Dimensionality Reduction of a Pathological Voice Quality Assessment System Based on Gaussian Mixture Models and Short-Term Cepstral Parameters

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Abstract—Voice diseases have been increasing dramatically in recent times due mainly to unhealthy social habits and voice abuse. These diseases must be diagnosed and treated at an early stage, especially in the case of larynx cancer. It is widely recognized that vocal and voice diseases do not necessarily cause changes in voice quality as perceived by a listener. Acoustic analysis could be a useful tool to diagnose this type of disease. Preliminary research has shown that the detection of voice alterations can be carried out by means of Gaussian mixture models and short-term mel cepstral parameters complemented by frame energy together with first and second derivatives. This paper, using the F-Ratio and Fisher's discriminant ratio, will demonstrate that the detection of voice impairments can be performed using both mel cesptral vectors and their first derivative, ignoring the second derivative.

Index Terms—Cepstral parameters, F-Ratio, fisher's discriminant ratio, Gaussian mixture models, short-term analysis, voice disorders.

I. INTRODUCTION

COUSTIC analysis is a noninvasive technique based on the digital processing of the speech signal, which is an efficient tool for the objective support of the diagnosis of voice disorders, the screening of vocal and voice diseases (and particularly their early detection), the objective determination of vocal function alterations and the evaluation of surgical as well as pharmacological treatments and rehabilitation. Its application is not restricted to the medical area alone, as it may also be of special interest in the control of voice quality for voice professionals such as singers, speakers, etc. It furthermore offers two main advantages: it is a noninvasive tool, and it provides an objective diagnosis. It can be a complementary tool for those methods based on the direct observation of the vocal folds using laryngoscopy, as this inspection technique is considered risky and it has to be carried out by a specialist under well controlled conditions. Moreover, it is expensive, time consuming

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and requires costly resources, such as special light sources, endoscopic instruments and specialized video-camera equipment.

This study is mainly focused on organic pathologies affecting the vocal folds, appearing as a modification of the morphology of the excitation (i.e., vocal folds, increasing the distribution of masses) and producing a more irregular vibration pattern. This group may include pathologies such as polyps, nodules, cysts, sulcus, edemas, carcinoma, etc.

Most of the approaches found in existing literature address the automatic detection of voice alterations by means of longtime signal analysis [1]-[11]. These long-time parameters are generally calculated by averaging local time perturbations measured from the speech, so providing estimations of the degree of normality. Amongst these parameters are the following: pitch, jitter, shimmer, amplitude perturbation quotient (APQ), pitch perturbation quotient (PPQ), harmonics to noise ratio (HNR), normalized noise energy (NNE), voice turbulence index (VTI), soft phonation index (SPI), frequency amplitude tremor (FATR), glottal to noise excitation (GNE), etc. Previous studies [12]–[14] indicate that the detection of voice alterations can be carried out by means of the above mentioned long-term acoustic parameters, enabling each individual voice utterance to be quantified by a single vector. However, some of these parameters are based on an accurate estimation of the fundamental frequency, a fairly complex task in the presence of certain pathologies [15], [16].

In recent years, more modern approaches have been devised which use short-time speech analysis [17]–[19] or electroglotographic (*EGG*) [20]–[22] signals. Some of these focus on the automatic detection of voice impairments from the excitation waveform collected with a laryngograph [5], [21] or extracted from the acoustic data by inverse filtering [18], [22]. However, due to the fact that *linear prediction coding (LPC)*-based inverse filtering is based on the assumption of a linear model, such methods do not behave well when pathology is present due to nonlinearities introduced by the pathology itself. Some of the authors concerned have also proposed nonlinear signal processing for the same task [19], [22].

Throughout this research the automatic detection of voice impairments is carried out by means of Gaussian mixture models (GMMs) using the well-known short-term mel frequency cepstral coefficients (MFCC) as an alternative to the above mentioned methods. The main advantage of this approach is that it does not exhibit dependency on previous pitch estimations. Since this parameter has demonstrated a good degree of reliability for this task in previous research [23], the main idea in this paper is to reduce the dimensionality of the feature vectors in order to minimize complexity without affecting the recognition

performance. For this task, two methods extracted from the classical statistics have been used: the *F-Ratio* and the *Fisher's discriminant ratio*. Both are regarded as feature selection methods in existing literature, providing an idea about the significance of every single feature. The *Fisher* criteria and *F-Ratio* have been used on previous occasions [24] in speech recognition to reduce the vector length in a speech recogniser based on hidden Markov models (*HMM*) and both linear prediction cepstral coefficients (*LPCC*) and *MFCC*. In both cases 12 features and energy were used complemented with the first and second derivatives (up to 38 components), and it was demonstrated that the second order derivatives were the less significant, with some exceptions for the low frequency related features and the differences of energy.

In short-term analysis, the detection is carried out on a frame basis. Each frame is quantified with a vector formed by a feature set. In this approach, each feature set consists of the following features: the energy, L mel-cepstral coefficients, L first temporal derivatives, L second temporal derivatives of these coefficients—known as delta (Δ) mel-cepstrum and delta-delta ($\Delta\Delta$) mel-cepstrum, respectively—and the first and second derivative of the energy (known as delta energy and delta-delta energy). The final decision about the presence or absence of pathology is taken assuming independence between observations by summing the log-probabilities obtained for each frame of a patient's utterance and establishing a threshold to separate both sets. This parameterization approach has been used previously for this task in [23].

II. DATABASE

The company Kay Elemetrics distributes a CD-ROM database of approximately 700 records originally developed by the Massachusetts Eye and Ear Infirmary (MEEI) Voice and Speech Labs [25]. The acoustic samples are the sustained phonation of the vowel /ah/ (1–3 s. long) from patients with normal voices and a wide variety of organic, neurological, traumatic, and psychogenic voice disorders in different stages (from early to mature). The speech samples were collected in a controlled environment and sampled with a 50- or 25-kHz sampling rate and 16 bits of resolution. A downsampling with a previous half band filtering has been carried out over some registers in order to adjust every utterance to the sampling rate of 25 kHz.

Since we are interested in pathologies which affect the vocal folds, the process has been carried out over sustained phonations of the vowel /ah/. This is to ensure that the vocal folds remain in motion throughout the entire utterance.

The database has been segmented according to the criteria explained in [14]. The subset taken from the database contains 53 normal and 173 pathological speakers with a wide range of organic, neurological, traumatic and psychogenic voice disorders (Table IV). The criteria used in [14] ensure that gender and age are uniformly distributed between the two classes (Table I).

III. METHODS

The digital speech signal (1–3 s. long) is framed and windowed (a 40 ms. Hamming window was used throughout the various experiments) with no preemphasis [26], [27]. The frames were extracted with a 50% frame shift. The framing was followed by an endpoint detector [28] allowing voiced and unvoiced segments or silences to be separated. The next step was feature extraction, which was necessary in order to reduce the

TABLE I
DISTRIBUTION OF NORMAL AND PATHOLOGICAL SPEAKERS [14]

	Number		Mean ag	e (years)	Age rang	ge (years)	Std. deviation (years)		
	8	9	8	9	3	9	3	9	
Normal	21	32	38.81	34.16	26-59	22-52	8.49	7.87	
Pathol.	70	103	41.71	37.58	26-58	21-51	9.38	8.19	

pattern dimensionality and complexity. These feature vectors fed a *Gaussian mixture model*-based detector enabling a final decision about the absence or presence of pathology for every frame to be reached.

A. Computation of Recognition Features (MFCC)

It is widely recognized that the acoustic signal itself contains information about the vocal tract and the excitation waveform. The basic idea in this research is to use a short-term nonparametric approach capable of modeling the effects of pathologies on both the excitation (vocal folds) and the system (vocal tract), although throughout this paper emphasis has been placed on pathologies which mainly affect the vocal folds.

Mel cepstral coefficients [27] can be estimated by using a parametric approach derived from linear prediction coefficients (*LPC*), or by using a nonparametric *fast Fourier transform* (FFT)-based approach. However, FFT-based *MFCCs* typically encode more information from excitation; whereas *LPC*-based *MFCCs* remove this. This idea is demonstrated in [29], were FFT-based *MFCCs* are found to be more dependent on high-pitched speech resulting from loud or angry speaking styles than *LPC*-based *MFCCs*, which were found to be more sensitive to additive noise in speech recognition tasks. This is the case because *LPC*-based *MFCCs* ignore the pitch-based harmonic structure seen in FFT-based *MFCCs*.

It is well known that pathological voice is induced by an increase of mass, a lack of closure, or a change in the elasticity of the vocal folds. The result is that the movement of the vocal folds is not balanced and an incomplete closure of the vocal folds may appear in some or all glottal cycles. This is the reason why changes appear over the whole harmonic structure, (increasing the inter-harmonic energy and the fundamental frequency perturbation), and it also explains why energy increases at higher energy components due to aerial turbulence induced by an incomplete closure of the glottal clef.

FFT MFCCs [29] were considered to be appropriate for our purpose because in the presence of voice disorders they demonstrate an inherent ability to model either an irregular movement of the vocal folds, or a lack of closure induced by an increase of mass or due to a change in the properties of the tissue covering the vocal folds. The alterations related with the mucosal waveform due to an increase of mass are reflected in the low bands of the MFCC, whereas the higher bands are able to model the noisy components due to a lack of closure. Both alterations are reflected as noisy components with poor outstanding components and wide band spectrums. The spectral detail given by the MFCC can be considered good enough for our purpose.

MFCC parameters [27], [30], [31] are obtained by calculating the discrete cosine transform (DCT) over the logarithm of the energy in several frequency bands as shown in

$$c_m = \sum_{k=1}^{M} \log(S_k) \cdot \cos\left[m \cdot (k - 0.5) \cdot \frac{\pi}{M}\right]$$
 (1)

where m = (1 : L); L being the order of the MFCC coefficients, and S_K given by

$$S_k = \sum_{j=1}^{NFFT} W_k(j) \cdot X(j)$$
 (2)

where k = (1:M); M being the number of the mel bands in the mel scale, $W_k(j)$ is the triangular weighting function associated with the k_{th} mel band in the mel scale, and X(j) is the NFFT-point magnitude spectrum (j = 1: NFTT). It is usual to fix $M = \text{round } (3 \cdot \ln(\text{sampling frequency}))$.

Each band in the frequency domain is bandwidth dependant on the central frequency of the filter. The higher the frequency is, the wider is bandwidth is. This method is based on the human perception system, establishing a logarithmic relationship between the real frequency scale (Hz) and the perceptual frequency scale (mels). The suggested formula that models this relationship is as follows [27]:

$$F_{\text{mel}} = 2595 \cdot \log_{10} \left(1 + \frac{F_{\text{Hz}}}{700} \right).$$
 (3)

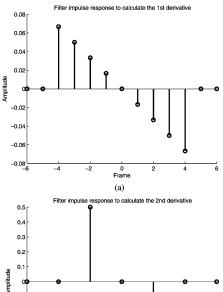
A representation providing an improved view of the dynamic behavior of speech can be obtained by extending the analysis to include information about the frame energy (quadratic sum of the amplitude divided by window length), and temporal derivatives of the parameters among neighbor frames [26], [27]. Both first (Δ) and second derivatives ($\Delta\Delta$) have been used in the present study. To introduce temporal order into the parameter representation, let us denote the m_{th} coefficient at time t by $c_m(t)$ (4))

$$\frac{\partial c_m(t)}{\partial t} = \Delta c_m(t) \approx \mu \cdot \sum_{i=-N}^{N} i \cdot c_m(t+i)$$
 (4)

where μ is an appropriate normalization constant and (2N+1) is the number of frames over which the computation is performed.

The first and second derivatives provide information about the dynamics of the time-variation in *MFCC* parameters. *A priori*, these features have been considered significant because, due to the presence of disorders there is a lower stability in the speech signal; therefore, larger time variations of the parameters may be expected in pathological speech relative to normal speech.

For each time frame t, the result of the analysis is a vector of L cepstral coefficients, L delta-



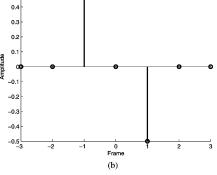


Fig. 1. Filter impulse responses used to calculate (a) the first derivative of the *MFCC* temporal sequence and (b) the second derivative of the *MFCC* temporal sequence. Both are anti-symmetric FIR filters.

delta coefficients, the energy, one delta energy and one deltadelta energy. This vector has $3 \cdot L + 3$ components and the dimensionality of the feature space is $D = 3 \cdot L + 3$, as follows:

$$o(t) = (E(t), c_1(t), c_2(t), \dots, c_L(t),$$

$$\Delta E(t), \Delta c_1(t), \Delta c_2(t), \dots, \Delta c_L(t),$$

$$\Delta \Delta E(t), \Delta \Delta c_1(t), \Delta \Delta c_2(t), \dots, \Delta \Delta c_L(t)) \quad (5)$$

where o(t) is a feature vector with D elements.

The calculation of delta (Δ) and delta-delta $(\Delta\Delta)$ was carried out by means of anti-symmetric moving-average *finite impulse response* (*FIR*) filters [Fig. 1(a) and (b)] to avoid phase distortion of the temporal sequence (length 9 for Δ , and 3 for $\Delta\Delta$).

The length of the *MFCC* vector used in the present study ranges from 10 to 26 in order to find the optimal dimensionality for the declared purposes.

B. GMM-Based Voice Pathology Detector

Let $x \in \Re^n$ be a random vector that has an arbitrary distribution. The distribution density of x is modeled as a Gaussian mixture density, a mixture of Q component densities, given by [32]

$$p\left(\frac{x}{\lambda}\right) = \sum_{i=1}^{Q} c_i \cdot p_i(x), \quad \sum_{i=1}^{Q} c_i = 1, \quad c_i \ge 0$$
 (6)

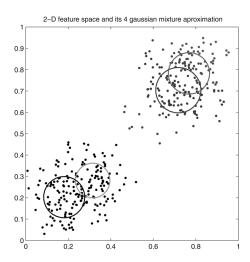


Fig. 2. Scatter plot of a two-dimensional (2-D) cepstral vector and its approximation by means of a 2-D Gaussian mixture.

where $p_i(x)$, i = 1, ..., Q are the component densities, and c_i , i = 1, ..., Q are the component weights. Each component density is an n-variate Gaussian function of the form

$$p_i(x) = \frac{1}{(2\pi)^{n/2} |C_i|^{1/2}} \exp\left[-\frac{1}{2} (x - \mu_i)^T C_i^{-1} (x - \mu_i)\right]$$

with μ_i the $n \times 1$ mean vector and C_i the $n \times n$ covariance matrix.

As the Gaussian components are acting together to model the overall probability density function (pdf), full covariance matrices are not necessary. For our purpose, the linear combination of diagonal covariance Gaussians has the capacity to model the correlation among feature elements. With this assumption, the parameters of the mixture can be represented by

$$\lambda = \{c_i, \mu_i, C_i\}, \qquad i = 1, \dots, Q$$
 (8)

where C_i is the covariance matrix.

The main motive for using the *GMM* as a representation of the acoustic space is that it has been demonstrated that a linear combination of Gaussian basis functions has a capacity to represent a large class of sample distributions [33], [34]. Fig. 2 shows how a Gaussian mixture is able to approximate the histogram of a given sequence.

The detector used is based on a Gaussian mixture model trained with an *expectation-maximization* (*EM*) algorithm [35]–[37]. The *EM* algorithm is an iterative method for learning maximum-likelihood parameters of a generative model where some of the random variables are observed, and some are hidden. The hidden random variables may represent quantities which we consider to be the underlying causes of the observables.

C. Dimensionality Reduction

There are a number of methods outlined in existing pattern recognition literature for reducing the dimensionality of a feature space. Several of these have been used in speech and speaker recognition with good results [24]. In this paper we study two feature selection methods: the F-Ratio, and F is discriminant ratio.

1) The F-Ratio: The F-Ratio [24], [38], [39] has been widely used as the figure of merit for feature selection in speaker recognition applications. It is defined as the ratio of the between-class variance and the within-class variance. In the context of feature selection for pattern classification, this ratio selects the features which maximize the scatter between classes. The following assumptions must be enforced: a) the feature vectors within each class must have Gaussian distribution; b) features should be uncorrelated; c) the variances within each class must be equal. Since the variances within each class are generally not equal, the pooled within-class variance is used to define the F-Ratio. To simplify, if the number of training patterns in each of the K classes is assumed to be the same (N), the F-Ratio is defined as

$$F_i = \frac{B_i}{W_i} \tag{9}$$

where B_i is the between-class variance and W_i the pooled within-class variance, both given by

$$B_i = \frac{1}{K} \sum_{k=1}^{K} (\mu_{ik} - \mu_i)^2$$
 (10)

$$W_i = \frac{1}{K} \sum_{k=1}^{K} W_{ik}$$
 (11)

where μ_{ik} and W_{ik} are the mean and variance of the *i*th feature for the *k*th class, and μ_i is the overall mean for the *i*th feature. These are given by

$$\mu_{ik} = \frac{1}{N} \sum_{k=1}^{K} x_{ikn} \tag{12}$$

$$W_{ik} = \frac{1}{N} \sum_{n=1}^{N} (x_{ikn} - \mu_{ik})^2$$
 (13)

$$\mu_i = \frac{1}{K} \sum_{k=1}^{K} \mu_{ik} \tag{14}$$

where x_{ikn} is the *i*th feature of the *n*th training pattern from the kth class.

- 2) Fisher's Discriminant Ratio: This ratio [24], [38], [39] represents the relationship between within-class and inter-class variances under the same assumptions as the F-Ratio. The following assumptions must be enforced: a) the feature vectors within each class must have Gaussian distribution; b) features should be uncorrelated; and, c) the variances within each class must be equal. Given a set of classes w_k , $k = 1, 2, \ldots, K$, twice scatter measurements can be defined as follows.
 - a) Within-class scatter (Sw): is a measurement of the scattering of the samples that belong to a class wk around their respective means

$$S_w = \frac{1}{K} \sum_{k=1}^K E\{(x - \mu_k)(x - \mu_k)^t\} \quad x \in w_k.$$
 (15)

b) *Interclass scatter (Sb):* measures the scattering of each class mean around the overall mean

$$S_b = \frac{1}{K} \sum_{k=1}^{K} (\mu_k - \mu_0)(\mu_k - \mu_0)^t$$
 (16)

where μ_k is the mean of class w_k and μ_0 is the mean value of the whole dataset without considering the class segmentation.

There exist several ways to quantify the discriminative power. The interclass separation measurement can be calculated comparing the relationship between within-class and inter-class scattering. The following is a standard computation:

$$J = |S_w^{-1} S_b|. (17)$$

A feature vector is said to be optimum if the inter-class separation is maximized. If computing these measurements is carried out for every single feature i alone, such measurements are known as Fisher's discriminant ratio F_i . The higher the value of F_i , the more important the feature is: this means that feature i has a low variance with respect to inter-class variance, and this is the reason why it is desirable to discriminate between them. These criteria adopt a special form in the one-dimensional, two-class problem, quantifying the separability capabilities of individual features

$$F_i = \frac{(\mu_{ic} - \mu_{i\overline{c}})^2}{\sigma_{ic}^2 - \sigma_{i\overline{c}}^2} \tag{18}$$

where subindex ${\cal C}$ corresponds to normal voices and $\overline{\cal C}$ to pathological.

D. Performance Measurements

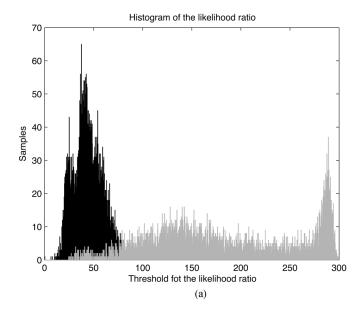
The likelihood ratio test is applied to determine whether the voice is to be classified as normal or not. Both models, λ_c and $\lambda_{\overline{c}}$, (normal and pathological) are obtained. For an utterance $X = \{\vec{x}_1, \vec{x}_3, \ldots, \vec{x}_T\}$ and a model λ_c , the likelihood ratio (applying Bayes' rule and disregarding the constant prior probabilities) in the log domain is

$$\Lambda(X) = \log \left[p\left(\frac{X}{\lambda_C}\right) \right] - \log \left[p\left(\frac{X}{\lambda_{\overline{C}}}\right) \right]. \tag{19}$$

The log-likelihood ratio is compared with a threshold θ and the voice is said to be pathological if $\Lambda(X)>\theta$ and normal if $\Lambda(X)<\theta$. The decision threshold is then set to adjust the tradeoff between rejecting pathological voices (false rejection) or accepting normal voices (false acceptance). Fig. 3(a) shows the histogram of the log-likelihood ratio for the false acceptance and false rejection of a typical case. Fig. 3(b) shows false acceptance and false rejection plots versus threshold θ . Both lines cross over the *equal error rate* point (*ERR*).

The log-likelihood of the utterance given the normal or pathological model is computed assuming independence between observations by summing the probabilities obtained for each frame over a patient's utterance

$$\log\left[p\left(\frac{X}{\lambda_C}\right)\right] = \frac{1}{T} \sum_{i=1}^{T} \log\left[p\left(\frac{x_i}{\lambda_{\overline{C}}}\right)\right]. \tag{20}$$



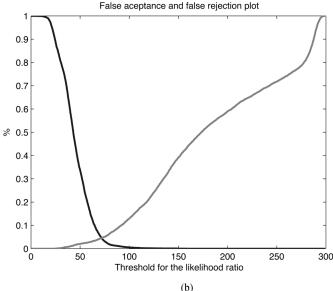


Fig. 3. (a) Histogram of the log-likelihood ratio for false acceptance and false rejection (frame accuracy). (b) Cumulative false acceptance (right) and false rejection (left).Both lines cross over the equal error rate point (ERR). The unit of analysis of the figure is frames.

The 1/T scale is used to normalize the likelihood for utterance duration (i.e., number of frames).

A block diagram of the detector is shown in Fig. 4. For a given input test utterance X, the choice is between s (X is a normal voice record) and n (X is a pathological voice record). Possible decisions are: S, the record is classified as normal; N, the record is classified as pathological.

E. Performance Evaluation

The k-fold cross-validation scheme [40] was used for estimating the classifier performance. The variance of the performance estimates was decreased by averaging results from multiple runs of cross validation where a different random split of the training data into folds is used for each run. In this study nine repetitions were used to estimate classifier performance figures.

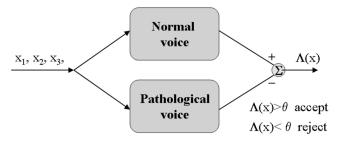


Fig. 4. Scheme of the detector. From each frame, the likelihood is calculated over both models.

TABLE II CONFUSSION MATRIX

EVENT							
		ABSENT	PRESENT				
DECISION		Efficiency (%)					
DECISION	ABSENT	TN	FN				
	PRESENT	FP	TP				

For each run of k-fold cross validation the total normal population and a randomly selected group of abnormals equal in size to the normal population was utilized. The performance was calculated by averaging the results obtained from each data set.

For each set, data files were split randomly into two subsets: the first for training (70%), and the second (30%) to simulate and validate results, keeping the same proportion for each class. The division into training and evaluation datasets was carried out on a file basis (not on a frame basis) in order to check and prevent the system from learning speaker-related features. Both male and female voices were mixed altogether in the training and validation sets. It is very important to use different speakers for training and validation in order to ensure that the system does not learn speaker dependent characteristics.

The training sets were used not only to extract the mixture parameters but also the value of θ . The threshold θ was calculated from the false acceptance and false rejection curves, averaging all the runs of the k-folds cross validation. According to the central limit theorem, the average of the log-likelihood ratio for false acceptance and false rejection follow a Gaussian distribution.

The number of speakers randomly selected from the subset of the database (segmented according to [14]) to build each set for cross-validation was 200 (53 normal and 147 pathological voices). This asymmetry is due to the fact that normal voice records last for approximately 3 s, whereas pathological voice records tend to be shorter as people with voice disorders have great difficulty holding a vowel over 2 or 3 s. Nine different sets were constructed for cross validation.

In order to evaluate the performance of the detector, and to enable comparisons to be made, several measurements (TP, TN, FN, and FP), ratios (SE, and SP) and curves [relative operating characteristic (ROC) and Detection Error Tradeoff (DET)] were taken into account (Table II).

- 1) True negative (TN): the detector found no event (normal voice) when indeed none was present.
- 2) True positive (TP): the detector found an event (pathological voice) when one was present
- 3) False rejection (FN): the classifier missed an event. Also called false negative

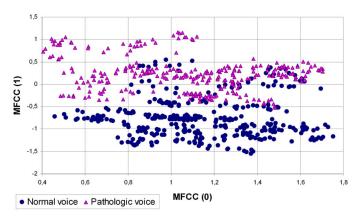


Fig. 5. The scatter plot represents the first *MFCC* versus the second *MFCC* coefficient for normal and pathological voices. (Color version available online at http://ieeexplore.ieee.org).

- 4) False acceptance (FP): the detector found an event when none was present. Also called false positive
- 5) Sensitivity (SE): likelihood that an event will be detected given that it is present

$$SE = 100 \cdot \frac{TP}{TP + FN}.$$
 (21)

6) Specificity (SP): likelihood that the absence of an event will be detected given that it is absent

$$SP = 100 \cdot \frac{TN}{TN + FP}.$$
 (22)

7) Efficiency (E): likelihood that the classification is correct

$$ET = 100 \cdot \frac{TN + TP}{TN + TP + FN + FP}.$$
 (23)

- 1) Relative Operating Characteristic (ROC) or Receiver Operating Characteristic: The ROC is a popular tool in medical decision-making [41], [42]. It reveals diagnostic accuracy expressed in terms of sensitivity (or true positive rate) against (1-specificity) (or false acceptance rate) at all possible threshold values in a convenient way. The area underlying the ROC can be used as an estimation of the probability that the perceived abnormality detected will allow a correct identification. This index varies from 0.5 (no apparent accuracy) to 1.0 (perfect accuracy) as the ROC curve moves towards the left and top boundaries.
- 2) Detection Error Tradeoff (DET): The DET [43], [43] has been used widely for the assessment of detection performance in speaker identification tasks. The DET curve plots error rates on both axes, giving uniform treatment to both types of error, and it uses a scale for both axes which spreads out the plot and better distinguishes different well performing systems and usually produces plots that are close to linear.

IV. RESULTS

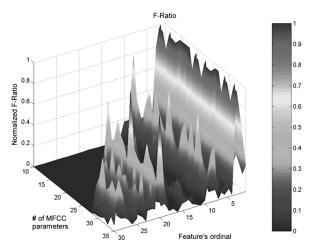
Fig. 5 represents the scatter plot of the first *MFCC* versus the second *MFCC* parameter. Although there is a clear overlapping,

	GMM 6 mixtures				GMM 8 mixtures				GMM 12 mixtures				
	AUC	Confussion Matrix		Efficiency (%)	AUC	Confussion Matrix		Efficiency (%)	AUC	Confussion Matrix		Efficiency (%)	
MFCC16	0.9976	86.4	13.5	93,2±3,8	0.9986	84.2	15.7	92.08±4.3	0.9957	84.2	15.7	92.10±4.5	
MI CCIO	0.5570	0	100	75.2-5,6	0.5500	0.01	99.9	72.00±4.3	0.5551	0	100	72.10=4.5	
MFCC22	0.9970	84.4	15.6	02.7+2.21	0.9970	82.8	17.2	02.115.22	0.9994	85.6	14.4	91.33±5.71	
		0	100	92.7±2.31		0	100	92.1±5.33		0	100		
VECC34	0.0007	88.2	11.8	04.07+2.20	0.0070	81	19	00 40 5 47	0.0005	72.5	27.5	05.06+5.1	
MFCC24 0	0.9997			94.07±3.28	0.9978^{-1}			90.48±5.47	0.9885			85.96±5.1	

0

100

TABLE III AREA UNDER ROC CURVE (AUC) AND CONFIDENCE MATRIX FOR SOME OF THE TEST PERFORMED USING $MFCC + E + \Delta + \Delta\Delta$ (FILE ACCURACY)



0

100

Fig. 6. F-Ratio for every test performed. F-Ratio is represented versus the number of MFCC parameters and the position of each feature in the feature vector, being E the first, MFCC(1) the second, MFCC(2) the third, ΔE the (L+2)th, $\Delta MFCC(1)$ the (L+3)th, $\Delta \Delta E$ the (2L+3)th, $\Delta \Delta MFCC(1)$ the (2L + 4)th, and so on.

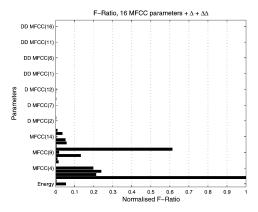
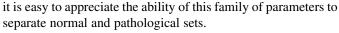
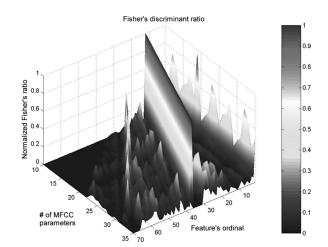


Fig. 8. Normalized F-Ratio plot using 16 MFCC features $+\Delta + \Delta\Delta$.



Training was carried out, with a number of MFCC static parameters ranging from 10 to 26. The energy (E) of the frame and dynamic features (Δ and $\Delta\Delta$) complemented every feature vector. GMM models were trained using 4, 6, 8, 10, 12, and 14 mixtures. The best performance was obtained using 16 MFCC features and a GMM with 6 centers. Independently of the combination of static and dynamic parameters used (E + MFCC +



0.5

99.5

Fig. 7. Fisher's discriminant ratio for every test performed, represented versus the number of parameters and the position of each feature in the feature vector, being E the first, MFCC(1) the second, MFCC(2) the third, ΔE the (L+2)th, $\Delta MFCC(1)$ the (L + 3)th, $\Delta \Delta E$ the (2L + 3)th, $\Delta \Delta MFCC(1)$ the (2L + 4)th, and so on.

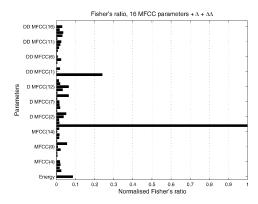


Fig. 9. Normalized Fisher's discriminant ratio plot using 16 MFCC features $+\Delta + \Delta \Delta$.

 $\Delta + \Delta \Delta$, E+MFCC+ Δ , or E+MFCC), the results are fairly similar. Table III represents some results for the most interesting working points: 16, 22, and 24 MFCC parameters, trained with a GMM of 6, 8, and 12 mixtures.

Fig. 6 shows the normalized F-Ratio for every test performed. The F-Ratio is represented in a three-dimensional (3-D) plot versus the number of MFCC parameters and each feature's position in the feature vector (its ordinal), being E

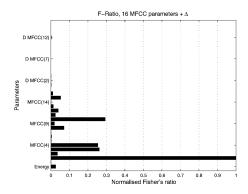


Fig. 10. Normalized F-Ratio plot using 16 MFCC features $+ \Delta$.

the first, MFCC(1) the second, MFCC(2) the third, ΔE the (L+2)th, $\Delta MFCC(1)$ the (L+3)th, $\Delta \Delta E$ the (2L+3)th, $\Delta \Delta MFCC(1)$ the (2L+4)th, and so on. It is easy to observe that the most important features following this criterion are concentrated in the first third of the feature vector (MFCC) static features). For simplicity reasons it was plotted only for the first 30 features. These results are shown better in Figs. 8 and 10 where a slice $(16\ MFCC)$ parameters) of the previous plot is represented. It is easy to observe that the first and second derivatives seem to be not significant under the assumptions of this criterion.

Under the assumption of Gaussian and uncorrelated feature vectors [24], Fig. 7 shows the normalized Fisher's discriminant ratio for every test performed. The Fisher's discriminant ratio is again represented in a 3-D plot versus the number of MFCC parameters and each feature's position in the feature vector. It can be observed that the most important features following this criterion are concentrated in the first and second third of the feature vector. These results are shown better in Figs. 9 and 11 where a slice (16 MFCC parameters) of the previous plot is represented. It is easy to observe that the second derivative seems to be not significant under the assumptions of this criterion, due mainly to the similarity among normal and pathological voices; however, first derivatives remain important.

In order to show the ability of the Δ and $\Delta\Delta$ features, Figs. 12 and 13 show, respectively, five different ROCs and DETs corresponding to different detectors developed from 16 MFCC + E. These detectors were developed by averaging the different runs of the k-folds cross validation scheme. The first detector was developed by feeding with E + MFCC features alone, leaving aside Δ and $\Delta\Delta$ parameters. The second detector was developed with input vectors made up of ΔE and $\Delta MFCC$ features alone. The third was formed with $\Delta\Delta E$ and $\Delta\Delta MFCC$ features alone. The fourth with MFCC + E and their Δs . And the fifth was constructed with MFCC + E, their Δs , and their $\Delta \Delta s$. For this purpose a 8 mixtures GMM was trained for every vector combination. The *ROCs* and *DETs* show that Δ and $\Delta\Delta$ features alone have no discrimination ability at all by themselves, whereas MFCC features alone achieve a good performance, slightly improved when complemented with Δ but with no statistical significance. It is clear that the classification abilities of the Δ and $\Delta\Delta$ alone are quite poor compared with

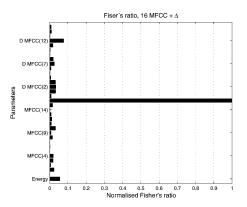


Fig. 11. Normalized Fisher's discriminant ratio plot using 16 MFCC features $\perp \Delta$

MFCC, but combined performance slightly improved the efficiency.

As a summary, Fig. 14 shows the efficiency of the test using an 8 mixture *GMM* model. Once again no relevant differences are observed when including either the first or second derivatives of the *MFCC* parameters. In view of this plot, no assumptions about the influence of Δ or $\Delta\Delta$ can be made due to the overlapped confidence intervals.

V. DISCUSSION

In speech, as in pattern recognition, the objective is not to obtain extremely representative models, but to eliminate recognition errors. The *GMM* provides a good estimation of the probability density function that corresponds to the hyperspace formed by the n-dimensional *MFCC* vectors, minimizing the number of misclassifications in the training data. It is well established that one of the main advantages of *GMM* is its ability to classify results correctly even when classes are similar. This methodology requires a shorter time for training than other approaches such *multilayer perceptron* (*MLP*) or *learning vector quantization* (*LVQ*) [44]. Furthermore, the *GMM* approach displays comparable accuracy with respect to *LVQ*, or *MLP*.

Looking at the F-Ratio plots it can be observed that the most important features are the independent parameters. The differences between the most and least important features are very clear. In view of the results, independent MFCC parameters can be considered with a higher significance than E, Δ , and $\Delta\Delta$. The first and second derivatives seem to be lacking in interest. Such a conclusion is reflected in the efficiency of the test where it is observed that the variations in efficiency are not significant when eliminating the energy, Δ and $\Delta\Delta$ (Figs. 12–14).

Fisher's criterion partially supports these conclusions. Independent parameters and Δ have a higher relevance, whereas $\Delta\Delta$ showed a lower significance.

VI. CONCLUSION

The proposed scheme may be used for laryngeal pathology detection. With respect to accuracy, it can be shown that each test yielded a false acceptance ratio lower than a false rejection ratio with an efficiency around $94 \pm 3.2\%$. In order to ensure that the features in the reduced space have physical meaning,

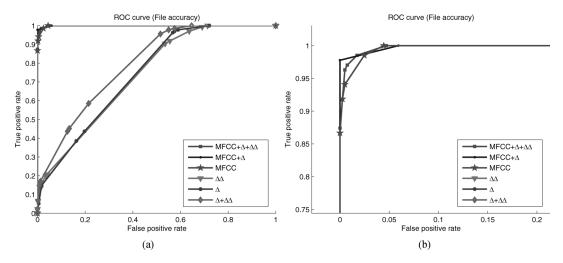


Fig. 12. (a) ROCs showing the true and false positive rate using 16 MFCC features alone (AUC = 0,996), Δ alone (AUC = 0,727), $\Delta\Delta$ alone (AUC = 0,721), $\Delta + \Delta\Delta$ (AUC = 0,797), MFCC + Δ (AUC = 0,998), and MFCC + $\Delta + \Delta\Delta$ (AUC = 0,998) and an 8 mixtures GMM. (b) Close-up of (a) plot close to TP = 1.

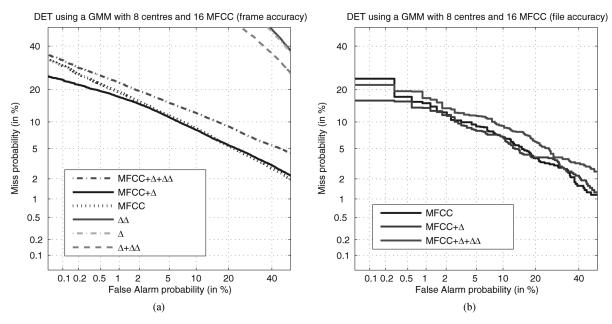


Fig. 13. DET plot calculated over 16 MFCC features alone, Δ alone, $\Delta\Delta$ alone, $MFCC + \Delta$, and $MFCC + \Delta + \Delta\Delta$ using an 8 mixtures GMM. The closer the plot is to the origin, the better the performance of the classifier is (a) frame accuracy and (b) file accuracy.

cepstral parameters complemented with their fist derivatives are considered the best solution for our purpose. We can conclude that the combination of the second derivatives do not show relevant influence on the results.

As in speaker and speech recognition [24], second temporal derivatives of *MFCC* parameters do not provide complementary information to the pattern recognition task of the automatic detection of voice impairments, increasing the computational complexity.

Regarding further future research, this scheme has to be tested using running speech. Preliminary studies have revealed that this approach could be used with text-dependent running speech maintaining performance. Only a small tuning would be required in the endpoint detector [28] to avoid not only silences, but also unvoiced frames.

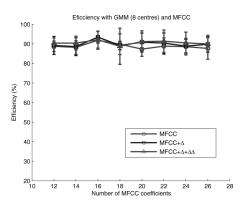


Fig. 14. Efficiency using a *GMM*-based detector with 8 mixtures and *MFCC* static parameters combined with Δ , with Δ and $\Delta\Delta$, and with energy Δ and $\Delta\Delta$.

TABLE IV FILES EXTRACTED FROM THE DATABASE

Normal		Pathological									
1. AXH1NAL.NSP	1.	ALB18AN.NSP	54.	JEG29AN.NSP	107.	MFC20AN.NSP	160.	TAB21AN.NSP			
2. BJB1NAL.NSP	2.	AMC14AN.NSP	55.	JFG08AN.NSP	108.	MPB23AN.NSP	161.	TDH12AN.NSP			
3. BJV1NAL.NSP	3.	AOS21AN.NSP	56.	JFN21AN.NSP	109.	MPF25AN.NSP	162.	TLP13AN.NSP			
4. CAD1NAL.NSP	4.	AXD19AN.NSP	57.	JHW29AN.NSP	110.	MPS09AN.NSP	163.	TLS09AN.NSP			
CEB1NAL.NSP	5.	AXT13AN.NSP	58.	JLD24AN.NSP	111.	MRB11AN.NSP	164.	TPP24AN.NSP			
6. DAJ1NAL.NSP	6.	BAH13AN.NSP	59.	JLS11AN.NSP	112.	MRC20AN.NSP	165.	TPS16AN.NSP			
7. DFP1NAL.NSP	7.	BEF05AN.NSP	60.	JMC18AN.NSP	113.	MWD28AN.NSP	166.	VAW07AN.NSP			
8. DJG1NAL.NSP	8.	BKB13AN.NSP	61.	JPP27AN.NSP	114.	MXC10AN.NSP	167.	WCB24AN.NSP			
DMA1NAL.NSP	9.	BLB03AN.NSP	62.	JRF30AN.NSP	115.	MXN24AN.NSP	168.	WFC07AN.NS			
10. DWS1NAL.NSP	10.	BPF03AN.NSP	63.	JTM05AN.NSP	116.	NFG08AN.NSP	169.	WJB06AN.NSP			
EDC1NAL.NSP	11.	BSG13AN.NSP	64.	JXC21AN.NSP	117.	NJS06AN.NSP	170.	WJP20AN.NSP			
12. EJC1NAL.NSP	12.	CAC10AN.NSP	65.	JXD30AN.NSP	118.	NKR03AN.NSP	171.	WPB30AN.NSP			
13. FMB1NAL.NSP	13.	CAK25AN.NSP	66.	JXF11AN.NSP	119.	NLC08AN.NSP	172.	WST20AN.NS			
GPC1NAL.NSP	14.	CLS31AN.NSP	67.	KAB03AN.NSP	120.	NMB28AN.NSP	173.	WXE04AN.NSP			
15. GZZ1NAL.NSP	15.	CMA06AN.NSP	68.	KAC07AN.NSP	121.	NMC22AN.NSP					
16. HBL1NAL.NSP	16.	CMR06AN.NSP	69.	KCG23AN.NSP	122.	NML15AN.NSP					
17. JAF1NAL.NSP	17.	CRM12AN.NSP	70.	KCG25AN.NSP	123.	NMV07AN.NSP					
18. JAN1NAL.NSP	18.	CTB30AN.NSP	71.	KDB23AN.NSP		OAB28AN.NSP					
19. JAP1NAL.NSP	19.	DAP17AN.NSP	72.	KJB19AN.NSP	125.	PAT10AN.NSP					
20. JEG1NAL.NSP	20.	DAS30AN.NSP	73.	KLC06AN.NSP	126.	PDO11AN.NSP					
21. JKR1NAL.NSP	21.	DBF18AN.NSP	74.	KLC09AN.NSP	127.	PGB16AN.NSP					
22. JMC1NAL.NSP	22.	DJP04AN.NSP	75.	KLD26AN.NSP	128.	PLW14AN.NSP					
23. JTH1NAL.NSP	23.	DMC03AN.NSP	76.	KMC22AN.NSP	129.	PMC26AN.NSP					
24. JXC1NAL.NSP	24.	DMP04AN.NSP	77.	KMS29AN.NSP	130.	PMD25AN.NSP					
25. KAN1NAL.NSP	25.	DRC15AN.NSP	78.	KMW05AN.NSP	131.	PMF03AN.NSP					
26. LAD1NAL.NSP	26.	DSC25AN.NSP	79.	KPS25AN.NSP	132.	RCC11AN.NSP					
27. LDP1NAL.NSP	27.	DSW14AN.NSP	80.	KTJ26AN.NSP	133.	RHP12AN.NSP					
28. LLA1NAL.NSP	28.	DVD19AN.NSP	81.	KXB17AN.NSP	134.	RJF22AN.NSP					
29. LMV1NAL.NSP	29.	DWK04AN.NSP	82.	KXH30AN.NSP	135.	RJL28AN.NSP					
30. LMW1NAL.NSP	30.	EAB27AN.NSP	83.	LAC02AN.NSP	136.	RJR15AN.NSP					
31. MAM1NAL.NSP	31.	EAS11AN.NSP	84.	LAD13AN.NSP	137.	RJZ16AN.NSP					
32. MAS1NAL.NSP	32.	EAS15AN.NSP	85.	LAI04AN.NSP	138.	RMB07AN.NSP					
33. MCB1NAL.NSP	33.	EEC04AN.NSP	86.	LAP05AN.NSP	139.	RPJ15AN.NSP					
34. MFM1NAL.NSP	34.	EED07AN.NSP	87.	LBA15AN.NSP	140.	RPQ20AN.NSP					
35. MJU1NAL.NSP	35.	EJH24AN.NSP	88.	LBA24AN.NSP	141.	RTL17AN.NSP					
36. MXB1NAL.NSP	36.	EMP27AN.NSP	89.	LES15AN.NSP	142.	RWC23AN.NSP					
37. MXZ1NAL.NSP	37.	ESS05AN.NSP	90.	LGM01AN.NSP	143.	RXM15AN.NSP					
38. NJS1NAL.NSP	38.	EWW05AN.NSP	91.	LJH06AN.NSP		RXP02AN.NSP					
39. OVK1NAL.NSP	39.	FMR17AN.NSP	92.	LJS31AN.NSP	145.	SAC10AN.NSP					
40. PBD1NAL.NSP	40.	FXC12AN.NSP	93.	LLM22AN.NSP	146.	SAE01AN.NSP					
41. PCA1NAL.NSP	41.	GDR15AN.NSP	94.	LNC11AN.NSP	147.	SAV18AN.NSP					
42. RHG1NAL.NSP	42.	GMM09AN.NSP	95.	LRD21AN.NSP	148.	SBF11AN.NSP					
43. RHM1NAL.NSP	43.	GMS05AN.NSP	96.	LVD28AN.NSP	149.	SCC15AN.NSP					
44. RJS1NAL.NSP	44.	GSB11AN.NSP	97.	LWR18AN.NSP	150.	SEC02AN.NSP					
45. SCK1NAL.NSP	45.	GXL21AN.NSP	98.	LXC01AN.NSP		SEF10AN.NSP					
46. SCT1NAL.NSP	46.	GXT10AN.NSP	99.	LXC06AN.NSP		SEG18AN.NSP					
47. SEB1NAL.NSP	47.	HJH07AN.NSP		LXR15AN.NSP		SEK06AN.NSP					
48. SIS1NAL.NSP	48.	HLM24AN.NSP		MAB06AN.NSP		SHD04AN.NSP					
49. SLC1NAL.NSP	49.	HXI29AN.NSP		MAM08AN.NSP		SJD28AN.NSP					
50. SXV1NAL.NSP	50.	HXL58AN.NSP		MCA07AN.NSP		SLC23AN.NSP					
51. TXN1NAL.NSP	51.	JAP02AN.NSP		MCW21AN.NSP		SLG05AN.NSP					
EQ VINCOLNIAL NICD	52.	JCC10AN.NSP	105	MEC06AN.NSP	158	SMA08AN.NSP					
52. VMC1NAL.NSP	52.	300107111.1101		MEC28AN.NSP		SWS04AN.NSP					

APPENDIX

Table IV shows the files included according the criteria used in [14] to segment the Kay Elemetrics Voice Disorders Database.

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