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Abstract

The relationship between sleep apnoea-hypopnoea syndrome (SAHS) severity and the regularity of nocturnal oxygen saturation (SaO\textsubscript{2}) recordings was analysed. Three different methods were proposed to quantify regularity: approximate entropy (AEn), sample entropy (SEn) and kernel entropy (KEN). A total of 240 subjects suspected of suffering from SAHS took part in the study. They were randomly divided into a training set (96 subjects) and a test set (144 subjects) for the adjustment and assessment of the proposed methods, respectively. According to the measurements provided by AEn, SEn and KEN, higher irregularity of oximetry signals is associated with SAHS-positive patients. Receiver operating characteristic (ROC) and Pearson correlation

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analyses showed that KEn was the most reliable predictor of SAHS. It provided an area under the ROC curve of 0.91 in two-class classification of subjects as SAHS-negative or SAHS-positive. Moreover, KEn measurements from oximetry data exhibited a linear dependence on the apnoea-hypopnoea index, as shown by a correlation coefficient of 0.87. Therefore, these measurements could be used for the development of simplified diagnostic techniques in order to reduce the demand for polysomnographies. Furthermore, KEn represents a convincing alternative to AEn and SEn for the diagnostic analysis of noisy biomedical signals.

Keywords: Oxygen saturation, Entropy rate, Approximate entropy, Sample entropy, Kernel entropy, Density estimation
1. Introduction

Regularity is defined as the consistency of subpattern recurrence in a time series [1]. Regularity has shown to be a useful property of biomedical signals to discriminate those either generated by pathological systems or by the same system under different conditions [2]. Regular signals are characterised by a predictable behaviour, with recognizable patterns that repeat. Regularity is associated with the amount of information in a series, which, in a probabilistic sense, is a measure of the unexpectedness in the data [3]. Shannon [4] proposed the concept of entropy to evaluate the information (or uncertainty) in a message, which is modelled as a finite collection of random variables. In the context of infinite sequences or series, the entropy rate has been employed for the quantification of the amount of information [2]. Several metrics have been proposed to estimate the entropy rate of a series, with approximate entropy (AEn) [5] and sample entropy (SEn) [6] being the most common ones.

A generalised entropy measure is given by the family of Renyi entropies ($R_q$), where $q$ denotes the entropy order [7]. Lake [8] analysed the incorporation of the Renyi entropy into the entropy rate framework, showing that AEn and SEn approximate the differential Renyi entropy rate for $q = 1$ and $q = 2$, respectively.

AEn and SEn are based on the computation of probabilities by counting matches between signal subsequences of length $m$ and $m + 1$. A match is found when the distance between two subsequences is lower or equal than a tolerance parameter $r$ [6]. A different procedure to obtain the Renyi entropy rate of a series consists of substituting probability terms in AEn and SEn algorithms by the corresponding probability density functions [8, 3]. Several
advantages are found in this approach. It suppresses the need of predefined rules for the choice of the tolerance parameter $r$, which can be freely varied in order to obtain confident estimates of the density functions. In addition, entropy estimates made with different values of $r$ measure the same inherent quantity and can be compared directly [8, 9].

This approach requires the approximation of the (unknown) probability density function of the data, for which a finite set of samples extracted from the underlying series is initially available. Non-parametric kernel density estimation based on the Parzen window method has been suggested for this purpose [8, 3]. Specifically, Gaussian kernels are of special interest since they result in a smooth and continuous profile of the approximated density [10]. Additionally, in the case of the quadratic entropy ($R_2$), i.e., the Renyi entropy of order $q = 2$, Gaussian kernels lead to the exact evaluation of the integral found in its definition [3]. In a preceding study, a kernel-based estimation of $R_2$ was adopted to assess the quadratic entropy rate of a time series [11]. The resulting measure, termed as kernel entropy (KEn), was proposed as an indicator of the irregularity of the series [11, 12].

Entropy analysis has yield successful results in several applications involving time series processing such as earthquake forecasting [13], exchange rating [14] or fault detection [15]. Furthermore, entropy measures of biomedical signals have been widely used to assess physiological differences between subjects [16, 17]. The present study focuses on this scenario. We explored the utility of entropy rate measurements of nocturnal oxygen saturation signals ($\text{SaO}_2$) in the context of sleep apnoea-hypopnoea syndrome (SAHS) diagnosis. Nowadays, a definitive diagnosis about SAHS is obtained from
in-hospital evaluation of the patient’s sleep through nocturnal polysomnography (PSG). This test enables the assessment of SAHS severity by means of the apnoea-hypopnoea index (AHI), which quantifies the number of apnoea and hypopnoea events per hour of sleep. To obtain the AHI of a patient, the sleep specialist must evaluate a large amount of clinical and physiological data that, in addition to SaO₂ series, include other signals such as the electrocardiogram (ECG), the electroencephalogram (EEG) or the respiratory airflow (AF) [18]. Therefore, PSG is a highly complex and time-consuming procedure.

Reliable indicators of SAHS severity automatically extracted from these data would enable an objective and simplified interpretation. Nocturnal oximetry recordings are of special interest as they reflect respiratory dynamics during sleep. Apnoeas and hypopnoeas are usually accompanied by hypoxaemia due to airflow reduction, which is reflected by a marked drop in the saturation value [19]. The diagnostic utility of oximetry signals has been previously evaluated through different methods. A straightforward approach is the use of oximetry parameters based on the computation of desaturation events or the time spent below a certain level of saturation [20, 21]. In addition, complex signal processing and pattern recognition techniques like neural networks or genetic algorithms have been employed for the extraction of useful descriptors from SaO₂ data [22, 23, 24]. According to the reported results, a higher diagnostic accuracy can be obtained through the combination of different features including statistical, spectral and non-linear ones. Correct diagnostic rates close to 90% have been reported for screening algorithms based on this approach [25, 26, 22].
Among other features, SaO₂ irregularity measured by the entropy rate has been employed as a descriptor of the influence of SAHS severity on its dynamic behaviour [27, 25]. The non-deterministic occurrence of apnoic episodes tends to increase the uncertainty in the SaO₂ signal and, equivalently, its amount of information. As a result, signals from subjects suffering from SAHS are expected to have a higher entropy rate than those from control subjects. Previously, AEn has been employed to measure SaO₂ irregularity [28, 27]. These preceding studies showed the relationship between higher irregularity of oximetry signals and SAHS severity, estimating that a correct diagnosis based on regularity analysis can be obtained for approximately 85% of the patients. However, AEn has proven to be a biased entropy estimator [6] and, thus, further analysis is required to extract robust conclusions on the relationship between SAHS severity and SaO₂ irregularity.

To this end, the present study proposes a comparative analysis between different entropy metrics. In addition to AEn, we suggest entropy analysis of SaO₂ series based on SEn and KEn, which provide two different approaches to estimate the quadratic entropy rate of a signal. The present study aims to determine to which extent the irregularity of SaO₂ data is related to SAHS severity, as well as the most accurate method to quantify this relationship.

We hypothesise that a more confident assessment of the entropy of SaO₂ recordings can be obtained by means of kernel-based approximations to probability density functions as implemented by KEn. This method represents a novel approach for entropy estimation with respect to conventional procedures like AEn and SEn. The framework implemented by KEn suitably adapts to oximetry analysis since SaO₂ samples can be interpreted as ob-
servations of a continuous variable. Thus, probability density functions may provide a more reliable description of their statistical behaviour. This hypothesis is evaluated through an exhaustive regularity analysis of SaO₂ data using AEn, SEn and KEn.

2. Materials and methods

2.1. Subjects and signals

A total of 240 subjects suspected of suffering from SAHS took part in the study. They underwent PSG in the Sleep Unit of Hospital Universitario Pío del Río Hortega, Valladolid, Spain. The Review Board on Human Studies approved the protocol and each subject gave their consent to participate in the study. To draw useful conclusions on the effect of SAHS on SaO₂ dynamics, subjects affected by any other relevant respiratory disorder were excluded. The selected patients were continuously monitored using a polysomnograph (Alice 5, Respironics, Philips Healthcare, The Netherlands). A medical expert analysed the PSG recordings according to the rules proposed by Rechtschaffen and Kales [29]. Once apnoeas and hypopnoeas were identified, the AHI was obtained as the total number of events (i.e., the sum of apnoeas and hypopnoeas) divided by the total sleep time. The resulting value is expressed as the number of events per hour of sleep [30]. A threshold given by AHI = 10 hr⁻¹ was used to determine a positive diagnosis of SAHS [31].

A Nonin PureSAT pulse oximeter (Nonin Medical Inc., USA) was used to record oximetry signals at a sampling frequency of 1 Hz. These signals were subsequently saved to separate files to be processed offline. A prepro-
cessing stage was initially applied to remove artefacts like marked drops or zero samples due to a bad contact of the probe during sleep. The criteria suggested by Magaling et al. [32] were taken into account to perform signal preprocessing. According to these criteria, all changes greater than 4%/s between consecutive sampling intervals and any sample lower than 20% were removed.

Figure 1 shows two oximetry recordings from our dataset once artefacts were removed. The signals correspond to a normal subject (AHI = 0.5 h⁻¹) and a subject with severe SAHS (AHI = 32.1 h⁻¹), respectively. In addition, a detailed view (12 minutes) of both recordings is provided. The differences between these signals reflect the influence of repeated apnoeas and hypopnoeas on SaO₂ dynamics. The signal from the normal subject is characterised by a near-constant saturation value along the night, with small fluctuations around the baseline level. This behaviour is confirmed when observed in detail, as it exhibits some variability without marked desaturation events. In contrast, the profile of the signal from the subject with severe SAHS reflects a significant instability as a consequence of repeated desaturations accompanying apnoeas and hypopnoeas. As observed in the zoomed segment of the signal, these desaturation events are more frequent when compared with the oximetry recording from the normal subject. Additionally, they are more pronounced and longer. Therefore, a distinct value of the entropy rate can be expected for these signals since they reflect different dynamics.

INSERT FIGURE 1 AROUND HERE

The hold-out method was used to prevent bias in the estimation of the performance of the three entropy metrics [10]. Therefore, the initial popu-
lation was randomly divided into a training set with 96 subjects (40%) and a test set with 144 subjects (60%). The former was used to adjust user-dependent parameters in AEEn, SEn and KEEn algorithms. Signals in the test set were used to assess the diagnostic capability of these methods. Table 1 summarises the demographic and clinical data for subjects in training and test sets. Note that a higher proportion of SAHS-positive subjects was obtained due to the initial suspicion of SAHS in the population under study. In addition, the higher percentage of older males is motivated by the increased prevalence of SAHS in this group, as reported in previous studies. In the landmark study of the Wisconsin Sleep Cohort, including 602 men and women, 24% of men and 9% of women had AHI \( \leq 5 \) hr\(^{-1}\), while 9% of men and 4% of women had AHI \( \leq 15 \) hr\(^{-1}\) [33]. In addition, it has been estimated that AHI increases with age for both men and women [34].

**INSERT TABLE 1 AROUND HERE**

### 2.2. Methods

We compared the utility of three different entropy metrics, namely AEEn, SEn and KEEn, to quantify the relationship between SAHS severity and the irregularity of \( \text{SaO}_2 \) data. The proposed methods represent distinct implementations of the Renyi entropy rate of a series. The mathematical definition of the latter is derived from the expression of the Renyi entropy, which is obtained as follows. Let \( \mathbf{x} = [x_1, x_2, \ldots, x_N] \) be a continuous stochastic process composed of a sequence of \( N \) random variables \( x_i \). The Renyi entropy of the process is given by [7]:

10
\[
R_{q,N} = \frac{1}{1-q} \log \left\{ \mathbb{E} \left[ f_{q-1}^N (x_1, \ldots, x_N) \right] \right\} \\
= \frac{1}{1-q} \log \int f_q^N (x_1, \ldots, x_N) \, dx_1 \ldots dx_N
\]

(1)

where \( q \) determines the entropy order and \( f (x) = f (x_1, \ldots, x_N) \) is the joint probability density function of the set of variables \( x_i \) that compose the process. The Renyi entropy evaluates the amount of information (uncertainty) in \( x \).

Adding new variables to the process \( x \) will contribute to increase its information content, showing the dependence of the entropy on the process length specified by \( N \) [2]. Thus, a measure of the variation of the entropy due to the inclusion of a new variable \( x_i \), i.e., the entropy rate, can be obtained. The differential Renyi entropy rate is defined by the following asymptotic limit [8]:

\[
D_{q,N} = \lim_{N \to \infty} (R_{q,N+1} - R_{q,N}) \\
- \lim_{N \to \infty} \left\{ -\log \left[ \int f_q^N (x^{(N+1)}) \, dx^{(N+1)} \right] + \log \left[ \int f_q^N (x^{(N)}) \, dx^{(N)} \right] \right\}
\]

(2)

where the superscripts \( (N) \) and \( (N+1) \) denote the length of the process \( x \). In the following sections, a description of AEn, SEn and KEn is provided, showing the connection between each of these metrics and \( D_{q,N} \) as expressed in (2).

2.2.1. Approximate entropy (AEn)

AEn is a family of metrics developed by Pincus [5] for the analysis of noisy data such as biomedical signals. Briefly, AEn estimates the entropy rate
of a series from the computation of the probability of repetition for subsequences of length \( m \). To this end, two subsequences are considered similar if the distance between them is lower than a threshold \( r \). Mathematically, the algorithm to compute AEN is defined for a finite time series of length \( N \) given by \( \mathbf{x} = [x_1, \ldots, x_N] \). From this series, extract overlapping \( m \)-length windows, from \( \mathbf{x}_i^{(m)} \) to \( \mathbf{x}_{N-m+1}^{(m)} \), defined as \( \mathbf{x}_i^{(m)} = [x_i, x_{i+1}, \ldots, x_{i+m-1}] \). The distance \( d[\mathbf{x}_i^{(m)}, \mathbf{x}_j^{(m)}] \) between two such vectors \( \mathbf{x}_i^{(m)} \) and \( \mathbf{x}_j^{(m)} \) is calculated as the maximum absolute difference between their respective scalar components. A tolerance value \( r \) is used to obtain the number of vectors \((j = 1, \ldots, N - m + 1) \) such that \( d[\mathbf{x}_i^{(m)}, \mathbf{x}_j^{(m)}] \leq r \), which is denoted as \( N^m(i) \). Then, the likelihood that a vector is within a distance \( r \) from vector \( \mathbf{x}_i^{(m)} \) \((i = 1, \ldots, N - m + 1) \) is estimated as:

\[
C_r^m(i) = \frac{N^m(i)}{N - m + 1} \tag{3}
\]

The term \( C_r^m(i) \) reflects the regularity of patterns of length \( m \) similar to \( \mathbf{x}_i^{(m)} \) for a tolerance \( r \). Equivalently, it estimates the probability of observing the \( m \)-length vector \( \mathbf{x}_i^{(m)} \), implementing a discrete approximation to the probability density function \( f(\mathbf{x}^{(m)}) \).

The average of the logarithmic likelihood over the complete set of samples is computed as:

\[
\phi^m(r) = \frac{1}{N - m + 1} \sum_{i=1}^{N-m+1} \log C_r^m(i) \tag{4}
\]

From the previous analysis, \( \phi^m(r) \) represents an estimation of the expected value of \( \log[f(\mathbf{x}^{(m)})] \), which in turn corresponds to the negative value of the Renyi entropy of order \( q = 1 \) [8]:

12
\[ \phi^m (r) \approx E \left\{ \log \left[ f \left( x^{(m)} \right) \right] \right\} = \int \log \left[ f \left( x^{(m)} \right) \right] f \left( x^{(m)} \right) dx^{(m)} = -R_{1,N} \]

\[ \text{(5)} \]

Therefore, AEn approximates \( D_{1,N} \), the differential Renyi entropy rate of order \( q = 1 \) [8], as expressed by its mathematical definition [5]:

\[ AEn (m, r) = \lim_{N \to \infty} \left[ \phi^m (r) - \phi^{m+1} (r) \right] \]

\[ \text{(6)} \]

The following statistic is adopted for the computation of AEn on finite time series:

\[ AEn (m, r, N) = \phi^m (r) - \phi^{m+1} (r) \]

\[ \text{(7)} \]

2.2.2. Sample entropy (SEn)

SEn adopts a similar approach to AEn for the estimation of the entropy rate, which is based on the estimation of the probability of repetition for a subsequence. As a substantial difference, self-matching is prevented in the SEn algorithm. It has been shown that self-matching, i.e., the comparison of a vector \( x_i^{(m)} \) with itself, introduces some bias in the computation of AEn. Richman and Moorman proposed SEn in order to avoid this problem [6].

For a time series \( \mathbf{x} = [x_1, \ldots, x_N] \), only the first \( N - m \) vectors of length \( m \) are considered for comparison in the SEn algorithm. Let \( U_r^m (i) \) denote the probability that a \( m \)-length vector \( x_i^{(m)} \) is within a distance \( r \) from vector \( x_i^{(m)} \).
\[ U_r^m (i) = \frac{N^m (i)}{N - m - 1} \] (8)

where \( N^m (i) \) denotes the number of vectors \( x_j^{(m)} \) \((j \neq i, j = 1, \ldots, N - m)\) such that \( d \left[ x_i^{(m)}, x_j^{(m)} \right] \leq r \) and \( r \) is a tolerance value. Similarly to the term \( C_r^m (i) \) in the AEn algorithm, \( U_r^m (i) \) plays the role of the density function \( f (x^{(m)}) \), since it approximates the probability of observing \( x_i^{(m)} \). The average of the set of \( U_r^m (i) \) values is given by the quantity \( U^m (r) \):

\[ U^m (r) = \frac{1}{N - m} \sum_{i=1}^{N-m} U_r^m (i) \] (9)

which provides an estimation of the expected value of the function \( f (x^{(m)}) \):

\[ U^m (r) \approx E \left[ f \left( x^{(m)} \right) \right] = \int f^2 \left( x^{(m)} \right) dx^{(m)} \] (10)

It can be observed that the integral in the previous equation corresponds to the argument of the logarithm found in the definition of the Renyi entropy of order \( q = 2 \) or quadratic entropy. As a result, SEn can be interpreted as an estimation of the quadratic entropy rate as expressed by the following equation [6]:

\[ SEn \left( m, r \right) = \lim_{N \to \infty} \left\{ \log \left[ U^m (r) \right] - \log \left[ U^{m+1} (r) \right] \right\} \] (11)

SEn is estimated by the statistics:

\[ SEn \left( m, r, N \right) = \log \left[ U^m (r) \right] - \log \left[ U^{m+1} (r) \right] \] (12)

2.2.3. Kernel entropy (KEn)
Unlike AEn and SEn algorithms, which involve the computation of probabilities, KEn is based on modelling the statistical properties of the generator of the data. Hence, the algorithm to obtain KEn estimates the density function of the signal subsequences. To this end, KEn approximates the quadratic entropy rate from the use of non-parameteric probability density estimation techniques. The KEn assumes that the set of \( m \)-length vectors \( x_i^{(m)} \) from the original series has been generated according to the density function \( f(x^{(m)}) \) [8]. Hence, the Parzen window method with Gaussian kernels is proposed to estimate this function [11, 35, 3], resulting in the following expression:

\[
\hat{f}(x^{(m)}) = \frac{1}{N-1} \sum_{i=1}^{N-m+1} G \left( x^{(m)} - x_i^{(m)}, \Sigma \right) 
\]  

(13)

where \( G(x^{(m)}, \Sigma) \) denotes the zero-mean Gaussian kernel with covariance matrix \( \Sigma \) evaluated at point \( x^{(m)} \). In our study, spherical Gaussians with a covariance matrix given by \( \Sigma = \sigma I \) will be considered. The scalar \( \sigma \) is referred to as the kernel bandwidth and \( I \) denotes the identity matrix.

Using \( \hat{f}(x^{(m)}) \), the expected value of the density function \( f(x^{(m)}) \) is estimated by \( J^m(\sigma) \), which is given by:

\[
J^m(\sigma) = \int \hat{f}^2(x^{(m)}) \, dx^{(m)} 
\]  

(14)

Note that the use of a Gaussian kernel estimator for \( f(x^{(m)}) \) enables the computation of the exact value of the integral:
\[ \int \hat{f}^2 (x^{(m)}) \, dx^{(m)} = \frac{1}{(N - m + 1)^2} \sum_{i=1}^{N-m+1} \sum_{i=1}^{N-m+1} G \left[ x_j^{(m)} - x_i^{(m)}, 2\sigma^2 I \right] \]  

(15)

where \( N - m + 1 \) is the total number of \( m \)-length vectors in the original time series. The negative logarithm of \( J^m (\sigma) \) approximates the quadratic entropy \( R_{2,N} \). Hence, \( KEn \) is an estimation of the quadratic entropy rate, which is obtained from the incorporation of this expression in the Renyi entropy rate framework [8]:

\[ KEn (m, \sigma) = \lim_{N \to \infty} \left\{ \log [J^m (\sigma)] - \log [J^{m+1} (\sigma)] \right\} \]  

(16)

In practice, the following estimation is used for finite series:

\[ KEn (m, \sigma, N) = \log [J^m (\sigma)] - \log [J^{m+1} (\sigma)] \]  

(17)

It is worth noting that different techniques can be used to select the kernel bandwidth parameter \( \sigma \) of the Parzen density estimator from the original data. In this study, the Bayesian approach proposed by Zhang et al. [36] was applied for this purpose. According to this procedure, the elements of the covariance matrix \( \Sigma = \sigma I \) are treated as parameters and the aim is to estimate their posterior distribution \( f (\sigma \mid D) \), where \( D \) denotes the training set.

From Bayes theorem [10], the posterior density \( f (\sigma \mid D) \) is proportional to the product of the prior density \( f (\sigma) \) and the likelihood \( f (D \mid \sigma) \):

\[ f (\sigma \mid D) \propto f (D \mid \sigma) f (\sigma) \]  

(18)
Note that the likelihood of observing the sample set $D$ for a given value of $\sigma$ can be approximated from the expression of $\hat{f}(x^{(m)})$. Therefore, the prior probability $f(\sigma)$ is the remaining factor to obtain $f(\sigma \mid D)$. Zhang et al. suggested the following functional form for the prior in order to make the sampling algorithm work properly [36]:

$$
f(\sigma) \propto \prod_{k=1}^{m} \frac{\sigma_k}{\lambda + \sigma_k^2}
$$

(19)

where $\lambda$ controls the shape of the function. This prior density function aims to avoid large values of $\sigma$, for which the associated probability is small.

The most probable value of $\sigma$ given the data in $D$ is selected as the optimum. This value is obtained by sampling from the posterior probability $f(\sigma \mid D)$ using Markov Chain Monte Carlo (MCMC) techniques. The Metropolis-Hastings algorithm was used for this purpose [37, 38].

2.3. Statistical analysis

To assess the performance of the proposed entropy methods in the quantification of SaO$_2$ irregularity derived from SAHS, two different approaches were used. First, a two-class classification model was defined by dividing the initial population into SAHS-negative and SAHS-positive subjects. Receiver operating characteristic (ROC) analysis was used to assess the capability of AEn, SEn and KEEn measurements for identifying SaO$_2$ signals from these two groups [39]. The area under the ROC curve (AUC), which represents the probability of correct classification for a randomly chosen pair of samples from the two possible categories [39], was used as a measure of classification performance.
The second model for SAHS diagnosis consists in estimating the AHI from the available oximetry signal. According to the value of this index, subjects are assigned to one of four severity groups [18]: no SAHS (AHI < 5 hr⁻¹), mild SAHS (5 hr⁻¹ ≤ AHI < 15 hr⁻¹), moderate SAHS (15 hr⁻¹ ≤ AHI ≤ 30 hr⁻¹) and severe SAHS (AHI > 30 hr⁻¹). The utility of the three entropy measures to rank SAHS severity was assessed by means of the Pearson’s correlation coefficient (ρ). It evaluates the linear relationship between AEn, SEn and KEn and the AHI. The value of ρ can be interpreted as the utility of a given method to predict the AHI.

3. Results

3.1. Selection of the input parameters

Entropy analysis based on the proposed methods require the prior selection of several parameters. These correspond to N, m, and r for AEn and SEn algorithms. In the case of KEn, only N and m are needed since a data-driven technique was used to optimise the value of σ. The parameter N denotes the length of the time series to be processed. In our study, the length of oximetry recordings was approximately 7 hours (i.e., more than 25000 samples), involving a large amount of data. However, as apnoeic events can take place at different moments during sleep and, in particular, during REM phases [18], the whole recording must be analysed for an objective assessment of SaO₂ dynamics. To reduce the computational load, we adopted the strategy suggested in preceding studies [40]. Hence, the original SaO₂ signal was divided into epochs of length N = 512 samples to estimate AEn, SEn and KEn. As the duration of apnoeas is typically between 10 seconds
and 2 minutes, the chosen epoch length (approximately, 8.5 minutes) is large
eough to include one or more complete events. The measurements obtained
from all the epochs were then averaged to determine the final estimation for
each method.

On the other hand, parameters \( m \) and \( r \) involve statistical considerations.
Setting \( m \) large and \( r \) too small would result in inaccurate estimates of the
probabilities in \( \text{AE}_n \) and \( \text{SE}_n \) algorithms. In contrast, a large value of \( r \)
and a small \( m \) is generally too coarse to distinguish processes. Thus, \( m \) and
\( r \) were set to the widely established values suggested by Pincus [5, 41] to
obtain a statistically valid estimate of the proposed entropy measures: \( m = 1 \) or 2, and \( r = 0.1, 0.15, 0.2 \) or 0.25 times the standard deviation (SD) of
the original series.

It is worth noting that, while \( \text{KE}_n \) enables the automatic selection of
the parameter \( \sigma \), the applied MCMC-based technique requires several design
parameters to be specified. Figure 2 represents the shape of the prior \( f (\sigma) \)
for \( \lambda = 0.1, 1, 5 \) and 10. This function tends to be more concentrated near
zero as the hyperparameter \( \lambda \) becomes smaller. Zhang et al. [36] did not
find significant differences in the estimated density functions resulting from
\( \lambda \) between 0.1 and 5. Thus, we set \( \lambda = 5 \) to avoid an excessive concentration
of the probability density in a small range of \( \sigma \) values, favouring a smooth
profile of \( f (\sigma^{(m)}) \). Furthermore, the variance of the proposal distribution
for the Metropolis-Hastings algorithm must result in an acceptance rate be-
tween 20\% and 30\% of the total number of samples [36]. In our study, this
requirement was satisfied by setting that variance to 0.015. Finally, in order
to ensure the convergence of the sampling process, the number of samples to
be omitted and the number of samples to be retained were set to 5000 while
the starting $\sigma$ value was set to 1% of the SD of the original series.

3.2. Training set

All the experiments in our study were conducted on Matlab 8.2.0. Ini-
itially, we evaluated different configurations of AEn, SEn and KEn on SaO$_2$
signals in the training set to find the optimum value of the input parameters
$m$ and $r$. For each method, the configuration with the highest performance
was selected. The results achieved on the training set are summarised in
Table 2. As can be observed, $m = 1$ and $r = 0.1$SD resulted in a more ac-
curate characterisation of SAHS for AEn and SEn. For the KEn algorithm,
the two evaluated configurations provided similar results. An AUC of 0.86
and a correlation coefficient of 0.82 were reached when $m$ was set to 1. This
configuration was slightly improved by setting $m = 2$ (AUC = 0.86 and $r =
0.83$). Therefore, $m = 2$ was finally selected as the optimum for KEn.

3.3. Test set

AEn, SEn and KEn were computed on oximetry signals in the test set us-
ing the selected configurations. For each of the three methods, the Lilliefors
test [42] was applied to assess the normality of the samples. The results
showed a level of significance ($p$-value) higher than 0.1 for the distribution
of AEn, SEn and KEn in both SAHS-negative and SAHS-positive groups,
reflecting a valid assumption for normality. Subsequently, ROC and correlation analysis were conducted for each method. Table 3 summarises the obtained results. KEn showed to be the most accurate predictor of SAHS from the methods evaluated in our study. It achieved $\text{AUC} = 0.91$ and $\rho = 0.87$, which were substantially higher than the results provided by AEn ($\text{AUC} = 0.67$ and $\rho = 0.34$) and SEn ($\text{AUC} = 0.74$ and $\rho = 0.45$). The experiments reveal that a finer characterisation of $\text{SaO}_2$ irregularity was obtained by using a density estimation technique for entropy quantification as implemented by KEn.

INSERT TABLE 3 AROUND HERE

For each method, Figure 3 depicts the ROC curves and boxplots in SAHS-negative and SAHS-positive groups. The $p$-value in the figure corresponds to the level of significance for the difference between the means of each entropy metric in both groups, as obtained from the one-way ANOVA test [43]. It confirms that SEn and KEEn provided statistically significant differences between both groups of subjects. In particular, a $p$-value close to 0 was obtained for KEEn, reflecting notably more significant differences between SAHS-negative and SAHS-positive samples than AEn and SEn. Moreover, the results reflect the utility of the evaluated methods in two-class classification of patients. As can be observed, the distributions of AEn, SEn and KEEn measurements reflect that higher irregularity, i.e., higher entropy rate, is associated with oximetry signals from SAHS-positive subjects.

INSERT FIGURE 3 AROUND HERE
On the other hand, correlation analysis indicates that SaO₂ regularity is
directly related to SAHS severity, with larger entropy values obtained from
signals corresponding to subjects with a higher AHI. Figure 4 depicts the
AHI versus AEn, SEn and KEn measurements as well as the boxplots of
these measurements in each severity group. The highest Pearson’s correlation
coefficient was obtained for KEn (ρ = 0.87), which showed a marked linear
trend with respect to AHI. The boxplots obtained for the four severity groups
show a higher dispersion of AEn and SEn measurements in each category
as well as smaller differences between their median values. As a result, a
significant overlap between different groups was observed for them. This
overlapping was substantially smaller for KEn measurements, which provided
a more accurate assessment of SAHS severity.

INSERT FIGURE 4 AROUND HERE

For a more rigorous evaluation of the differences achieved by each entropy
metric between the four groups, we applied the one-way ANOVA test. The
obtained results are summarised in Table 4. As can be observed, KEn pro-
vided statistically significant differences for any pair of severity groups under
evaluation, but for the comparison between mild and moderate subjects. In
the case of SEn, significant differences were found between normal subjects
and patients with moderate SAHS, as well as between normal subjects and
patients with a severe diagnosis of SAHS. Finally, AEn only achieved signif-
icant differences when subjects in normal and severe groups were compared.
Thus, the analysis confirms the higher capability of KEn to capture differ-
ences in SaO₂ dynamics due to SAHS severity. In addition, the experiment
also shows that signals corresponding to subjects with mild and moderate SAHS tend to reflect a similar behaviour, as no significant differences was observed for any of the evaluated entropy metrics.

INSERT TABLE 4 AROUND HERE

3.4. Comparison with conventional statistical features

To complete our study, we assessed the diagnostic utility of regularity analysis of SaO2 data with respect to other statistical features commonly employed for the evaluation of biomedical signals. Conventionally, these features include the mean ($f_{avg}$), standard deviation ($f_{sd}$), coefficient of variation ($f_{cv}$), interquartile range ($f_{iqr}$) and dispersion indices ($f_{sd1}$ and $f_{sd2}$) derived from the Poincare plot. The results achieved by these features are summarised in the Table 5.

INSERT TABLE 5 AROUND HERE

As can be observed, conventional features capture relevant diagnostic information about SAHS from nocturnal oximetry recordings. In particular, they achieved significantly high AUC values, showing a good capability to discriminate between SAHS-negative and SAHS-positive subjects. Our experiments reflect that all the evaluated features but $f_{avg}$ achieved AUC higher than 0.95, which improves the AUC results provided by the three entropy metrics assessed in our research. However, correlation analysis reveals a lower ability of the conventional statistical features to detect small variations in the AHI. The obtained $\rho$ values are significantly smaller than that achieved by KEn. The latter reached $\rho = 0.87$, whereas the highest correlation coefficient among the conventional features was $\rho = 0.77$, which was provided by
Therefore, correlation results show a stronger correspondence between SAIHS severity and SaO₂ irregularity when compared to the signal properties evaluated by the proposed conventional methods.

4. Discussion and conclusions

Regularity analysis of SaO₂ recordings was performed using three different entropy algorithms: AEn, SEn and KEn. The obtained measurements show that more irregular signals are associated with SAIHS-positive subjects, reflecting the influence of apnoea events on SaO₂ dynamics. Nevertheless, there were substantial differences between the diagnostic performances of AEn, SEn and KEn. This reveals a distinct reliability of the estimators implemented by these algorithms. The latter showed to be the most consistent entropy estimator, outperforming conventional entropy algorithms like AEn and SEn. Specifically, we found that KEn measurements from oximetry data could be used to estimate the AHI of a patient \( (\rho = 0.87) \).

The KEn method represents a novel strategy for entropy estimation. The main difference between KEn and the conventional AEn and SEn algorithms is the use of probability density functions for modelling the statistical distribution of the data. According to our results, this approach has shown to be a more suitable procedure when continuous variables like SaO₂ are analysed. In addition, the use of probability density functions involves other advantages for entropy estimation. First, a data-driven method as that proposed by Zhang et al. [36] can be applied to determine the value of the kernel bandwidth. As shown in our experiments, the variance \( \sigma \) of the Gaussian kernels is optimised for the series under evaluation instead of taking an arbitrary
value fixed by the user. The automatic optimisation of \( \sigma \) suppresses one
of the user parameters in AEn and SEn algorithms, reducing by four the
number of KEn configurations to be evaluated. Second, it must be taken
into account that kernel density estimation results in the exact computation
of the integral in the definition of the Renyi entropy, as expressed in \( (15) \).
Hence, the KEn algorithm avoids one of the approximations adopted in AEn
and SEn, which corresponds to the expectation operator.

It is worth noting that the use of density functions overcomes the strong
dependency of conventional entropy metrics like AEn or SEn on the tolerance
parameter \( r \). Small values of \( r \) lead to higher and less confident entropy
estimates due to the reduced number of matches of length \( m \) and \( m + 1 \). In
the context of density estimation, the value of \( r \) is chosen in order to obtain an
accurate approximation of the target probability density function \([8, 3, 9]\).
As a result, any value of \( r \) can be used for any time series, enabling the
comparison between entropy results computed for distinct \( r \) \([9]\). As a matter
of further study, the influence of the kernel chosen for density approximation
(e.g., Gaussian, uniform, triangular or cosine kernels) on the resulting entropy
estimates should be assessed.

From our experiments, we have found that SaO\(_2\) irregularity is more
closely related to SAHS severity than conventional statistical properties in-
cluding mean, variance or dispersion indices extracted from Poincaré plots. It
is reflected by the Pearson’s correlation coefficient, which was close to 0.90
in the case of KEn. This results shows the utility of the entropy rate of
oximetry data to discriminate subjects with subtle differences between their
AHI. Nevertheless, conventional statistical features should be taken into ac-
count for the implementation of accurate methods based on oximetry data. As reported in preceding studies, multivariate analysis including several uncorrelated features can yield higher diagnostic performance than univariate measures of a given signal property [22, 24].

Several limitations can be found in our study. We have demonstrated that KEn is a valuable approach to perform regularity analysis of SaO$_2$ data in the context of SAHS diagnosis. However, the information captured by KEn is not sufficient to provide a definitive diagnosis about SAHS as reflected by AUC < 1 and $\rho < 0.9$. KEn analysis should be then considered as a tool for the interpretation of SaO$_2$ data. In this vein, the role of nocturnal pulse oximetry in SAHS diagnosis must be analysed. As in our study, the results reported by other researchers reveal suboptimal diagnostic performance of oximetry-based methods, with sensitivity and specificity lower than 100% [26, 22, 24]. Hence, the number of false negative and false positive cases would prevent the use of these methods as an alternative for PSG. Instead, home unattended pulse oximetry could be adopted as a screening tool to reduce the number of required PSG tests, contributing to minimise the waiting time for a diagnosis about SAHS. On the other hand, KEn is computationally more expensive than AEn and SEn. We have estimated that the time required to compute KEn is approximately a hundred times that of AEn and SEn. Specifically, we estimated that the time required to compute AEn, SEn and KEn on a signal epoch (512 samples) was 0.72, 0.91 and 96.78 seconds, respectively, using Matlab 8.2.0 on an Intel i7 CPU at 3.4 GHz. Mainly, this difference is due to the MCMC-based procedure for automatic adjustment of $\sigma$. This procedure requires thousands of samples to converge, with several operations.
carried out for each of these samples. In order to reduce the computational load, the value of $\sigma$ could be previously set by the user for the estimation of \textit{KEn}. Nevertheless, this does not ensure the choice of the optimum $\sigma$ for the underlying data.

In summary, our study confirms that \textit{SaO$_2$} signals from patients suffering from \textit{SAHS} tend to be more irregular. This result suggests the unpredictable occurrence of apnoeas and hypopnoeas during sleep. Hence, regularity analysis could be used for the interpretation of nocturnal oximetry dynamics in the context of \textit{SAHS} diagnosis. In particular, \textit{KEn} showed to be a reliable predictor of \textit{SAHS}. It could be considered to build new methods for automatic assessment of \textit{SAHS} severity in order to reduce the demand for conventional \textit{PSG}. Furthermore, it has been proved that \textit{KEn} is a valuable metric for regularity analysis of biomedical data, representing an alternative to conventional methods such as \textit{AEn} and \textit{SEn}.

\textbf{Acknowledgments}

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Figure captions

Figure 1: Two examples of the SaO₂ signals analysed in our study. (a) SaO₂ signal from a subject with \text{AHI} = 0.5 \text{ h}^{-1}; (b) SaO₂ signal from a subject with \text{AHI} = 32.1 \text{ h}^{-1}; (c) detailed view of the signal corresponding to the subject with \text{AHI} = 0.5 \text{ h}^{-1}; (d) detailed view of the signal corresponding to the subject with \text{AHI} = 32.1 \text{ h}^{-1}.

Figure 2: Prior probability distribution $f(\sigma)$ of the bandwidth parameter $\sigma$ for $\lambda = 0.1, 1, 5, 10$.

Figure 3: Analysis of the results on two-class classification of subjects. ROC curves computed from the measurements of (a) AEn, (c) SEn and (e) KEn. Boxplots in SAHS-negative and SAHS-positive groups for (b) AEn, (d) SEn and (f) KEn.
Figure 4: Utility of the evaluated entropy metrics to rank SAHS severity. AHI versus SaO₂ regularity quantified by (a) AEn. (c) SEn and (e) KEEn. Boxplots in the four SAHS severity groups for (b) AEn. (d) SEn and (f) KEEn.

Table captions
<table>
<thead>
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<th>SAHS-positive</th>
<th>SAHS-negative</th>
</tr>
</thead>
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<td><strong>Subjects</strong></td>
<td>96</td>
<td>64</td>
<td>32</td>
</tr>
<tr>
<td><strong>Age (years)</strong></td>
<td>52.35 ± 13.76</td>
<td>54.88 ± 14.53</td>
<td>47.31 ± 10.59</td>
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<tr>
<td><strong>Males (%)</strong></td>
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<td>84.38</td>
<td>62.50</td>
</tr>
<tr>
<td><strong>BMI (kg/m²)</strong></td>
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<td>30.61 ± 3.86</td>
<td>28.27 ± 4.38</td>
</tr>
<tr>
<td><strong>Recording Time (h)</strong></td>
<td>7.25 ± 0.33</td>
<td>7.25 ± 0.35</td>
<td>7.25 ± 0.29</td>
</tr>
<tr>
<td><strong>AHI (h⁻¹)</strong></td>
<td>24.75 ± 25.19</td>
<td>35.01 ± 25.16</td>
<td>4.23 ± 2.22</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th></th>
<th>All</th>
<th>SAHS-positive</th>
<th>SAHS-negative</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Subjects</strong></td>
<td>144</td>
<td>96</td>
<td>48</td>
</tr>
<tr>
<td><strong>Age (years)</strong></td>
<td>52.19 ± 13.73</td>
<td>54.71 ± 13.35</td>
<td>47.17 ± 13.20</td>
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<tr>
<td><strong>Males (%)</strong></td>
<td>77.78</td>
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<tr>
<td><strong>BMI (kg/m²)</strong></td>
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<td>30.98 ± 4.65</td>
<td>27.54 ± 3.26</td>
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<tr>
<td><strong>Recording Time (h)</strong></td>
<td>7.24 ± 0.66</td>
<td>7.22 ± 0.78</td>
<td>7.30 ± 0.33</td>
</tr>
<tr>
<td><strong>AHI (h⁻¹)</strong></td>
<td>26.39 ± 26.74</td>
<td>37.71 ± 26.17</td>
<td>3.75 ± 2.51</td>
</tr>
</tbody>
</table>

Table 1: Clinical and demographic features for subjects in training and test sets. Data are presented as mean ± standard deviation. BMI: body mass index; AHI: apnoea-hypopnoea index.
<table>
<thead>
<tr>
<th>Configuration</th>
<th>AUC</th>
<th>$\rho$</th>
</tr>
</thead>
<tbody>
<tr>
<td>AEn ($m = 1, r = 0.1$ SD)</td>
<td>0.68</td>
<td>0.32</td>
</tr>
<tr>
<td>AEn ($m = 1, r = 0.15$ SD)</td>
<td>0.57</td>
<td>0.19</td>
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<tr>
<td>AEn ($m = 1, r = 0.2$ SD)</td>
<td>0.50</td>
<td>0.18</td>
</tr>
<tr>
<td>AEn ($m = 1, r = 0.25$ SD)</td>
<td>0.48</td>
<td>0.15</td>
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<tr>
<td>AEn ($m = 2, r = 0.1$ SD)</td>
<td>0.56</td>
<td>0.18</td>
</tr>
<tr>
<td>AEn ($m = 2, r = 0.15$ SD)</td>
<td>0.55</td>
<td>0.13</td>
</tr>
<tr>
<td>AEn ($m = 2, r = 0.2$ SD)</td>
<td>0.51</td>
<td>0.13</td>
</tr>
<tr>
<td>AEn ($m = 2, r = 0.25$ SD)</td>
<td>0.52</td>
<td>0.14</td>
</tr>
<tr>
<td>SEn ($m = 1, r = 0.1$ SD)</td>
<td>0.74</td>
<td>0.40</td>
</tr>
<tr>
<td>SEn ($m = 1, r = 0.15$ SD)</td>
<td>0.68</td>
<td>0.31</td>
</tr>
<tr>
<td>SEn ($m = 1, r = 0.2$ SD)</td>
<td>0.63</td>
<td>0.33</td>
</tr>
<tr>
<td>SEn ($m = 1, r = 0.25$ SD)</td>
<td>0.63</td>
<td>0.34</td>
</tr>
<tr>
<td>SEn ($m = 2, r = 0.1$ SD)</td>
<td>0.67</td>
<td>0.22</td>
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<td>SEn ($m = 2, r = 0.15$ SD)</td>
<td>0.61</td>
<td>0.13</td>
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<tr>
<td>SEn ($m = 2, r = 0.2$ SD)</td>
<td>0.58</td>
<td>0.17</td>
</tr>
<tr>
<td>SEn ($m = 2, r = 0.25$ SD)</td>
<td>0.60</td>
<td>0.19</td>
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<tr>
<td>KEn ($m = 1$)</td>
<td>0.86</td>
<td>0.82</td>
</tr>
<tr>
<td>KEn ($m = 2$)</td>
<td>0.86</td>
<td>0.83</td>
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</table>

Table 2: Selection of the input parameters using training data. Results achieved on the training set by the evaluated configurations of AEn, SEn and KEn. AUC: area under the ROC curve; $\rho$: Pearson's correlation coefficient; $m$: vector length parameter; $r$: tolerance; SD: standard deviation.
<table>
<thead>
<tr>
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<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>AEn (m = 1, r = 0.1 SD)</td>
<td>&gt; 0.001</td>
<td>&gt; 0.5</td>
<td>&gt; 0.05</td>
<td>&gt; 0.005</td>
<td>&lt; 0.0001</td>
<td>&gt; 0.05</td>
</tr>
<tr>
<td>SEn (m = 1, r = 0.1 SD)</td>
<td>&gt; 0.001</td>
<td>&gt; 0.5</td>
<td>&gt; 0.01</td>
<td>&lt; 0.0001</td>
<td>&lt; 0.0001</td>
<td>&gt; 0.001</td>
</tr>
<tr>
<td>KEk (m = 2)</td>
<td>&lt; 0.0001</td>
<td>&gt; 0.001</td>
<td>&lt; 0.0001</td>
<td>&lt; 0.0001</td>
<td>&lt; 0.0001</td>
<td>&lt; 0.0001</td>
</tr>
</tbody>
</table>

Table 3: Assessment of the entropy metrics on test samples. Results achieved on the test set by the selected configurations of AEn, SEn and KEk. AUC: area under the ROC curve; $\rho$: Pearson’s correlation coefficient; $m$: vector length parameter; $r$: tolerance; SD: standard deviation.

<table>
<thead>
<tr>
<th></th>
<th>AUC</th>
<th>$\rho$</th>
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<tbody>
<tr>
<td>$f_{avg}$</td>
<td>0.85</td>
<td>0.59</td>
</tr>
<tr>
<td>$f_{sd}$</td>
<td>0.96</td>
<td>0.76</td>
</tr>
<tr>
<td>$f_{cv}$</td>
<td>0.96</td>
<td>0.71</td>
</tr>
<tr>
<td>$f_{iqr}$</td>
<td>0.96</td>
<td>0.74</td>
</tr>
<tr>
<td>$f_{sd1}$</td>
<td>0.97</td>
<td>0.77</td>
</tr>
<tr>
<td>$f_{sd2}$</td>
<td>0.96</td>
<td>0.76</td>
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Table 4: One-way ANOVA test to evaluate the difference between the means of each entropy metric in the four SAHS severity groups: normal, mild, moderate and severe.

Table 5: Diagnostic results obtained for conventional statistical features of oximetry samples on the test set. AUC: area under the ROC curve; $\rho$: Pearson’s correlation coefficient; $f_{avg}$: mean; $f_{sd}$: standard deviation; $f_{cv}$: coefficient of variation; $f_{iqr}$: interquartile range; $f_{sd1}$: first-order dispersion index from Poincare plot; $f_{sd2}$: second-order dispersion index from Poincare plot.