tag{33}$$

In addition it's possible to rewrite dual problem as minimization of:

$$\hat{\mathbf{L}}(\alpha) = -\mathbf{J}(\alpha) = \frac{1}{2}\alpha^T\ \mathbf{H}\ \alpha - \alpha^T\mathbf{1} \tag{34}$$

and it can be minimized by L-BFGS-B algorithm. After that we have calculated the optimal α we can compute w^*.

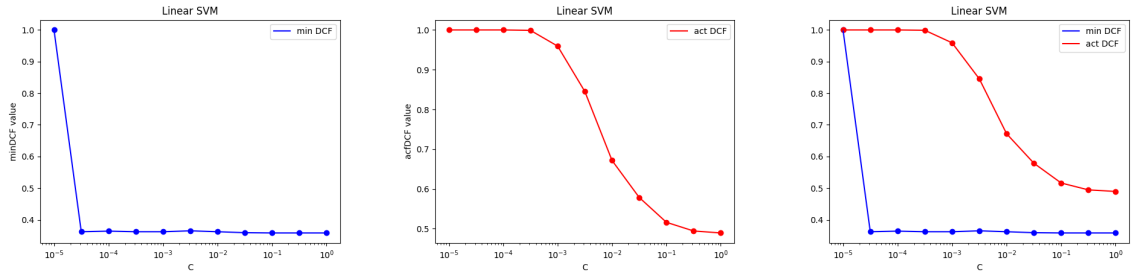


Figure 13: Shows minDCF and actDCF for Linear SVM

Linear SVM $K = 1.0$		
C	**minDCF**	**actDCF**
10^{-5}	1.0000	1.0000
10^{-4}	0.3640	1.0000
10^{-3}	0.3620	0.9593
10^{-2}	0.3620	0.6718
10^{-1}	0.3582	0.5162
10^{0}	0.3582	0.4894

Table 9: Show minDCF and actDCF for Linear SVM

5.3.2 Kernel Support Vector Machines

It's possible in Support Vector Machines to use kernels to allow nonlinear classification. In this case there is no explicit expansion of the feature space; we only can calculate the scalar product between the expanded features:$k(x_1, x_2) = \phi(x_1)_T\phi(x_2)$ where k is the kernel function. To do this we need to go and replace H as we saw in the previous section with $\hat{H} = z_i z_j k(x_1, x_2)$. During our project we see two different types of kernels:

- **Polynomial kernel of degree d**: $k(x_1, x_2) = (x_1^T x_2 + c)^d$
- **Radial Basis Function kernel(RBF)**: $k(x_1, x_2) = e^{-\gamma||x1-x_2||^2}$

We can now apply the polynomial kernel to the SVM with $d = 2, \quad c = 1, \quad \xi = 0$ and see how minDCF and actDCF vary as C changes in Figure 14 and Table 10.

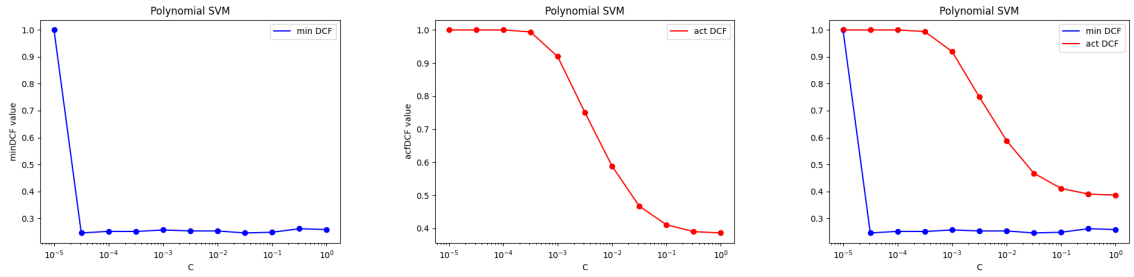


Figure 14: Shows minDCF and actDCF with polynomial kernel

Polynomial Kernel $d = 2, c = 1, \xi = 0$		
C	**minDCF**	**actDCF**
10^{-4}	0.2513	1.0000
10^{-3}	0.2565	0.9196
10^{-2}	0.2528	0.5884
10^{-1}	0.2480	0.4109
10^{0}	0.2582	0.3861

Table 10: Show minDCF and actDCF for SVM with polynomial kernel

In this case apply a RBF kernel to the SVM with $\xi = 1$, the first thing we consider is which value of γ can give us a better outcome in terms of minDCF and actDCF. Looking at Figure 15, we sse that the best solution is given to us by $\gamma = 0.1$.

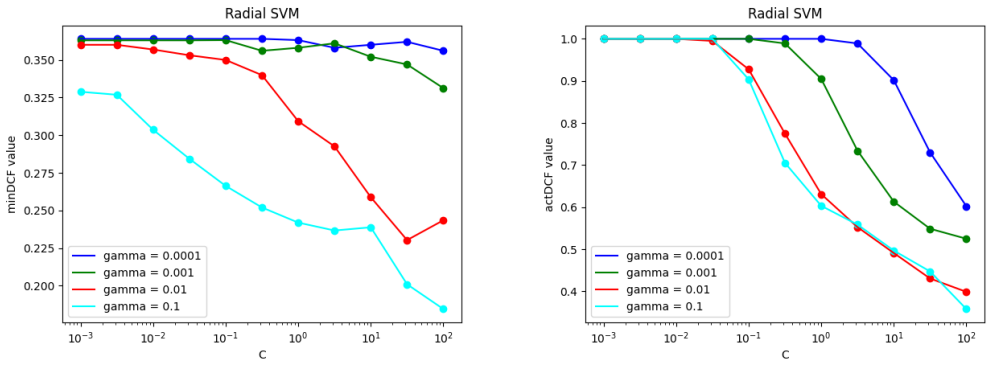


Figure 15: Shows minDCF and actDCF with polynomial kernel

Now that this consideration has been made, we can specifically represent the values of minDCF and actDCF for values of $\gamma = 0.1$. We can observe this result in Figure 16 and Table 11.

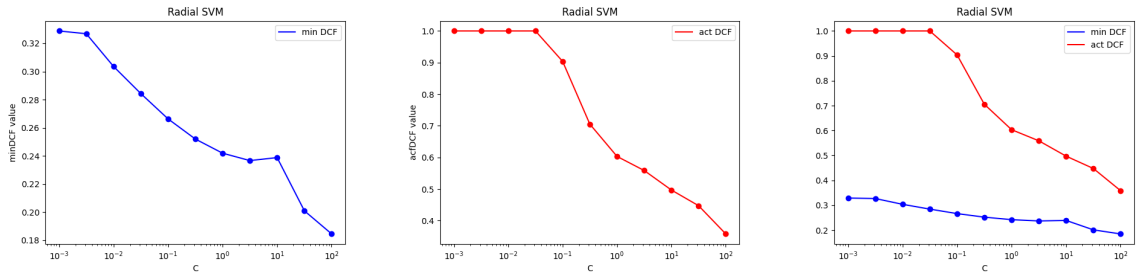


Figure 16: Shows minDCF and actDCF with best RBF kernel

Radial Kernel $\xi = 1, \gamma = 0.1$		
C	**minDCF**	**actDCF**
10^{-3}	0.3288	1.0000
10^{-2}	0.3036	1.0000
10^{-1}	0.2663	0.9038
10^{0}	0.2419	0.6033
10^{1}	0.2388	0.4970
Best result		
10^{2}	0.1845	0.3581

Table 11: Show minDCF and actDCF for SVM with best RBF kernel

Summarize

Looking at the outcomes, one can see:

- when parameter C increases, the outcomes become better and also better calibrated because the difference between minDCF and actDCF decreases.

- we can see that outcomes improve when switching from a linear to a polynomial kernel, which confirms what we have identified with QLR, namely modelling quadratic relationships between features to improve outcomes.
- the case with RBF kernels also confirms that more benefits can be gained from non-linear mapping
- As already mentioned, cases with higher C values improve outcomes and are better calibrated than cases with lower C values. But in general, a calibration operation could improve classification even for high C values.

5.4 Gaussian Mixture Models Classifier

The last model we are going to consider is a generative model, the Gaussian Mixture Model (GMM). This model is based on the assumption that the data is generated by a mixture of K Gaussian distributions. The GMM density consists of a weighted sum of K Gaussians:

$$\mathbf{X} \sim GMM(\mathbf{M}, \mathcal{S}, w) \implies f_x(x) = \sum_{c=1}^{K} \mathcal{N}(x|\mu_g, \Sigma_g) w_g \tag{35}$$

where $M = [\mu_1 ... \mu_k]$, $\mathcal{S} = [\Sigma_1 ... \Sigma_k]$ and $w = [w_1 ... w_k]$ of Equation 35 are the parameters of the model. Gaussian components can be viewed as clusters to which the samples belong (hard or soft), and the cluster label is an unobserved latent random variable. We can also define in this case the responsability term, defined as Equation 36, that represents the posterior probability that a sample belongs to a certain cluster:

$$\gamma(z_{n,i}) = P(G_i = g|X_i = x) = \frac{f_{x_i,G_i}(x_i, g)}{f_{x_i}(x_i)} = \frac{\mathcal{N}(x_i|\mu_g, \Sigma_g) w_g}{\sum_{g'} \mathcal{N}(x_i|\mu_{g'}, \Sigma_{g'}) w_{g'}} \tag{36}$$

Then assign the sample to the cluster label for which the responsability is highest and re-estimate the model parameters based on the cluster assignment. We can apply the Expectation-Maximization algorithm to estimate the model parameters. The EM algorithm is an iterative algorithm that consists of two steps:

- Expectation stage: estimation of the responsability (given the model parameters (M_t, S_t, w_t))
- Maximization step: estimation of new model parameters using the above statistics, estimation continues from an initial value of the model parameters until a certain criterion is met.

The EM algorithm then requires an initial estimate for the GMM parameters, so we use the LBG algorithm to incrementally construct a GMM with 2G components from a GMM with G components. The starting point will be $(1, \mu, \sigma)$, so we use the empirical mean and covariance matrix of the data set. Then it builds a 2-component model starting from one and from each of the new components 2 more components are generated and so on. GMM can have differents versions as:

- **The full covariance model:** in this case each component has a full covariance matrix, which means that all possible covariances between the variables are considered.
- **The diagonal covariance model:** in this setup, the covariance matrix of each component is assumed to be diagonal, which means the variables are considered independent.

As an initial point in our project, we considered having a maximum component number of 32 and using the two methods mentioned above. So at first we evaluated the model where the number of components for class 0 and class 1 were equal, obtaining the results shown in Table 12. Instead, a graphical view of these results can be seen for the Full Covariance method in Figure 17 and in Figure 18 for the Diagonal Covariance method.

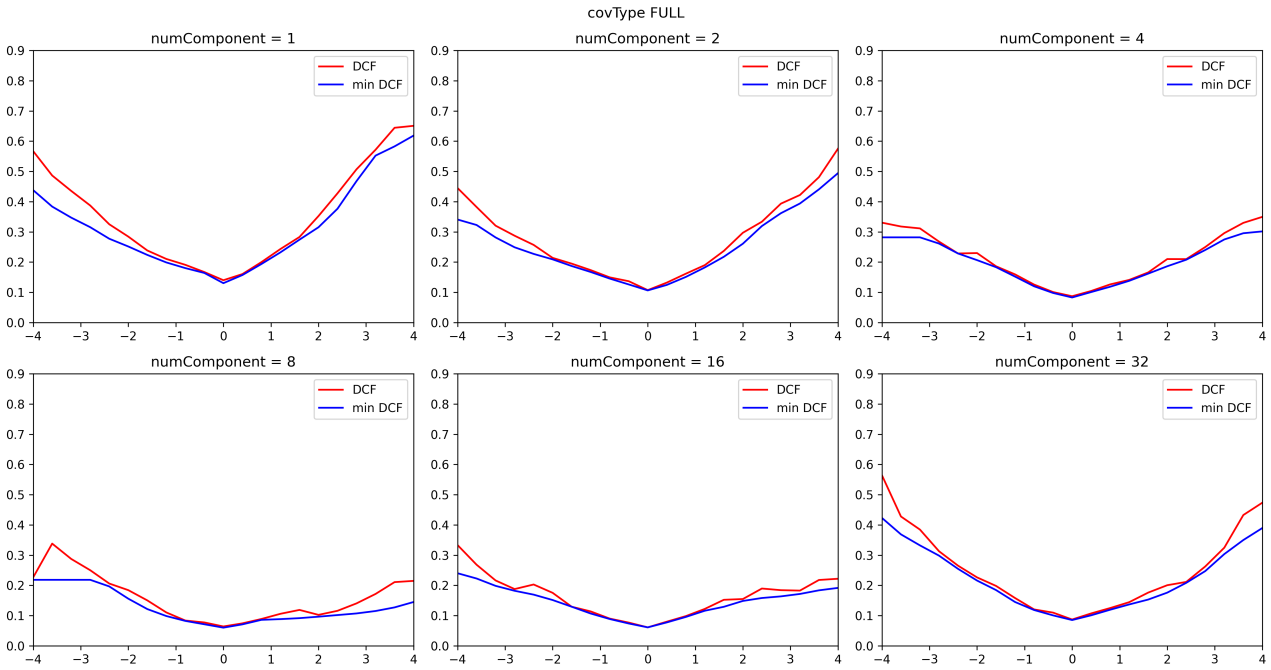


Figure 17: Shows minDCF and actDCF for GMM with Full Covariance and same number of components

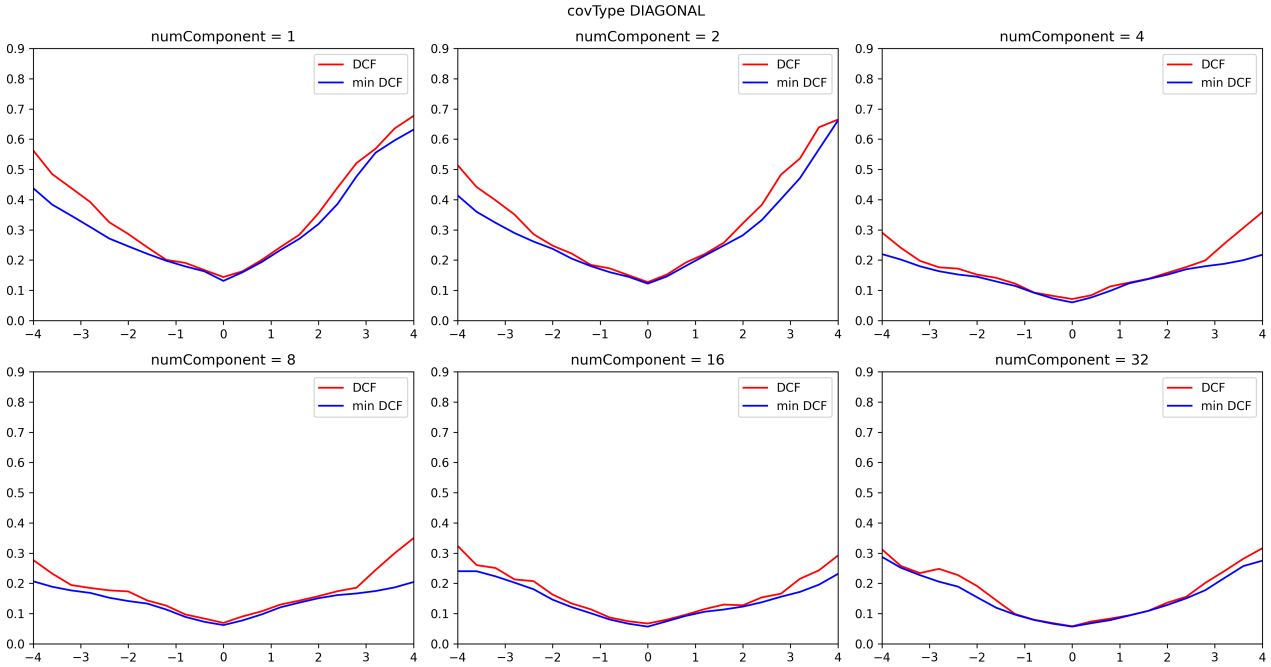


Figure 18: Shows minDCF and actDCF for GMM with Diagonal Covariance and same number of components

Subsequently, all possible combinations within the maximum value indicated above were tested. Table 13 shows what may be the most significant values. I chose these combinations because (1,16) and (2,16) are the two best combinations for the Full Covariance method, whereas (8,32) and (8,16) are the best for Diagonal Covariance method. Their representation is shown in Figure 19 and Figure 20.

GMM Model				
(nc0, nc1)	**Full Cov**		**Diag Cov**	
	minDCF	**actDCF**	**minDCF**	**actDCF**
(1, 1)	0.2629	0.3051	0.2570	0.3022
(2, 2)	0.2170	0.2337	0.2489	0.2674
(4, 4)	0.2161	0.2395	0.1481	0.1687
(8, 8)	0.1786	0.1928	0.1463	0.1809
(16, 16)	0.1631	0.1766	0.1622	0.1769
(32, 32)	0.2337	0.2499	0.1766	0.1989

Table 12: Show minDCF and actDCF for same number of components in GMM

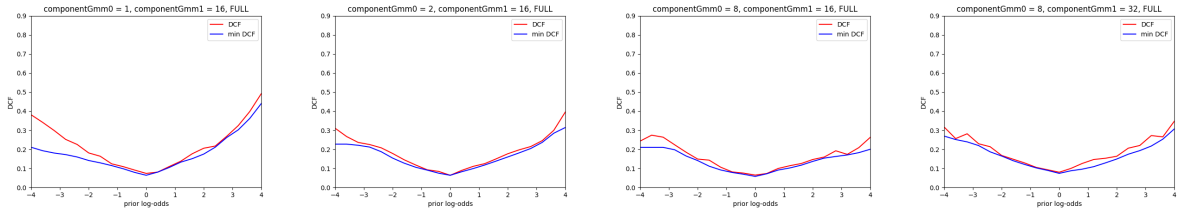


Figure 19: Shows minDCF and actDCF for Full Covariance method

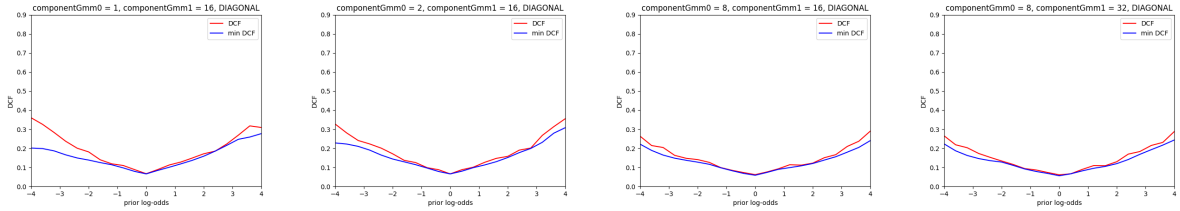


Figure 20: Shows minDCF and actDCF for Diagonal Covariance method

GMM Model				
(nc0, nc1)	**Full Cov**		**Diag Cov**	
	minDCF	**actDCF**	**minDCF**	**actDCF**
(1, 16)	0.1495	0.2055	0.1433	0.1807
(2, 16)	0.1701	0.1980	0.1536	0.1731
(8, 16)	0.1526	0.1725	0.1324	0.1487
(8, 32)	0.1745	0.1903	0.1312	0.1517

Table 13: Show minDCF and actDCF for same number of components in GMM

Summarize

It can be observed from the results obtained that:

- Too high or to low number of components leads to a worsening of the metrics considered.
- The best outcomes show that class 0 requires fewer components, meaning it requires a simpler model. In contrast, for class 1 it is observed that the number of components is greater this requires a more complex

model to avoid underfitting.

- Comparing the outcomes with previously applied methods shows that the performance is better but also better calibrated since the gap between minDCF and actDCF is low.
- GMM seems to prove to be the best performing model on our data set for several reasons. Because the combination of multiple Gaussian distributions allows for better modeling of more complex distributions. Also by estimating Gaussian parameters we can effectively model the distributions of the data we have, even if they are complex. Finally, through cross-validation we can choose the number of components we think is most correct for each class. For all these reasons we can say that in our case this seems to be the model that best fits.

6 Calibration

6.1 Calibration Consideration On Selected Models

Best Models

For each method, the best configuration was chosen by looking at the minDCF parameter, so the best configurations are:

- **Logistic Regression**: $\lambda = 3.162 * 10^{-2}$, minDCF is 0.2436 and actDCF is 0.4972
- **Support Vector Machine**: with RBF Kernel with $\gamma = 0.1$ and $C = 100$, minDCF is 0.1845 and actDCF is 0.3581
- **Gaussian Mixture Model**: with Diagonal convolution and $(nc0 = 8, nc1 = 32)$ where $nc0$ is the number of component for class 0 and $nc1$ is number of component for class 1, minDCF is 0.1312 and actDCF is 0.1517

In Figure 21 we see the behaviour of the 3 best models we have selected, now we will apply the calibration to these.

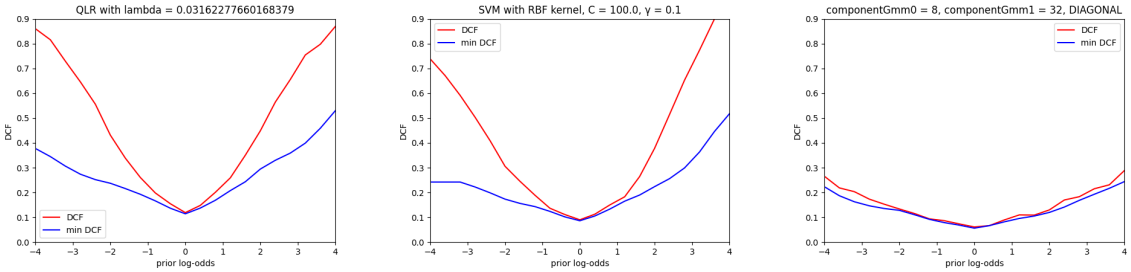


Figure 21: Shows minDCF and actDCF for three best models

6.2 Calibration Scores For Best Models

The evaluation is based on the theoretical threshold:

$$t = -\log \frac{\tilde{\pi}}{1 - \tilde{\pi}} \tag{37}$$

The threshold expressed in the Equation 37 does not guarantee that it is the optimal one, so a score analysis is applied. The function f that maps the uncalibrated score to the calibrated score has the form:

$$f(s) = \alpha s + \beta \tag{38}$$

Equation 39 can be interpreted as log-likelihood of the two classes:

$$f(s) = \log \frac{f_{S|C}(s \mid H_T)}{f_{S|C}(s \mid H_F)} = \alpha s + \gamma \tag{39}$$

Instead, class posterior probability for prior $\tilde{\pi}$ is:

$$\log \frac{P(C = H_T \mid s)}{P(C = H_F \mid s)} = \alpha s + \gamma + \log \frac{\tilde{\pi}}{1 - \tilde{\pi}} \tag{40}$$

From the previous equations we can write that:

$$\beta = \gamma + \log \frac{\tilde{\pi}}{1 - \tilde{\pi}} \tag{41}$$

Equation 39 can be rewritten as:

$$f(s) = \alpha s + \gamma = \alpha s + \beta - \log \frac{\tilde{\pi}}{1 - \tilde{\pi}} \tag{42}$$

6.3 Result Calibration

We apply calibration to the models we have chosen, so as to improve the actual performance by making them more similar to the predicted performance. Basically, we try to reduce the gap between minDCF and actDCF . In the Table 14 shows the results of the different models after calibration; the calibration was applied using different $\tilde{\pi}$ as shown in the table. The best results are obtained by applying GMM with a prior of 0.1 0.5

	Uncalibrated Models [minDCF - actDCF]		
QLR	0.2436 - 0.4972		
SVM	0.1845 - 0.3581		
GMM	0.1312 - 0.1517		
	Calibrated Models [minDCF - actDCF]		
	$\tilde{\pi} = 0.1$	$\tilde{\pi} = 0.5$	$\tilde{\pi} = 0.9$
QLR	0.2486 - 0.2721	0.2496 - 0.2609	0.2480 - 0.2657
SVM	0.1794 - 0.1894	0.1814 - 0.2032	0.1881 - 0.2020
GMM	0.1324 - 0.1518	0.1314 - 0.1518	0.1283 - 0.1559
Fusion	0.1304 - 0.1568	0.1373 - 0.1639	0.1382 - 0.1731

Table 14: Show minDCF and actDCF for different models before and after calibration

Figure 22 shows the Bayes error graphs of the different methods with a calibration prior of 0.1. We can see that an improvement can be observed for all configurations after calibration. Instead, by applying fusion, the scores of the different models are combined in an attempt to have an overall improvement, but it can be observed that the values of this method are very similar to those of GMM and are also less calibrated.

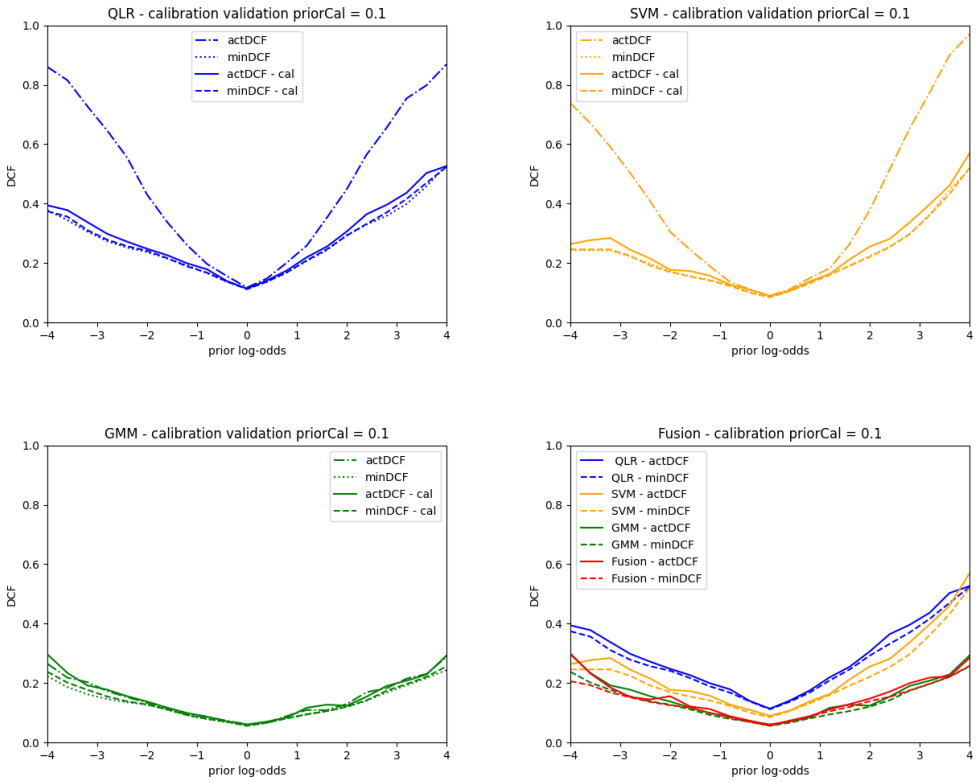


Figure 22: Shows result of each model before and after calibration

Summarize

Looking at the various outcomes, it is determined that the best method is GMM by applying a diagonal covariance matrix, as it is the one with the best minDCF and actDCF values.

7 Experimental Results

7.1 Evaluation

Starting with the model already tested, we go on to see how it behaves with another dataset which is the evaluation dataset.

The Figure 23 shows the result of each model before and after calibration with the evaluation dataset and a $\tilde{\pi} = 0.1$.

The Table 15 shows the minDCF and actDCF for different models before and after calibration on the evaluation dataset. From the values obtained, it is confirmed that GMM is again the method that gives the best performance, but Fusion also gives good results that do not deviate much from those of GMM.

As seen above GMM is the best method for our case study, it might be interesting to see what happens when changing the type of convolution and the number of components on the evaluation dataset, leading to a better result. Testing other strategies, we observe that the one with the best result is the one that applies a diagonal convolution, however, we observe that the number of components that improve the result change, becoming nc0=4 and nc1=16. This could be because the distribution of the data in the evaluation dataset is slightly

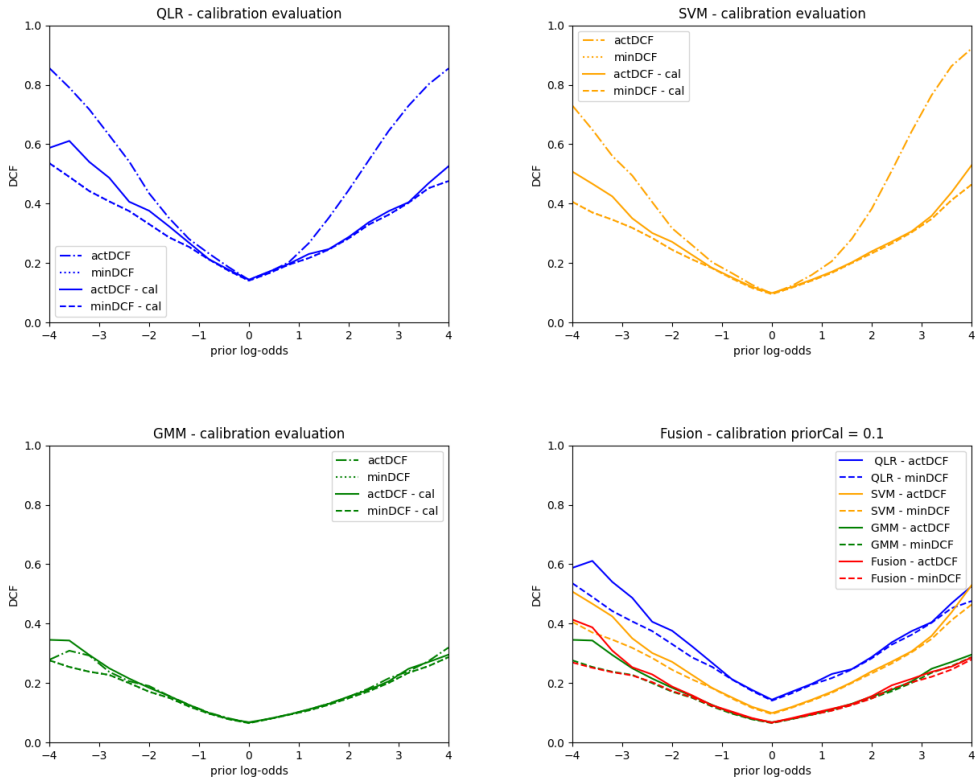


Figure 23: Shows result of each model before and after calibration with the evaluation dataset

	Uncalibrated Models [minDCF - actDCF]		
QLR	0.3515 - 0.4935		
SVM	0.2636 - 0.3634		
GMM	0.1838 - 0.2253		
Fusion	0.1979 - 0.2362		
	Calibrated Models [minDCF - actDCF]		
	$\tilde{\pi} = 0.1$	$\tilde{\pi} = 0.5$	$\tilde{\pi} = 0.9$
QLR	0.3515 - 0.3896	0.3515 - 0.3717	0.3515 - 0.3648
SVM	0.2636 - 0.2939	0.2636 - 0.2714	0.2636 - 0.2695
GMM	0.1838 - 0.2053	0.1838 - 0.2023	0.1838 - 0.2106
Fusion	0.1865 - 0.2046	0.1831 - 0.2056	0.1828 - 0.2089

Table 15: Show minDCF and actDCF for different models before and after calibration on eval dataset

different from the training dataset resulting in a better outcome on a different number of components to be used, the results are shown in the Table 16.

GMM chosen (Diag, nc0 = 8, nc1 = 32)		
	Uncalibrated	**Calibrated**
minDCF - actDCF	0.1838 - 0.2253	0.1838 - 0.2053
GMM other (Diag, nc0 = 4, nc1 = 16)		
minDCF - actDCF	0.1782 - 0.2400	0.1782 - 0.1970

Table 16: Show minDCF and actDCF for GMM chosen and another

8 Conclusion

Concluding our experiment, it can be observed that quadratic models perform better. Furthermore, among these models, it can be seen that those that make assumptions about the independence of the features don't lose performance, which means that the features are indeed not particularly correlated, and this is also confirmed by the GMM model, the best performing, that the features can be considered independent of each other.
Following on from what has been said about features independence, it can be seen that PCA did not lead to any real improvement, these might be possible because the original features are already independent and contribute uniformly to the variance of the data, hence PCA might not improve performance. This is because PCA may be useful when there are relationships between features that could be exploited to reduce dimensionality, but when this relationship does not exist, each component carries with it a small part of the total variance, thus making the application of PCA of little use. In conclusion, it can be said that the considerations made on our model, based on the training dataset, also proved useful on the evaluation dataset. In fact, the model chosen was the one that performed best on the latter dataset as well.