

# Deep-learning in Identification of Vocal Pathologies

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**Keywords:** Vocal Acoustic Analysis, Leave-one-out, Deep Neural Network, Architecture of Deep-NN, Dysphonia, Vocal Fold Paralysis, Laryngitis Chronica.

**Abstract:** The work consists in a classification problem of four classes of vocal pathologies using one Deep Neural Network. Three groups of features extracted from speech of subjects with Dysphonia, Vocal Fold Paralysis, Laryngitis Chronica and controls were experimented. The best group of features are related with the source: relative jitter, relative shimmer, and HNR. A Deep Neural Network architecture with two levels were experimented. The first level consists in 7 estimators and second level a decision maker. In second level of the Deep Neural Network an accuracy of 39,5% is reached for a diagnosis among the 4 classes under analysis.

## 1 INTRODUCTION

Voice is a sound resulting from a set of events in the vocal apparatus and along the vocal tract, with a certain force, sound, duration, speed and rhythm, subconsciously regulated by the information sent from the brain (PANEK, SKALSKI, GAJDA, & TADEUSIEWICZ, 2015).

The vocal acoustic analysis allows quantifying some characteristics of a sound signal. Using this technique for the study of voice, it is possible, in a non-invasive way, to determine and quantify the vocal quality of the individual through the different acoustic parameters that make up the signal (Guimarães, 2004).

Only the self-parameters obtained with the vocal acoustic analysis are not very conclusive, however, if they are properly associated with an Artificial Intelligence (IA) tool, the results are considerably better (Guimarães, 2004; Matuck, 2005; Teixeira J. P. et al, 2017)

The tools of IA are, therefore, an added value to take into account, since analyzing a large number of data, with several variables, it becomes difficult for the human being. After performing the training of an IA system, it is expected that the system be able to generalize. This means, for a new situation, never seen before, the system should be able to make a decision based on similar parameters seen before (Teixeira F., et al, 2018; Guedes et al, 2019).

A Deep Learning technic is based on the so-called artificial neural network. However, they have a greater number of hidden layers in their architecture, which helps in the processing of information and it can still have different activation functions in each hidden layer. The purpose of additional hidden layers is to have some more objective but non-final ‘image’ of the output.

In this work it was intended to do the same study that Teixeira F. et al (2018) did, however, instead of Support Vector Machine, a Deep Learning approach is used.

The main objective of this work is to distinguish between pathological and healthy subjects and to distinguish the different pathologies under study.

In this work, three groups of subjects were used. Subjects with Dysphonia, Vocal Fold Paralysis and subjects with the Laryngitis Chronica pathology. These diseases are the ones that most often cause disturbances in the human voice (University, 2018), being sometimes undetectable to the human ear.

Dysphonia is a disorder of the voice, often caused by abnormalities that affect the vibration of the vocal chords. This affects the ability to speak easily and clearly. Dysphonia is characterized by the symptoms of hoarseness, weak voice, changes in voice tone and it may arise suddenly or gradually (Teixeira J. P. & Fernandes P. O., 2015).

Laryngitis Chronica consists of an inflammation that can result from inhalation of irritants or by the intensive use of voice. Symptoms include gradual

loss of voice, hoarseness, and sore throat (Huche, F., & Allali, A., 2005; Kumar et al., 2010).

Paralysis of the Vocal Folds is the total interruption of the nervous impulse. Being this total, it happens in the two Vocal Folds, or being partial, occurs only in one of the Folds. This situation can occur at any age and the problems associated with this condition correspond to voice change, respiratory problems and swallowing problems.

For the analysis, the following parameters are used: relative jitter, relative shimmer, Harmonic to Noise Ratio (HNR), Noise to Harmonic Ratio (NHR), Autocorrelation and Mel Frequency Cepstral Coefficients (MFCC). These parameters were extracted from sustained vowels /a/, /i/ and /u/ in high, low and normal tones. The MFCC's were also extracted from continuous speech.

Next section describes the used methodologies. In this section, the parameters are presented, the materials used are identified and the methodology, based on deep neural network, is also presented. The section 3 presents the results and discussion. Finally, the last section summarizes the work developed and the conclusions.

## 2 MATERIAL AND METHODOLOGY

The parameters used in this work combine source parameters that are related with vocal folds and therefore with low frequency components (jitter, shimmer, HNR and Autocorrelation), with parameters related with vocal folds and vocal tract, spreading a large bandwidth of frequencies (MFCCs). The first set of parameters were extracted from a sustained speech of vowel. The MFCCs were extracted both from the same vowels but also from continuous speech.

The parameter used were just retrieved from the cured database (Fernandes, J. et al, 2019) with the parameters extracted from the Saarbrücken Voice Database (SVD) as detailed below. This cured database contains only the subjects diagnosed with just one pathology/condition, been rejected all subjects with more than one pathology/condition.

The speech samples were classified into 4 classes (3 pathologies/conditions and control) using one Deep Artificial Neural Network with architecture projected for this specific purpose.

### 2.1 Parameters

For this work, it was necessary to extract a set of parameters from acoustic speech files to build the cured database of speech parameters (Fernandes, J. et al, 2019). These parameters are relative jitter, relative shimmer, HNR, NHR, Autocorrelation and 13 MFCC's.

The jitter analysis for a speech signal is the mean absolute difference between consecutive periods, divided by the mean period and expressed as a percentage (Eq. 1).

$$jitter = \frac{1}{N-1} \frac{\sum_{i=1}^{N-1} |T_{i+1} - T_i|}{\frac{1}{N} \sum_{i=1}^N T_i} \times 100, \quad (1)$$

where  $T_i$  is the length of the glottal period  $i$ , and  $N$  the number of glottal periods.

The relative shimmer is defined as the mean absolute difference between magnitudes of consecutive periods, divided by the mean amplitude, expressed as a percentage (Eq. 2).

$$Shim = \frac{1}{N-1} \frac{\sum_{i=1}^{N-1} |A_{i+1} - A_i|}{\frac{1}{N} \sum_{i=1}^N A_i} \times 100, \quad (2)$$

where  $A_i$  is the magnitude of the glottal period  $i$ , and  $N$  the number of glottal periods.

The HNR is a parameter in which the relationship between harmonic and noise components provides an indication of overall periodicity of the speech signal by quantifying the relationship between the periodic component (harmonic part) and aperiodic (noise) component. The overall HNR value of a signal varies because different vocal tract configurations imply different amplitudes for harmonics. HNR can be given by Eq. 3.

$$HNR = 10 \times \log_{10} \frac{H}{1-H}, \quad (3)$$

where  $H$  is the normalized energy of the harmonic components and  $1-H$  is the remaining energy of signal considered the non-periodic components (noise).

In the autocorrelation function, the similarity of periods along the signal is evaluated. The greater the similarities, the greater the autocorrelation value (Guedes et al, 2019).

According to (Fernandes J. et al, 2018), mathematically the autocorrelation can be determined in 3 steps.

In the first step (equation 4), having a signal  $x(t)$  uses a segment of the same signal of duration  $T$ , centered on  $tméd$ . From the selected part, the average

of  $\mu x$  is subtracted and the result is multiplied by a window function  $w(t)$  to obtain a signal window:

$$a(t) = \left( x \left( tméd - \frac{1}{2}T + t \right) - \mu x \right) w(t), \quad (4)$$

The window function  $w(t)$  is symmetrical around  $t$  and  $0$  everywhere outside the time interval  $[0, T]$ . (Boersma, 1993) mentions that the window must be a sine or Hanning window, given by equation 5.

$$wt=12-12 \cos 2\pi tT \quad (5)$$

Then the normalized autocorrelation  $ra(\tau)$  of the selected signal part is calculated (equation 6). This is a symmetrical function of delay  $\tau$ :

$$ra(\tau) = ra(-\tau) \frac{\int_0^{T-\tau} a(t)a(t+\tau)dt}{\int_0^T a^2(t)dt} \quad (6)$$

Finally, it is necessary to calculate the normalized autocorrelation  $rw(\tau)$  of the window function used. Using the Hanning window, autocorrelation is obtained through equation 7.

$$rw(\tau) = \left( 1 - \frac{|\tau|}{T} \right) \left( \frac{2}{3} + \frac{1}{3} \cos \frac{2\pi\tau}{T} \right) + \frac{1}{2\pi} \sin \frac{2\pi|\tau|}{T}, \quad (7)$$

To estimate the autocorrelation  $rx(\tau)$  of the original signal segment, the autocorrelation  $ra(\tau)$  of the signal window is divided by the autocorrelation  $rw(\tau)$  of the window used (Eq.8).

$$rx(\tau) = \frac{ra(\tau)}{rw(\tau)}, \quad (8)$$

where  $ra(t)$  is the normalized autocorrelation of the signal and  $rw(t)$  is the normalized autocorrelation of a window used.

Therefore, the autocorrelation function of a sustained speech signal displays the local maxima for multiples of  $\tau$ , so it is only necessary to identify the first local maxima, which will correspond to the harmonic part.

The NHR parameter quantifies the relationship between the aperiodic component (noise) and the periodic component (harmonic part). Although it is the inverse of HNR, it is not measured in the logarithmic domain, so the values are not the inverse (Fernandes, J. et al, 2018). The NHR parameter can be given by equation 9.

$$NHR = \frac{N}{H} = \frac{H - Autocorr}{H} = \frac{1 - Autocorr}{1 - Autocorr} \quad (9)$$

The MFCC's parameters are based in the spectrogram of the signal. The extract process can be divided in 7 steps (Lindasalwa Muda, 2010).

The first step (Pre-Emphasis) is to emphasize the higher frequencies by increasing the signal energy at these same frequencies. The calculation for Pre-Emphasis is given by equation 10 where  $x[n]$  is the speech signal.

$$y[n] = x[n] - 0,95x[n - 1], \quad (10)$$

The second step (Framing) consists of dividing the signal into small frames, where it is advised that each frame should be between 20 and 40 milliseconds. The third step (window) is to multiply a window function by each frame of the signal.

In the fourth step, it is necessary to convert the  $N$  samples of each frame from the time domain to the frequency domain, using the discrete Fourier transform (Lindasalwa Muda, 2010).

Equation 11 is used for its calculation, where  $X(k)$  are the spectral coefficients,  $x(n)$  the signal frame. It should be noted that the values of  $n$  and  $k$  must be greater than or equal, to zero and less than or equal, to  $N-1$ .

$$X(k) = \sum_{n=0}^{N-1} x(n)e^{-\frac{i2\pi nk}{N}}, \quad (11)$$

The fifth step serves to make the transformation to the Mel scale.

To make this transformation, equation 12 is used. In this process, triangular filters are applied to the spectrum to make the conversion.

$$Mel(f) = 2595 * \log_{10} \left( \frac{f}{700} + 1 \right), \quad (12)$$

In the sixth stage comes the discrete cosine transform, which consists in transforming the Mel spectrum into the time domain. This transformation can be referred to as Mel Frequency Cepstrum Coefficient, where the lowest order coefficients represent the vocal tract shape and the higher order coefficients represent the waveform periodicity (Lindasalwa Muda, 2010; Tiwari, V., 2010).

Finally, in the last step the signal energy of a signal is calculated for a segment at time  $t1$  to time  $t2$  using equation 13.

$$Energy=x^2[t] \quad (13)$$

In this work 13 MFCC were used, and the first coefficient is the energy of the signal.

## 2.2 Materials

In this work, we used the German Saarbrücken Voice Database (SVD) (Barry, W.J., Pützer, n.a.) to extract the parameters associated with the various subjects used for the study. In this database is possible to find more than 2000 subjects. For each subject it is available a recording for 3 sustained vowels (/a/, /i/, /u/) in three different tones. There is also the German phrase: "Guten Morgen, wie geht es Ihnen?" (Good morning, how are you?). The sampling frequency of voice signals is 50 kHz. The file size is between 1 and 3 seconds with a resolution of 16 bits.

Subjects with Dysphonia, Laryngitis Chronica and Vocal Fold Paralysis were used because they are the pathologies with higher number of subjects. Subjects with more than one pathology/condition were rejected.

The parameters were extracted for 473 subjects, according to the groups shown in table 1.

In the work of (Teixeira J. P. et al., 2018) it was proved that there is no difference between the male and female gender for the parameters relative jitter, relative and absolute shimmer, HNR, NHR and autocorrelation for control subjects and Laryngitis Chronica subjects. Therefore, there was no separation made by gender.

The subjects available in SVD limited the size and mean age of each pathological group.

Table 1: Dataset ages by each pathologic group and control group.

Groups	Sample size	Average age	Standard deviation of ages
Control	194	38,1	14,4
Dysphonia	69	47,4	16,4
Laryngitis Chronica	41	49,7	13,5
Vocal Fold Paralysis	169	57,8	13,8
Total	473		

For every subject the parameters were extracted from sustained vowels, and the MFCC's parameters were also extracted from the continuous speech sentence. Three groups of parameters were organized. The first group contains only parameters related with the source. This group I, actually was structured into group I(a) and I(b).

Group I(a) contains the parameters Jitter, Shimmer and HNR for the 9 vowels by subject, in a total of 27 features.

The group I(b) contains the parameters Jitter, Shimmer, HNR, NHR and Autocorrelation for the 9 vowels by subject, in a total of 45 features.

The group II contains the 13 MFCC's extracted from the 9 sustained vowels, in a total of 117 features by subject.

The group III contains 13 MFCC's extracted from 50 overlapped segments of the continuous speech, in a total of 650 features by subject.

The jitter and shimmer parameters were extracted using the algorithm developed by Teixeira and Gonçalves (2016). The HNR, NHR, Autocorrelation and MFCC's parameters were extracted with the work developed by Fernandes J. et al (2019).

## 2.3 Deep Learning Architecture

The basic structure of the developed architecture is to have two levels. The first level consists in a set of multi-layer-perceptron neural networks (NN). Each one will give a guess if the sample corresponds to the class 1, class 2 or none (3 output classes). The 4 classes (3 pathologies and control) were combined into 6 NN. One additional NN were considered to classify between control/pathologic, only with binary classification. Therefore, the first level consists into 7 NN, presented in Table 2. The organizations of NN with 3 classes were used to allow the use of the all dataset to trains each NN.

The second level consists in a NN with 7 nodes in the input, hidden layers, and one output. The input receives the output of the 7 NN of first level. The output is the classification of one of the 4 classes.

Figure 1 presents a representation of the Deep NN developed.

It is expected that each NN of first level be specialized in the classification between two pathologies. The second level would receive the guess of this 7 specialized NN and take a final decision.

Table 2: Neural Networks for first level of Deep Learning.

NN 1	Healthy / Pathological
NN 2	Healthy / Dysphonia / Other
NN 3	Healthy / Laryngitis / Other
NN 4	Healthy / Paralysis / Other
NN 5	Dysphonia / Paralysis / Other
NN 6	Dysphonia / Laryngitis / Other
NN 7	Laryngitis / Paralysis / Other

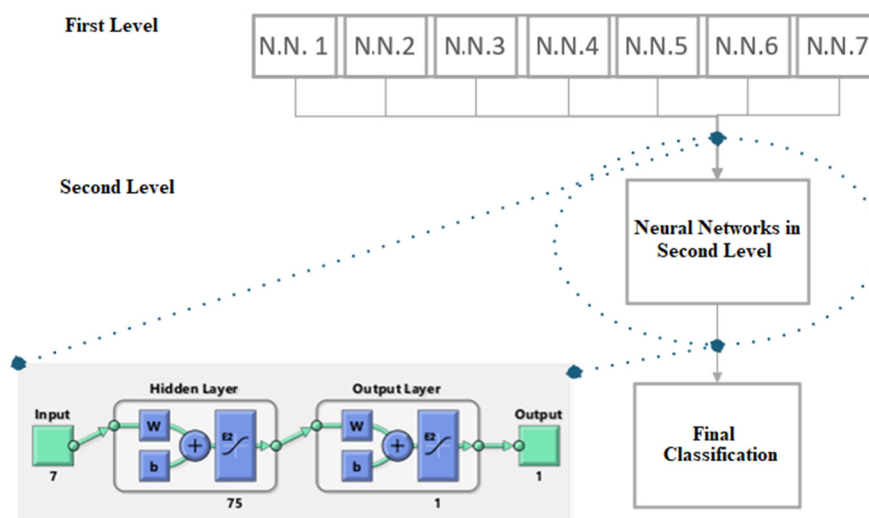


Figure 1: Deep Learning architecture.

For each parameters group, different NN Architectures were experimented.

The best architecture for the second level was achieved experimentally. It has 7 nodes in input layer (outputs of first level), 75 nodes in hidden layer and the training function is Levenberg-Marquardt (Marquardt, D., 1963).

The second level of Deep- Learning is responsible for classify the pathology of each subject. It is possible to classify between four classes, three pathological and one healthy.

The number of hidden layers were also experimented. One architecture with 2 hidden layers, where the second layer had 4 nodes, was expected to achieve better accuracy, supposing that each of these 4 nodes will classify one of the 4 classes. But the experimental result didn't show what was expected.

### 3 RESULTS AND DISCUSSION

The “leave-one-out” method has been implemented to train the Deep-NN. This method consists of testing all subjects, performing as many training sessions as subjects exist in the sample size. In N sessions of training, N-1 subjects were used to train and the remaining used to test. At the end, all subjects were tested and never used in the training process. This methodology is a time-consuming process but allows using a larger number of subjects to train and to test.

The result presented was measured for the subject in the test.

#### 3.1 First Level

Table 3 shows the results for the NN of the first level to classify between control/pathologic, using the tree groups of parameters. Acc is the Accuracy, Pre is the Precision, Sen corresponding to Sensibility, Spe is the Specificity and F1 corresponding to F1-Score. It can be seen that the higher accuracy is 72.7% using the parameters of group I(a).

Table 4 shows the results obtained in one of the remaining 6 NN of the first level as an example (Healthy/Paralysis/Others). The result for the remaining 5 NN has similar numbers. The measures are presented grouping the 3 output classes into groups of two classes to allow the determination of all the measures presented.

It can be seen that parameters of group I(a) achieve again, generally, higher accuracy.

Table 3: Measured values to classify healthy / pathological subjects.

Group of Parameters	I(a)	I(b)	II	III
Acc (%)	72.7	72.3	67.2	71
Pre (%)	69.6	68	53	57.2
Sen (%)	65.9	65.7	61.7	67.3
Spe (%)	78	77.2	70.3	73
F1	67.7	66.8	57	61.8

Table 4: Obtained values for classification between Healthy/Paralysis/Others with the different parameter groups.

		Healthy / Paralysis / Others				
		Parameters	Group I (a)	Group I (b)	Group II	Group III
Healthy / Others	Accuracy (%)		61,3	58,2	54,6	59
	Precision (%)		41,2	32,5	36	46,9
	Sensibility (%)		74,8	80,8	62,5	65,5
	Specificity (%)		55,6	51,7	51,2	55
	F1		53,2	46,3	45,8	54,7
Paralysis / Others	Accuracy (%)		63,7	60,8	59,2	60,5
	Precision (%)		60,1	71,1	67,4	63,1
	Sensibility (%)		45	37,9	35,5	38,5
	Specificity (%)		81,2	84,2	82,8	80
	F1		54,5	49,4	46,5	47,8
Others / (Healthy + Paralysis)	Accuracy (%)		49	43,6	45,3	49,8
	Precision (%)		60,9	69,1	63,6	55,5
	Sensibility (%)		26,2	24,9	25,7	26,4
	Specificity (%)		78,4	78,9	76,5	76,1
	F1		36,6	36,6	36,7	35,8

### 3.2 Complete Deep-NN

As mentioned previously, “Leave-one-Out” method has used, therefore, each subject was classify by eight neural networks (seven in first level and one in second leve), this process was repeat 473 times (number of subjects).

Table 5 presents the accuracy of the complete Deep-NN using three alternatives for the number of nodes in the hidden layer of the second level. It also presents the time consuming to train and test the all dataset using the “Leave-one-out” method. This values was obtained with processor Intel ®™ i5-3337u CPU@ 1.80GHz.

Table 5: Accuracy obtained and time consuming for the Deep-NN.

Nodes in hidden layer	Accuracy (%)	Time (minutes)
50	30.4	141.1
75	39.5	168.5
80	36.4	116.4

Table 5 shows the best accuracy to classify between the 4 classes is 39,5% using 75 nodes in the hidden layer.

Table 6 presents the confusion matriz for the best classification situation. The column of ‘Target’ corresponding the real situation of the subject, in ‘Classification’ line is mentioned the Deep-NN classification. The diagonal line has the number of subjects correctly classified. Where H is the Healthy subjects, D the Dysphonia, L the Laryngitis and P corresponds to Paralysis.

The confusion matriz shows that there is no problem with unbalance of the dataset.

Table 6: Confusion matrix.

		Classification			
		H	D	L	P
Target	H	98	62	27	7
	D	20	21	13	15
	L	9	13	8	11
	P	24	38	47	60

### 3.3 Discussion

Although several methodologies were tried, this article only contains the best results achieved.

The results shown that the neural network is a promising tool for voice pathology classification. In the work presented in (Teixeira F., Fernandes J., Guedes V., Junior A., & Teixeira J. P., 2018) the Support Vector Machines (SVM's) were used to classify between control/pathologic with the same dataset with best results about 70% accuracy. Comparing with present results of the first level NN an accuracy of about 73% was achieved, demonstrating the improvements introduced using ANN.

Guedes et al, 2019 developed a system to classify between the same 4 classes but using parameters from continuous speech and using transfer-learning techniques to do the classification. Very similar results were achieved, with F1-score of 40% for classify between four categories.

No separation by gender was used because for Laryngitis Chronica it was reported in (Teixeira J. P. et al., 2018) no gender difference for the relative jitter, relative and absolute shimmer, HNR, NHR and Autocorrelation. Anyhow, other conditions like Dysphonia and Vocal Fold Paralysis were used, and maybe some gender difference can exist for these subjects. Therefore, it is recommended to experiment a gender separation in future work.

## 4 CONCLUSIONS

This article describes the experience of using neural networks for diagnosis between healthy subjects and subjects with one of the three pathologies under study (Laryngitis Chronica, Dysphonia or Vocal Fold Paralysis). The parameters used in this analysis were extracted from sustained vowels or from a sentence.

A Deep NN was developed in two levels to classify between 4 classes.

The best results in the first level between control/pathologic subjects were an accuracy of 73%.

The best results between the 4 classes in the second level were an accuracy of about 40%.

The best results, in some cases, were not obtained with the same parameters group, however generally the parameters of group I(a) demonstrate the best results in higher number of cases. Therefore, for the uniformity reasons it was considered that group I(a) is the best group of parameters experimented. These parameters are relative jitter, relative shimmer and HNR.

As a final conclusion, the accuracy of about 40% to make the identification between healthy subjects and 3 pathologies is still below the requirements to become a real application. These results demand more research on this type of classification, experimenting different models of classification, different type of features, more subjects and maybe gender separation.

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