

1 تصنيف الاضطرابات الصوتية

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7 الملخص:

8 تقدم هذه المقالة تصنيف الأصوات المرضية، وهو مجال حيوي في الصحة الصوتية.

9 يقدم عملنا منهجية تدمج معالجة الإشارات، بما في ذلك تحليل الموجات واستخراج المتغيرات، ويكشف

10 عن نتائج وإعادة تم تحقيقها باستخدام نموذج نموذج آلة دعم المتجهات للتصنيف. الهدف الأساسي

11 هو تعزيز تشخيص وإدارة الاضطرابات الصوتية من خلال توفير أدوات أكثر فعالية لأخصائيي الرعاية

12 الصحية، استنادًا إلى قاعدة بيانات (Voiced). يؤكد هذا العمل على الأهمية الحاسمة للصحة

13 الصوتية وضرورة الاستثمار في طرق تشخيصية أكثر دقة وسهولة في الوصول إليها، مع فتح آفاق

14 جديدة للرعاية وجودة حياة الأفراد المصابين.

15 **الكلمات المفتاحية:** التصنيف، الأصوات المرضية، الإشارات الصوتية، تحليل الموجات، استخراج

16 المَعْلَمَات، نموذج SVM، قاعدة بيانات VOICED.



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سورية، يحتفظ المؤلفون بحقوق

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Classification of Voice Disorders	24
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Abstract:	30
This article introduces the classification of pathological voices, a vital area in vocal health.	31
Our work presents a methodology integrating signal processing, including wavelet	32
analysis and parameter extraction, and unveils promising results achieved using a Support	33
Vector Machine (SVM) model for classification. The primary objective is to enhance the	34
diagnosis and management of vocal disorders by providing more effective tools to	35
healthcare professionals, based on the VOICED database. This work underscores the	36
critical importance of vocal health and the necessity of investing in more precise and	37
accessible diagnostic methods, while also opening up new prospects for care and the	38
quality of life of affected individuals.	39
Keywords: Classification, Pathological Voices, Vocal Signal, Wavelet Analysis,	40
Parameter Extraction, SVM Model, VOICED Database.	41
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44 2.1 Introduction:

45 Brief Speech processing is a vast area of research
46 that requires the intervention of experts from several
47 specialties. Despite the remarkable development of
48 computer tools and programs, voice-controlled
49 systems have only become successful in recent
50 years. Early detection and precise of vocal
51 pathologies is crucial for appropriate medical
52 intervention. Traditionally, this detection relies on
53 the clinical expertise of health professionals, who
54 subjectively assess vocal alterations. However, this
55 approach has limitations in terms of objectivity,
56 reproducibility and precision. The advent of machine
57 learning technologies provides an unprecedented
58 opportunity to develop more objective and accurate
59 pathological voice classification methods.

60 Thus, the central question guiding our research is:
61 How can advances in machine learning be used to
62 improve the detection and classification of
63 pathological voices?

64 Advances in learning and classification technologies
65 have significantly simplified the development of
66 diagnostic methods and tools for vocal conditions.

67 The automatic detection and classification of
68 pathologies is a current field and still explored by the
69 research community (Malak & Ghulam & Mansour,
70 p. 571) A wide range of acoustic parameters was
71 used for pathology detection, namely pitch, jitter,
72 shimmer, p harmonic to noise ratio (HNR:
73 Harmonics to Noise Ratio), normalized noise energy
74 (NNE: Normalized Noise). Energy), cepstral
75 coefficients (MFCC: Mel-Frequency Cepstral
76 Coefficients), etc. In the field of automatic voice
77 pathology detection, various classifiers have been
78 proposed such as multi-layer perceptron, Gaussian
79 mixture model, probabilistic neural network, linear
80 discriminant analysis, k-nearest neighbor classifier (
81 KNN: K-Nearest Neighborhood), support vector
82 machines (SVM: Support Vector Machine), etc.

83 In (Lotfi & Haytham & et Adnène, 2009, p.3)
84 presented a method which is based on the use of a
85 multilayer neural network for the detection of voice
86 pathologies. The results highlight that the pitch
87 (fundamental frequency) and the first three formants

88 prove to be the most effective input parameters for
89 the distinction and identification of voices affected
90 by pathologies, thanks to the use of networks of
91 neurons. (Nesrine & Amina, 2016, p. 26) uses a
92 technique that relies on combining the continuous
93 wavelet transform with higher order statistics.
94 Classification is then carried out using support
95 vector machines (SVM).

96 (Hammami& Salhi & Labidi, 2020, p. 162)
97 developed a method to identify pathological voices
98 based on higher order features derived from analysis
99 of empirical modal decomposition and discrete
100 wavelet transform. The methodology of this study is
101 divided into three major steps. First, speech signals
102 undergo empirical mode decomposition (EMD),
103 followed by discrete wavelet transform (DWT).
104 From the coefficients from the DWT, several
105 characteristics relevant features are extracted,
106 including higher order features such as skewness,
107 kurtosis and variance, as well as other features such
108 as mean value, energy and entropy. These features
109 are then used to form data vectors representing the
110 speech signals. Finally, an SVM classifier is used for
111 the detection and classification of voices affected by
112 pathologies. The results presented in (MAROUA &
113 RAHMA,2020, p. 25) show the effectiveness of
114 SVM classifier to accomplish pathological voice
115 detection and classification. The accuracy rate
116 obtained using the SVM method and the use of Mel-
117 Frequency Cepstral Coefficients (MFCC)
118 characteristics are considered satisfactory for the
119 database used. On the other hand, (BOUDJELLABA
120 & BOUDJERIDA, 2021, p. 64) presents a method
121 that sets up a system for the identification of vocal
122 disorders using machine learning techniques. They
123 proposed two classification methods, namely
124 Support Vector Machine (SVM) and K Nearest
125 Neighbour (KNN), to evaluate their effectiveness in
126 detecting and classifying various vocal pathologies.
127 This system could be used by speech therapists to
128 perform an objective assessment of their patients'
129 voices, based on acoustic and aerodynamic
130 measurements such as Mel-Frequency Cepstral
131 Coefficients (MFCC), Jitter, Shimmer and
132 Harmonic-to-Noise Ratio (HNR).

133 As a result of the analysis of the results obtained, it
134 was observed that this method is very effective for
135 the detection of pathological and normal voices, as
136 well as for the classification of different types of
137 pathologies. The central goal of our work is to
138 significantly improve the diagnostic and treatment
139 capabilities of voice disorders. We aim to open new
140 perspectives for healthcare professionals by
141 providing them with more precise and objective
142 tools to assess and manage these conditions. At the
143 same time, we seek to raise awareness of the vital
144 importance of vocal health and highlight the
145 pressing need to invest in more accessible and
146 accurate diagnostic methods. Our commitment
147 focuses specifically on the effectiveness of voice
148 classification pathologies as a promising approach to
149 improve the quality of care and the quality of life of
150 the individuals concerned.

151 2.2 Literature Review:

152 2.1 Discrete wavelet transform

153 Describe any materials you used in your research,
154 and methods developed in your research. Discrete
155 wavelet, also known as discrete wavelet transform
156 (DWT), represents an analysis method applied to
157 discrete signals. Compared to the continuous
158 wavelet transform (CWT).

159 The steps to perform a simple TOD on a discrete
160 signal are:

- 161 • Selection of the base wavelet: We choose a
162 base wavelet which will serve as a pattern
163 for the transformation. Wavelets like Haar,
164 Daubechies, and many others are commonly
165 used.
- 166 • Decomposition steps:
 - 167 ➤ Step 1: Calculation of approximation and
168 detail coefficients: The first step consists of
169 calculating the approximation coefficients
170 (describing the large-scale components) and
171 the detail coefficients (describing the fine
172 details) from the original signal.
 - 173 ➤ Step 2: Downsampling: The calculated
174 approximation coefficients are then

- 175 downsampling by removing some samples to
176 achieve lower resolution.
- 177 ➤ Step 3: Repetition: The previous steps are
178 repeated on the approximation coefficients
179 obtained at each step to obtain
180 approximation and detail coefficients at
181 different scales.
- 182 • Repetition and scaling pyramid: The
183 decomposition steps are usually repeated
184 several times to obtain a scaling pyramid of
185 approximation and detail coefficients at
186 different resolutions
- 187 • The mathematical formula of DWT involves
188 convolution and subsampling of the parent
189 wavelet and the input signal. Here is how
190 this can be expressed generally:

$$191 DW(f, \varphi) = \sum_k a_{k,j} \cdot \varphi_{j,k}(t) + \sum_{j,k} d_{j,k} \cdot \phi_{j,k}(t)$$

192 Or :

193 $f(t)$ is the input signal.

- 194 • $a_{k,j}$ are the approximation coefficients at
195 scale j and position k .
- 196 • $d_{j,k}$ are the detail coefficients at scale j and
197 position k .
- 198 • $\varphi_{j,k}(t)$ is a dilated and translated version of
199 the parent wavelet $\varphi(t)$.
- 200 • $\phi_{j,k}(t)$ is a dilated and translated version of
201 the scaling function $\phi(t)$, which is related to
202 the mother wavelet and is used to calculate
203 the approximation coefficients.

204 The coefficients $a_{k,j}$ et $d_{j,k}$ can be calculated using
205 convolution filters and downsampling operations.

206 2. 2 Support vector machines

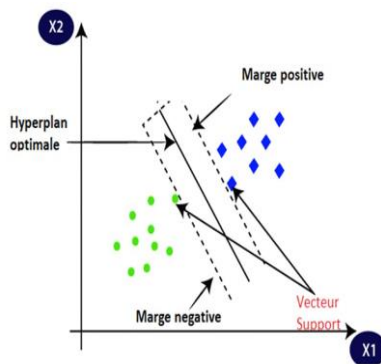
207 SVMs (Support Vector Machines) were invented by
208 scientists Vladimir Vapnik and Alexey
209 Chervonenkis in the 1960s. They have become a
210 popular tool for discriminative classification. An
211 exciting area of recent application of SVMs is in
212 speech processing. These models have a distinctly
213 different modeling strategy in detecting voice
214 disorders, compared to other classification methods
215 reported in the literature (Godino Llorente, &

216 Gómez-Vilda &. Sáenz-Lechón & Blanco-Velasco
 217 & CruzRoldán, Ferrer-Ballester & Angel, 2005, p.
 218 222). Supervised learning models called SVM are
 219 used to separate data points into different groups
 220 (classes) by determining an optimal hyperplane in a
 221 higher-dimensional feature space. The goal is to
 222 discover a dimensional hyperplane that maximizes
 223 the distance between data points of various classes.
 224 In its elementary form, when the two classes are
 225 linearly separable as illustrated in Figure. 1. this
 226 approach aims to identify a discriminating
 227 hyperplane represented by the equation next:

228
$$w \cdot x + b = 0$$

229 The elements of this equation are defined as follows:

- 230 ▪ w : Vector of weights ($w_1, w_2, w_3, \dots, w_n$).
- 231 ▪ x : Vector of attributes ($x_1, x_2, x_3, \dots, x_n$).
- 232 ▪ b : Threshold of the linear separator.



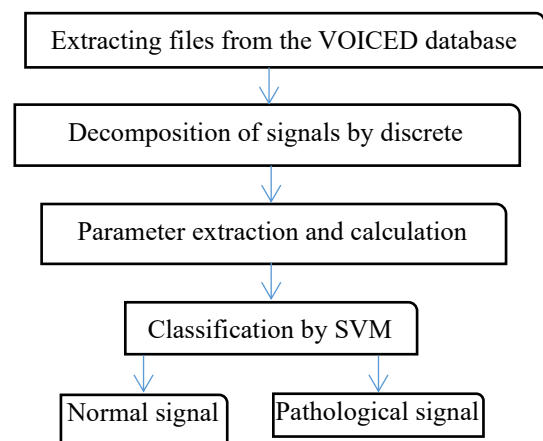
239 Figure 1- Example of classification by SVM

- 241 ✓ Hyperplane: The hyperplane divides the
 242 data into different classes. The task is to find
 243 the ideal hyperplane that minimizes
 244 classification errors while maximizing the
 245 margin between data points of different
 246 classes.
- 247 ✓ Margin: The distance between the
 248 separation hyperplane and the closest
 249 support vectors is represented by the margin
 250 of an SVM, and the optimization aims to
 251 find the hyperplane that maximizes this
 252 distance in order to obtain better separation
 253 between classes and better generalization of
 254 predictions.

- 255 ✓ Support vector: The data points closest to
 256 the separation hyperplane between the
 257 different classes are called support vectors.
 258 The position and orientation of the optimal
 259 hyperplane are determined by these
 260 vectors.

261 2.3 Proposed method

262 Our method goes through five steps; the
 263 following figure illustrates the block
 264 diagram of the classification of speech
 265 signals.



275 Figure 2 - block diagram of the classification of speech
 276 signals

277 a. Database:

278 The proposed algorithm is tested on the physionet
 279 database. This database includes 208 voice samples,
 280 from 150 pathological, and 58 healthy voices (Cesari
 281 & De Pietro & Marciano & Niri & Sannino, and
 282 Verde, 2018, p. 310). In detail, there were 73 male
 283 and 135 female participants. There is a prevalence of
 284 pathological voices compared to healthy ones, the
 285 former numbering 150 (52 male and 98 female), the
 286 latter 58 (21 male and 37 female).

287 b. Decomposition of wavelet signals

288 Our approach relies on the application of
 289 discrete wavelet decomposition to shape the
 290 feature vector of our speech signal. This
 291 decomposition is accomplished by being
 292 guided by the specific choice and order of the
 293 wavelets used in the calculation of the
 294 coefficients. The chunks of the speech signal

295 are fragmented into four levels using the
296 Daubechies wavelet 'db4', creating four details
297 and an approximation for each signal. Here are
298 the steps:

- 299 • Step 1: Signal Preparation: We take a
300 one-dimensional signal to analyze.
- 301 • Step 2: Filtering by the Low Pass Filter
302 (h): We apply a low pass filter (h) to the
303 signal to extract the low frequency
304 components. The result of the filtering
305 is the approximation signal (cA), which
306 contains the overall characteristics of
307 the signal at this scale.
- 308 • Step 3: Filtering by the High Pass Filter
309 (g): We apply a high pass filter (g) to
310 the signal to extract the high frequency
311 components. The result of the filtering
312 is the signal detail (cD), which contains
313 the fine variations and details specific
314 to the signal at that ladder.
- 315 • Step 4: Undersampling: We reduce the
316 temporal resolution of the
317 approximation signal (cA) and the
318 detail signal (cD) by retaining each
319 second sample.
320 This subsampling reduces the size of
321 the signals but preserves essential
322 characteristics at different scales.
- 323 • Step 5: Repeating the Process
324 (Iterations): We repeat steps 2 to 4 for
325 the approximation signal (cA) obtained
326 in the previous step. Each iteration
327 divides the signal into new
328 approximation (cA) and detail (cD)
329 components on a finer scale.
- 330 • Step 6: Repeat Until Desired Level: We
331 repeat steps 2 to 5 for a certain
332 predefined number of iterations
333 (decomposition levels) or until the
334 desired temporal resolution is reached.
- 335 • Step 7: Approximation and Detail
336 Coefficients: At the end of the process,
337 we obtain approximation (cA) and
338 detail (cD) coefficients at different
339 levels of decomposition.
340 The approximation coefficients contain
341 the overall characteristics of the signal

342 at different scales. Detail coefficients
343 contain the fine variations and specific
344 details of the signal at different scales.

345 c. Extracting parameters

346 After completing the wavelet transform, we
347 undertook the crucial step of extracting the
348 parameters. This phase is essential to capture the
349 distinctive characteristics of the signals vocal,
350 necessary to distinguish between pathological and
351 normal voices. We calculated a set of parameters that
352 provide information about different properties of the
353 speech signal, thus contributing to the construction
354 of a solid classification model. Before extracting the
355 parameters, we normalized the speech signal to bring
356 all signal values into a specific range, often between
357 -1 and 1. This step is crucial to avoid that the extreme
358 values do not bias subsequent calculations.
359 Normalization is performed as follows:

$$360 \text{ normalized signal} = \text{original signal} / \max(\text{original signal})$$

361 This normalization ensures that all signal amplitudes
362 fall between -1 and 1, making it easier to compare
363 and extract features.

364 ❖ A4 Approximation parameters:

365 • Energy: The energy of the signal is an
366 indicator of its overall intensity at that
367 frequency scale. It is calculated by summing
368 the squares of the amplitudes of the
369 normalized signal. Energy is an essential
370 characteristic for quantifying signal strength
371 or intensity at this scale.

$$372 \text{ Energy} = \sum_{i=1}^N (\text{signal } A4(i))^2$$

373
374 • Average: The average of the signal
375 amplitudes reflects its average level and can
376 be related to the overall loudness of the
377 normalized signal. A significantly different
378 average value could indicate a change in
379 signal level.

380 • Standard Deviation (SD): The standard
381 deviation measures the dispersion of
382 amplitudes around the mean. It provides
383 information on the variability of the
384 normalized signal. A high standard
385 deviation may indicate significant variations

386 in the signal, while a low standard deviation
 387 may indicate greater consistency.

388 • $SD = \sqrt{\frac{1}{N} \sum_{i=1}^N (signal\ A4(i)^2 - average)^2}$

389
 390 • Variance(V): The variance measures the
 391 dispersion of the A4 signal values around
 392 their mean.

393 $V = \frac{1}{N} \sum_{i=1}^N (signal\ A4 - average)^2$

394 • Average Frequency: The average
 395 frequency represents the weighted
 396 center frequency of the approximation
 397 signal. It provides information on the
 398 distribution of frequencies in the
 399 normalized signal.

400 ❖ Detail D4 Parameters:
 401 Similar steps were followed to extract
 402 the parameters from detail D4:

403 • Dominant Frequency: The dominant
 404 frequency in the signal is the one with
 405 the highest amplitude. It captures fine
 406 high-frequency variations of the
 407 normalized signal.

408 • Frequency Band Ratio: The frequency
 409 band ratio measures the energy
 410 distribution between the low and high
 411 frequency bands of the signal.

412 • Spectral entropy: Spectral entropy
 413 measures the complexity of the signal in
 414 terms of energy distribution over
 415 different frequencies.

416 **d. Classification (normal or pathological)**

417 To accomplish our goal of classifying voices into
 418 pathological and normal categories, we adopt a
 419 machine learning-based approach, specifically using
 420 an SVM. This method has proven effective in many
 421 classification tasks, including ours. We extracted
 422 relevant features from the voice recordings of the
 423 database used.

424 These features will be used as inputs for our SVM
 425 model. By adjusting the SVM parameters, we will
 426 seek to obtain the best possible separation, thus
 427 making it possible to efficiently classify new voices.
 428 Our choice of an SVM is based on its simplicity and
 429 effectiveness for moderately sized data sets, like the

430 one we handle with VOICED. Although the specific
 431 details of preprocessing, feature extraction have
 432 already been carried out, we will detail these steps in
 433 the next sections.

434 In addition, we will present our evaluation criteria
 435 which will measure the performance of our model.
 436 By capitalizing on the discriminative properties of
 437 the SVM, our objective is to create a tool for accurate
 438 and reliable classification to differentiate between
 439 pathological and normal voices. These advances
 440 open the way to various potential applications, both
 441 in the medical field and in other related fields.

442 ❖ **Measurement parameters**

443 The confusion matrix is a fundamental tool in
 444 classification model evaluation, and it is particularly
 445 useful in the context of pathological voice
 446 classification. It allows us to quantify the
 447 performance of our model by comparing its
 448 predictions with the actual values of the voice
 449 samples.

450 This matrix sorts all cases in the model into
 451 categories, determining whether the predicted value
 452 matched the actual value. All cases in each category
 453 are displayed in the matrix [22]. It has four main
 454 entries: TP, TN, FP and FN, organized as follows:
 455

456 Table 1- Confusion Matrix

	Negative Prediction	Positive prediction
Negative Label	True Negatives (TN)	False Negative (FN)
Positive Label	False Negative (FN)	True Positives (VP)

- 457 • Positive Label: The voice sample is indeed
 458 pathological.
 459 • Negative Label: The voice sample is indeed
 460 normal.
 461 • Positive Prediction: The model predicts that
 462 the voice sample is pathological.
 463 • Negative Prediction: The model predicts
 464 that the voice sample is normal.
 465 • TP represents the number of samples that
 466 actually belong to the positive class

467 (pathological voices) and that were correctly
 468 classified as such by the model.
 469 • TN represents the number of samples that
 470 actually belong to the negative class (normal
 471 voices) and that were correctly classified as
 472 such by the model.
 473 • FP represents the number of samples that
 474 actually belong to the negative class (normal
 475 voices), but were incorrectly classified as
 476 belonging to the positive class (pathological
 477 voices) by the model.
 478 • FN represents the number of samples that
 479 actually belong to the positive class
 480 (pathological voices), but were incorrectly
 481 classified as belonging to the negative class
 482 (normal voices) by the model.
 483 ❖ The sensibility:

484 Expressed as a true positive rate, sensitivity
 485 measures the ability of our model to correctly detect
 486 pathological voice samples. It is calculated as
 487 follows:

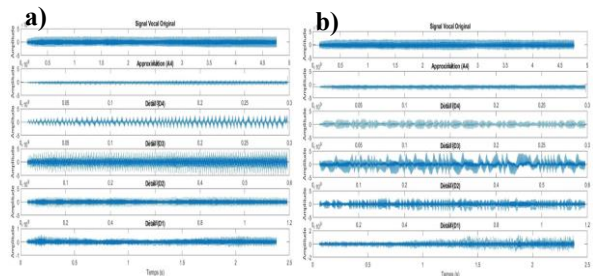
488
$$\text{Sensibility} = \frac{VP}{(VP+FN)} \times 100$$

489 **3. Results and Discussion:**

490 Our method was implemented in Matlab and used by
 491 a Physionet database. It goes through
 492 three stages; Decomposition of wavelet signals,
 493 Extracting parameters and Classification.
 494 Decomposition of wavelet signals is made up of
 495 seven stages. Figure 3 shows show that
 496 the wavelet decomposition up to the fourth level
 497 (A4, D4) for vocal signals: normal and
 498 pathological gives better frequency resolution.
 499 After, the extraction of the parameters are calculated
 500 for the approximation signals A4 and detail D4, both
 501 for a pathological voice and for a normal voice. The
 502 results obtained are presented in tables 2 and 3.

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520 Figure 3- Discrete 4-level wavelet decomposition of : a)
 521 normal voice b) pathological voice

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Table 2- Parameters of a normal voice signal

Parameters to calculate	A4	Parameters to calculate	D4
Energy	125.1481	Dominant Frequency	1835.9375
Average	-0.0788	Frequency Band Ratio	0.4799
Standard Deviation	0.2151	Spectral entropy	0.2617
Variance	0.0463		
Average frequency	2650.3964		

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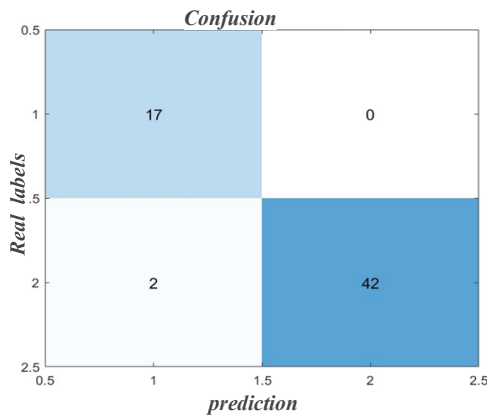
Table 3- Parameters of a pathological vocal signal

Parameters to calculate	A4	Parameters to calculate	D4
Energy	560.2596	Dominant Frequency	1101.5625
Average	-0.4033	Frequency Band Ratio	0.0213
Standard Deviation	0.2686	Spectral entropy	0.1851
Variance	0.0722		

528
 529

530 These parameters, calculated from the details and the
 531 normalized approximation, will serve as
 532 characteristics for our SVM classification model,
 533 aimed at differentiating pathological voices normal
 534 voices.

535 The following figure gives the results of the
 536 Confusion Matrix by the application of classification
 537 by SVM.



548 Figure 4 - Results of the confusion matrix

- 551 • This number (42) represents the number of
 552 pathological voice samples that were correctly
 553 identified as pathological by our model. In
 554 other words, these are the cases where our
 555 model made a positive (pathological)
 556 prediction that was correct.
- 557 • The number (2) of false negatives indicates
 558 the number of pathological voice samples that
 559 were misclassified as normal by our model.
 560 These samples were actually pathological, but
 561 our model misinterpreted them.
- 562 • This number (17) represents the number of
 563 normal speech samples that were correctly
 564 identified as normal by our model. In other
 565 words, these are the cases where our model
 566 made a negative (normal) prediction that was
 567 correct.
- 568 • The number (0) of false positives indicates
 569 that our model did not make errors in
 570 misclassifying normal voice samples as
 571 pathological. There are no cases where our
 572 model incorrectly predicted a normal voice
 573 sample as pathological.

574 The results of true negative (TN), true positive (TP),
 575 false negative (FN), false positive (FP), sensitivity,
 576 specificity and classification rate are shown in Table
 577 4:

578 Table 4 - Classification results obtained using
 579 SVM

Inputs	Results
TN	17
FN	2
TP	42
FP	0
Sensitivity	95.45%
Specificity	100%
Classification rate	96.72%

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 591 The obtained sensitivity means that our model
 592 correctly identified almost 95.45% of all
 593 pathological voice samples in our test set. In other
 594 words, our model demonstrated a high ability to
 595 accurately detect pathological cases among truly
 596 pathological samples.

597 The specificity of 100% indicates that our model
 598 correctly identified all normal speech samples in our
 599 test set. This means that our model made no errors in
 600 identifying normal speech samples, which is very
 601 positive.

602 The classification rate represents the overall
 603 accuracy of our model in terms of correct
 604 classification. It encompasses both true positives,
 605 true negatives, false positives, and false negatives,
 606 providing an overview of the overall performance of
 607 our model. Our model correctly classified almost
 608 96.72% of all samples, both pathological and
 609 normal, in our test set, demonstrating its overall
 610 effectiveness.

611 Table 5 presents a comparative study between our
 612 method and the methods of Variations of
 613 Pitch+neuron network [18], continuous wavelet
 614 transform + SVM [16], discrete wavelet transform +
 615 SVM [19], Mel-Frequency Cepstral Coefficients
 616 (MFCC) +SVM [20] and MFCC+SVM+ Variations
 617 of Pitch [21].

618 Reading Table 5 by line shows that the classification
 619 rate of our method is very high compared to other
 620 methods. The results obtained show the
 621 effectiveness of the proposed method.

622
 623

624
625 Table 5- Comparative study of our work with
626 other methods
627

References	Classification rate
[18]	85%
[16]	95.17%
[19]	93.1%
[20]	97%
[21]	95.70%
Our method	96.72%

628
629 **4. Conclusion**
630

631 This article was dedicated to the classification of
632 pathological voices, a crucial discipline for vocal
633 health. Through an in-depth exploration of the
634 foundations of the speech signal, a methodology
635 integrating signal processing and classification, as
636 well as the presentation of promising results
637 obtained thanks to an SVM model, we pursued the
638 major objective of improving the diagnosis and
639 treatment of vocal disorders.
640 Our investigations have highlighted the complexity
641 of the production of human speech and the diversity
642 of sounds it generates, whether sonorous or non-
643 sonorous. We highlighted the importance of
644 understanding vocal alterations and presented
645 methods for identifying vocal disorders,
646 highlighting voice parameters as key elements for
647 the analysis and classification of pathological
648 voices.
649 The methodology we developed, using the Physionet
650 database and the SVM model, has proven to be
651 effective, as evidenced by the high classification rate
652 of 96.72% that
653 we obtained. These promising results confirm the
654 relevance of our approach in the detection of vocal
655 pathologies.
656 However, it is essential to note that our work does
657 not stop here. Prospects for improvement remain, in
658 particular the use of richer and more diversified
659 databases to refine our models. Furthermore, the
660 exploration of advanced machine learning
661 techniques and the integration of artificial

662 intelligence could open new avenues in the field of
663 pathological voice classification.
664 Ultimately, this work represents a significant
665 contribution to improving the diagnosis and
666 management of voice disorders. It highlights the
667 need for increased attention to vocal health and
668 continued investment in more accurate and
669 accessible diagnostic methods. With a focus on the
670 effectiveness of pathological voice classification, we
671 Let us open up promising perspectives for the care
672 and quality of life of affected individuals.
673 We hope this work will stimulate further research in
674 the field of vocal health and help provide a brighter
675 future for people with vocal disorders.

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