Testing Stationarity with Surrogates: A Time-Frequency Approach

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Abstract—An operational framework is developed for testing stationarity relatively to an observation scale, in both stochastic and deterministic contexts. The proposed method is based on a comparison between global and local time-frequency features. The originality is to make use of a family of stationary surrogates for defining the null hypothesis of stationarity and to base on them two different statistical tests. The first one makes use of suitably chosen distances between local and global spectra, whereas the second one is implemented as a one-class classifier, the time-frequency features extracted from the surrogates being interpreted as a learning set for stationarity. The principle of the method and of its two variations is presented, and some results are shown on typical models of signals that can be thought of as stationary or nonstationary, depending on the observation scale used.

Index Terms—Stationarity Test, Time-Frequency Analysis, Support Vector Machines, One-Class Classification.

I. INTRODUCTION

CONSIDERING stationarity is central in many signal processing applications, either because its assumption is a pre-requisite for applying most of standard algorithms devoted to steady-state regimes, or because its breakdown conveys specific information in evolutive contexts. Testing for stationarity is therefore an important issue, but addressing it raises some difficulties. The main reason is that the concept itself of “stationarity”, while uniquely defined in theory, is often interpreted in different ways. Indeed, whereas the standard definition of stationarity refers only to stochastic processes and concerns the invariance of statistical properties over time, stationarity is also usually invoked for deterministic signals whose spectral properties are time-invariant. Moreover, while the underlying invariances (be they stochastic or deterministic) are supposed to hold in theory for all times, common practice allows them to be restricted to some finite time interval [3], [4], [5], possibly with abrupt changes in between [6], [7]. As an example, we can think of speech that is routinely “segmented into stationary frames”, the “stationarity” of voiced segments relying in fact on periodicity structures within restricted time intervals. Those remarks call for a better framework aimed at dealing with “stationarity” in an operational sense, i.e., with a definition that would both encompass stochastic and deterministic variants, and include the possibility of its test relatively to a given observation scale.

Several attempts in this direction can be found in the literature, mostly based on concepts such as local stationarity [5]. Most of them however share the common philosophy of comparing statistics of adjacent segments, with the objective of detecting changes in the data [6], [7] and/or segmenting it over homogeneous domains [3] rather than addressing the afore-mentioned issue. Other attempts have nevertheless been made in this direction too by contrasting local properties with global ones [4], [8], but not necessarily properly phrased in terms of hypothesis testing. Among more recent approaches, we can mention those reported in [9], [10] which share some ideas with this work, but with the notable difference that they are basically model-based (whereas ours is not). Early works [11], [12] proposed a global test of stationarity based on approximate statistics of evolutionary spectra [13], which is performed as a two-step analysis of variance. The assumption of independence of time-frequency bins that are used is necessary. It is obviously and openly understood to be wrong, and may lead to stationary decision errors due to biased estimations. It is therefore the purpose of this contribution to propose and describe a different approach, using a resampling method called surrogate data, to obtain robust statistics under the hypothesis of stationarity given one observation only. It is aimed at deciding whether an observed signal can be considered as stationary, relatively to a given observation scale, and, if not, to give an index as well as a typical scale of nonstationarity.

In a nutshell, the purpose of this paper is to recast the question of stationarity testing within an operational framework elaborating on three main ideas: (i) stationarity as an operational property relying on one observation has to be understood in a relative sense including some observation scale as part of its definition; (ii) both stochastic and deterministic situations should be embraced by the approach so as to meet common practice; (iii) significance should be given to any test in order to provide users with some statistical characterization of the null hypothesis of operational stationarity.

The paper is organized as follows. In Sect. II, the general framework of the proposed approach is outlined, detailing the
time-frequency rationale of the method and motivating the use of surrogate data for characterizing the null hypothesis of stationarity and constructing stationary tests. A first test, from which both an index and a scale of nonstationarity can be derived, is proposed in Sect. III on the basis of spectral distance measures and of a parametric modeling of surrogates distributions. A second, non-parametric, test is introduced in Sect. IV by considering surrogates as a learning set and using a one-class classifier. In both cases, implementation issues are discussed, together with some examples supporting the efficiency of the methods. Finally, some of the many possible variations and extensions are briefly outlined in Sect. V.

II. Framework

Second order stationary processes are a special case of the more general class of (nonstationary) harmonizable processes, for which time-varying spectra can be properly defined [14]. When the analyzed process happens to be stationary, those time-varying spectra may reduce to the classical (stationary, time-independent) Power Spectrum Density (PSD) when suitably chosen (this holds true, e.g., for the Wigner-Ville Spectrum (WVS) [14]). In the case of more general definitions that can be considered as estimators of the WVS (e.g., spectrograms), the key point is that stationarity still implies time-independence, the time-varying spectra identifying, at each time instant, to some frequency smoothed version of the PSD. The basic idea underlying the approach proposed here is therefore that, when considered over a given duration, a process will be referred to as stationary relatively to this observation scale if its time-varying spectrum undergoes no evolution. In other words, stationarity in this acceptance happens if the local spectra at all different time instants are statistically similar to the global spectrum obtained by marginalization. This idea has already been pushed forward [4], but the novelty is to address the significance of the difference “local vs. global” by elaborating from the data itself a stationarized reference serving as the null hypothesis for the test (see Sect. II-C).

A. A time-frequency approach

As far as only second order evolutions are to be tested, quadratic time-frequency (TF) distributions and spectra are natural tools [14]. Well-established theories exist for justifying the choice of a given TF representation. In the case of stationary processes, the WVS is not only constant as a function of time but also equal to the PSD at each time instant. From a practical point of view, the WVS is a quantity that has however to be estimated. In this study, we choose to make use of multitaper spectrograms [15] defined as

\[ S_{x,K}(t, f) = \frac{1}{K} \sum_{k=1}^{K} S_{x}^{(h_k)}(t, f), \]  

where the \( \{S_{x}^{(h_k)}(t, f), k = 1, \ldots, K\} \) stand for the K spectrograms computed with the K first Hermite functions as short-time windows \( h_k(t) \):

\[ S_{x}^{(h_k)}(t, f) = \left| \int x(s) h_k(s-t) e^{-i 2 \pi f s} ds \right|^2. \]

The reason for this choice is that spectrograms can be both interpreted as estimates of the WVS for stochastic processes and as reduced interference distributions for deterministic signals [14].

In (2), the Hermite functions \( h_k(t) \) are defined by

\[ h_k(t) = ((t-D)^k g) (t)/\sqrt{\pi^{1/2} k!}, \]

with \( g(t) = \exp(-t^2/2) \). In practice, such functions can be computed recursively, according to

\[ h_k(t) = g(t) H_k(t)/\sqrt{\pi^{1/2} k!}, \]

where the \( \{H_k(t), t \in \mathbb{N}\} \) stand for the Hermite polynomials that obey the recursion

\[ H_k(t) = 2tH_{k-1}(t) - (2k - 2)H_{k-2}(t), \ k \geq 2 \]

with the initialization \( H_0(t) = 1 \) and \( H_1(t) = 2t \). Being orthonormal and maximally concentrated in TF domains with elliptic symmetry, they are a preferred family of windows for the multitaper approach which is adopted here in order to reduce estimation variance without some extra time-averaging which would be unappropriate in a nonstationary context. In practice, the multitaper spectrogram is evaluated only at N time positions \{\( t_n, n = 1, \ldots, N\} \), with a spacing \( t_{n+1} - t_n \) which is an adjustable fraction—typically, one half—of the temporal width \( T_h \) of the K windows \( h_k(t) \).

B. Relative stationarity

The TF interpretation suggesting that suitable representations should undergo no evolution in stationary situations, stationarity tests can be envisioned on the basis of some comparison between local and global features. Relaxing the assumption that stationarity would be some absolute property, the basic idea underlying the approach proposed here is that, when considered over a given duration, a process will be referred to as stationary relatively to this observation scale if its time-varying spectrum undergoes no evolution. Quantitatively, this corresponds to the fact that the local spectra \( S_{x,K}^{(n)}(t_n, f) \) at all different time instants are statistically similar to the global (average) spectrum

\[ \langle S_{x,K}(t_n, f) \rangle_n := \frac{1}{N} \sum_{n=1}^{N} S_{x,K}(t_n, f) \]  

obtained by marginalization. In practice, fluctuations in local spectra will always exist, be the signal stationary or not. The whole point of the paper is therefore to develop a comprehensive and operational methodology able to assess the significance of observed fluctuations.

C. Surrogates

Revisiting stationarity within the TF perspective has already been pushed forward [4], but the novelty is to address the significance of the difference “local vs. global” by elaborating from the data itself a stationarized reference serving as the null hypothesis for the test. Indeed, distinguishing between stationarity and nonstationarity would be made easier if, besides the analyzed signal itself, we had at our disposal some
Fourier transform known in discrete-time (surrogate signals as phase randomizations are operated. Then inverse Fourier transformed, leading to as many stationary signal while being “stationarized”. In practice, this is achieved signal, so that each of them has the same PSD as the original signal, while being “stationarized”. In practice, this is achieved by destroying the organized phase structure controlling the nonstationarity, if any. To this end, the signal is first Fourier transformed, and the magnitude of the spectrum is then kept unchanged while its phase is replaced by a random sequence, uniformly distributed over \([-\pi, \pi]\). This modified spectrum is then inverse Fourier transformed, leading to as many stationary surrogate signals as phase randomizations are operated.

To be more precise, let us assume that the observed signal is known in discrete-time \(x[n], n = 1, \ldots, N\) and has a discrete Fourier transform \(X[k]\) such that

\[
x[n] = \frac{1}{N} \sum_k X[k] e^{i2\pi nk/N}.
\]

Expressing \(X[k]\) in terms of its magnitude \(A[k]\) and phase \(\Phi[k]\) such that \(X[k] = A[k] e^{i\Phi[k]}\), a surrogate \(s[n]\) is constructed from \(X[k]\) by replacing \(\Phi[k]\) with an i.i.d. phase sequence \(\Psi[k]\) drawn from a uniform distribution over \([-\pi, \pi]\):

\[
s[n] = \frac{1}{N} \sum_k A[k] e^{i\Psi[k]} e^{i2\pi nk/N},
\]

from which it follows that

\[
s[n] s^*[m] = \frac{1}{N^2} \sum_k \sum_l A[k] A[l] e^{i(\Psi[k]-\Psi[l]+2\pi(nk+ml)/N)},
\]

Making explicit the contributions in the above double sum as:

\[
\sum_k \sum_l G[k,l] = \sum_k \left( G[k,k] + \sum_{l \neq k} G[k,l] \right)
\]

and evaluating expectations regarding random quantities, we end up for the first term with:

\[
\frac{1}{N^2} \sum_k \mathbb{E}\{A^2[k]\} e^{i2\pi(n-m)k/N}
\] (4)

and, for the second one, with

\[
\frac{1}{N^2} \sum_k \sum_{l \neq k} \mathbb{E}\{A[k]A[l]\} e^{i2\pi(nk+ml)/N} \mathbb{E}\{e^{i(\Psi[k]-\Psi[l])}\}.
\]

Given that the \(\Psi[k]\)'s are i.i.d. and uniformly distributed over \([-\pi, \pi]\), their difference has the triangular distribution:

\[
\Lambda(\psi) = \frac{1}{4\pi^2} (1 - |\psi|/2\pi)
\]

over \([-2\pi, 2\pi]\) and, therefore:

\[
\mathbb{E}\{e^{i(\Psi[k]-\Psi[l])}\} = \int_{-2\pi}^{2\pi} e^{i\psi} \Lambda(\psi) d\psi = 0.
\]

It follows that the covariance function of the surrogate \(s[n]\) reduces to (4) and, as a function of \(n - m\) only, it is therefore stationary. In the practical case where \(A[k]\) is taken as the magnitude of the Fourier transform of the observed signal (i.e., of one specific realization of the process \(x[n]\)) and kept strictly unchanged for all phase randomizations, the stationary covariance of the surrogates identifies furthermore with the Fourier transform of the (global) empirical spectrum of this observation.

The effect of the surrogate procedure is illustrated in Fig. 1, displaying both signal and surrogate spectrograms, together with their marginals in time and frequency. It clearly appears from this figure that, while the original signal undergoes a structured evolution in time, the recourse to phase randomization in the Fourier domain ends up with stationarized (i.e., time unstructured) surrogate data with identical spectrum.

As compared to more standard uses of surrogate data, we underline that we are primarily interested here in second order properties in a transformed domain. In this respect, the fact that phase randomization not only destroys nonstationarity features (as expected) but also other properties in the signal by a well-known Gaussianization effect, does not play the same dramatic role as, e.g., in testing nonlinearity in some reconstructed phase-space.

**III. A DISTANCE-BASED TEST**

Once a collection of stationarized surrogate data has been synthesized, different possibilities are offered. The first one is to extract from them some features such as *distances* between local and global spectra, and to characterize the null hypothesis

![Fig. 1. Surrogates. This figure compares the TF structure of a nonstationary FM signal (1st column), of one of its surrogates (2nd column) and of an ensemble average over \(J = 40\) surrogates (3rd column). The spectrogram is represented in each case on the 2nd line, with the corresponding marginal in time on the 3rd line. The marginal in frequency, which is the same for the three spectrograms, is displayed on the far right of the 2nd line.](https://example.com/figure1.png)
of stationarity by the statistical distribution of their variation in time. This first approach is the purpose of this Section.

A. Principle

Given an observed signal \( x(t) \) and its (multitaper) spectrogram \( S_{x,K}(t_n, f) \), it is proposed to compare local and global frequency features according to

\[
\{c_n^{(x)} := D(S_{x,K}(t_n, f), S_{x,K}(t_n, f))_n, n = 1, \ldots, N \},
\]

where \( D(\cdot, \cdot) \) stands for some dissimilarity measure (or “distance”) in frequency.

If we now label \( \{s_j(t), j = 1, \ldots, J\} \) the \( J \) surrogate signals obtained as described previously and analyze them the same way, we end up with a new collection of distances which are a function of both time indices and randomizations, namely:

\[
\{c_n^{(s_j)} := D(S_{s_j,K}(t_n, f), S_{s_j,K}(t_n, f))_n, n = 1, \ldots, N \}
\]

with \( j = 1, \ldots, J \).

As far as the intrinsic variability of surrogate data is concerned, the dispersion of distances under the null hypothesis of stationarity can be measured by the distribution of the \( J \) empirical variances

\[
\{\Theta_0(j) = \text{var}(c_n^{(s_j)})_{n=1,...,N}, j = 1, \ldots, J\}. \tag{7}
\]

This distribution allows for the determination of a threshold \( \gamma \) above which the null hypothesis is rejected. The effective test is therefore based on the statistics

\[
\Theta_1 = \text{var}(c_n^{(x)})_{n=1,...,N} \tag{8}
\]

and takes on the form of the one-sided test:

\[
d(x) = \begin{cases} 
1 & \text{if } \Theta_1 > \gamma : \text{“nonstationarity”;} \\
0 & \text{if } \Theta_1 < \gamma : \text{“stationarity”}. \tag{9}
\end{cases}
\]

The choice of the threshold \( \gamma \) will be discussed in Sect. III-E from the distribution of \( \Theta_0 \) obtained with a collection of surrogates.

Assuming that the null hypothesis of stationarity is rejected, an index of nonstationarity can furthermore be introduced as a function of the ratio between the test statistics \( (8) \) and the mean value (or the median) of its stationarized counterparts \( (7) \):

\[
\text{INS} := \sqrt{\frac{\Theta_1}{\left\langle \Theta_0(j) \right\rangle_j}} \tag{10}
\]

If the signal happens to be stationary, INS is expected to take a value close to unity whereas, the more nonstationary the signal, the larger the index.

Finally, it has to be remarked that, whereas the tested stationarity is globally relative to the time interval \( T \) over which the signal is chosen to be observed, the analysis still depends on the window length \( T_h \) of the spectrogram. Given \( T \), the index INS will therefore be a function of \( T_h \) and, repeating the test with different window lengths, we can end up with a typical scale of nonstationarity SNS defined as:

\[
\text{SNS} := \frac{1}{T} \arg \max_{T_h} \{\text{INS}(T_h)\}, \tag{11}
\]

with \( T_h \) in the range of window lengths such that the prescribed threshold is exceeded in \( (9) \).

The principle of the test having been outlined, its actual implementation depends on a number of choices that have to be made and justified, regarding distances, surrogates, thresholds, etc. Many options are however offered, that are moreover intertwined. A complete investigation of all possibilities and their combinations will not be envisioned here but, nevertheless, key features that are important for the test to be used in practice will be highlighted in the following.

B. Test signals

Setting specific parameters in the implementation is likely to end up with performance depending on the type of nonstationarity of the signal under test. Whereas no general framework can be given for encompassing all possible situations, two main classes of nonstationarities can be distinguished, which both give rise to a clear picture in the time-frequency plane: amplitude modulation (AM) and frequency modulation (FM).

We will base the following discussions on such classes. In the first case (AM), a basic, stochastic representative of the class can be modelled as:

\[
x(t) = (1 + m \sin 2\pi t/T_0) \, e(t), t \in T, \tag{12}
\]

with \( m \leq 1 \) and where \( e(t) \) is white Gaussian noise, \( T_0 \) is the period of the AM and \( T \) the observation duration. In the second case (FM), a deterministic model can be defined as:

\[
x(t) = \sin 2\pi (f_0 t + m \sin 2\pi t/T_0), t \in T, \tag{13}
\]

with \( m \leq 1 \) and where \( f_0 \) is the central frequency of the FM, \( T_0 \) its period and \( T \) the observation duration. To this FM model, a white Gaussian noise can be added if one wants to obtain different realizations of the signal.

C. Distances

Within the chosen time-frequency perspective, the proposed test \( (9) \) amounts to compare local spectra with their average over time thanks to some “distance” \( (5) \), and to decide that stationarity is to be rejected if the fluctuation of such descriptors (as given by \( (8) \)) is significantly larger than what would be obtained in a stationary case with a similar global spectrum. The choice of a distance (or dissimilarity) measure is therefore instrumental for contrasting local vs. global features.

Many approaches are available in the literature \([18]\) that, without entering into too much details, can be broadly classified in two groups. In the first one, the underlying interpretation is that of a probability density function, one of the most efficient candidate being the well-known Kullback-Leibler (KL) divergence defined as

\[
D_{KL}(G, H) := \int_{\Omega} (G(f) - H(f)) \log \frac{G(f)}{H(f)} \, df, \tag{14}
\]

where, by assumption, the two distributions \( G(\cdot) \) and \( H(\cdot) \) to be compared are positive and normalized to unity over the domain \( \Omega \). In our context, such a measure can be envisioned for (always positive) spectrograms thanks to the probabilistic
the maximum value (over \( \lambda \) with \( \lambda = 1 \)) in the AM case (right, with \( m = 0.5 \)). A better balanced performance is obtained when using the combined distance (dots) defined in (10): in the AM case, this measure performs best, and in the AM case it achieves a good contrast when \( \lambda \geq 1 \). In the AM case, the boxplots resulting from 10 realizations of the process are displayed.

A second group of approaches, which is more of a spectral nature, is aimed at comparing distributions in both shape and level. One of the simplest examples in this respect is the log-spectral deviation (LSD) defined as

\[
D_{LSD}(G, H) := \int_{\Omega} \left| \log \frac{G(f)}{H(f)} \right| df. 
\]

Intuitively, the KL measure (14) should perform poorer than the LSD (15) in the AM case (12), because of normalization. It should however behave better in the FM case (13), because of its recognized ability at discriminating distribution shapes. In order to take advantage of both measures, it is therefore proposed to combine them in some ad hoc way as

\[
D(G, H) := D_{KL}(\tilde{G}, \tilde{H}) (1 + \lambda D_{LSD}(G, H)),
\]

with \( \tilde{G} \) and \( \tilde{H} \) the normalized versions of \( G \) and \( H \), and where \( \lambda \) is a trade-off parameter to be adjusted. In practice, the choice \( \lambda = 1 \) ends up with a good performance, as justified in Fig. 2 (the performance measure used in this figure is the inverse of the maximum value (over \( T_h \)) of the index of nonstationarity \( INS \) defined in (10), i.e., an inverse measure of contrast).

### D. Distribution of Surrogates

The basic ingredient (and originality) of the approach is the use of surrogate data for creating signals whose spectrum is identical to that of the original one while being stationarized by getting rid of a well-defined structuration in time. Since those surrogates can be viewed as distinct, independent realizations of the stationary counterpart of the analyzed signal, the central part of the test is based on the statistical distribution of the \( J \) variances given in (7).

When using the combined distance suggested above in Sect. III-C, an empirical study (on both AM and FM signals) has shown that such a distribution can be fairly well approximated by a Gamma distribution. This makes sense since, according to (7), the test statistics basically sums up squared, possibly dependent quantities which themselves result from a strong mixing likely to act as a Gaussianizer. An illustration of the relevance of this modeling is given in Fig. 3, where Gamma fits are superimposed to actual histograms in the asymptotic regime (\( J = 5000 \) surrogates). Assuming the Gamma model to hold, it is possible to estimate its 2 parameters directly from the \( J \) surrogates, e.g., in a maximum likelihood sense. In this respect, Fig. 3 also supports the claim that the “theoretical” probability density function (more precisely, its estimate in the asymptotic regime) can be reasonably well approached with a reduced number of surrogates (typically, \( J \approx 50 \)). Finally, the value of the test (8), computed on the actual signals under study, is also plotted and shown to stand in the middle of the distribution in the stationary case while clearly appearing as an outlier in the considered nonstationary situation.

As a general remark, let us emphasize that modeling the distribution is important in two (related) respects. First, based on the results of Fig. 3, this allows for using much less surrogates than a crude histogram. Second, given the estimated parameters of the model, it becomes much easier to precisely define the chosen threshold for the test (see Section III.E). One could have think of other models for the distribution, such as, e.g., the simpler situation of a \( \chi^2 \). Experimental attempts in this direction proved however unsatisfactory, and the specific
analyzed signal consists of one realization of an AM process (top row), nonstationary at mesoscale (middle row) and stationary again at microscale (bottom row). The threshold (dotted line) of the stationarity test (9) is calculated with a confidence level of 95% and represented in term of INS as $\sqrt{j/(\Theta_{0}(j))}$, with $J = 50$. In the nonstationary case, the position of the maximum of INS also gives an indication of a typical scale of nonstationarity.

choice of a Gamma model was therefore guided by the fact that, with two degrees of freedom, it is obviously more flexible than a $\chi^2$, while keeping the idea of resulting from summations of squared Gaussian-like quantities.

E. Threshold and Reproduction of the Null Hypothesis

Given the Gamma model for the distribution of $\Theta_0$ based on surrogates, it becomes straightforward to derive a threshold above which the null hypothesis of stationarity is rejected with a given statistical significance. The effectiveness of the procedure at reproducing the null hypothesis of stationarity has been considered elsewhere [19], and it will not be reproduced here. Based on Monte-Carlo simulations with stationary AR processes, the main finding of the results reported in [19] is that an actual number of false positives of about 6.5 % is observed for a prescribed false alarm rate of 5 %. The test appears therefore as slightly pessimistic, yet in a reasonable agreement with what expected.

F. Illustration

In order to illustrate the proposed approach and to support its effectiveness, a simple example is given in Fig. 4. The analyzed signal consists of one realization of an AM process of the form (12). Depending on the relative values of $T_0$ and $T$, three regimes can be intuitively distinguished:

1) if $T \gg T_0$ (macroscale of observation), many oscillations are present in the observation, creating a sustained, well-established quasi-periodicity that corresponds to a form of operational stationarity;

2) if $T \approx T_0$ (mesoscale), emphasis is put on the local evolutions due to the AM, suggesting to rather consider the signal as nonstationary, with some typical scale;

3) if $T \ll T_0$ (microscale), no significant difference in amplitude is perceived, turning back to stationarity.

What is shown in Fig. 4 is that such interpretations of operational and relative stationarity are precisely evidenced by the proposed test. One difference between theoretical (second-order) stationarity of random processes and operational stationarity that we study, and that aims at being consistent with the physical and time-frequency interpretation of what stationarity means, is that both situations where there is no change in time of second-order statistics (here at microscale), and situations where there is a regular repetition if the same feature (here at macroscale of observation) are considered as being stationary in its operational acceptance. They are moreover quantified in the sense that, when the null hypothesis of stationarity is rejected (middle diagram), both an index and a scale of nonstationarity can be defined according to (10) and (11). In the present case, the maximum value of INS is obtained for $SNS = T_h/T \approx 0.2$, in qualitative accordance with the 4 AM periods entering the observation window.

At this point, it is worth stressing the fact that allowing the window length to vary is an extra degree of freedom that is part of the methodology, since it permits to give sense to a notion of stationarity relatively to an observation scale. There is therefore no prior “proper” window length for a given signal, but varying it allows for the determination of a scale of stationarity (if any).

In this specific example, the data could have been referred to as cyclostationary and analyzed by tools dedicated to such processes [20]. However, it has to be stressed that no such a priori modeling is assumed in the proposed methodology, and that the existence of a typical scale of stationarity (related to the periodic correlation) naturally emerges from the analysis.
A second example, given in Fig. 5, reports the same analysis done on a realization of the FM signal of the form (13). The same dependence on the relative values of $T_0$ and $T$ is evident and the same three possible behaviors (observation at macroscale, mesoscale or microscale of the signal) are obtained. Also, let us further comment about a specificity of operational stationarity as tested here. For the microscale $T \ll T_0$, the model turns out to be almost a pure sine function (with some added noise). The time-frequency spectrum of this signal being constant (a spectral line at frequency $f_0$), this signal is considered as stationary from an operational point of view, and relatively to an observation scale $T$ that is much larger than the period of this sine $1/f_0$ and much lower than period of the frequency modulation $T_0$. Finally, let us remark that the INS in this last case is very small as compared to the threshold: this is due to the very small variability of a sine as compared to its surrogates. This feature will be further commented when turning to the non-parametric test of the next Section.

G. Comparison to other approaches

Using surrogate data to provide users with some statistical characterization of the null hypothesis of stationarity raises the question of simply testing if the phase of the Fourier transform of $x(t)$ is an i.i.d. sequence, as is the case with surrogates. This can be performed with classical i.i.d. Portmanteau tests such as Ljung and Box, and McLeod and Li (see, e.g., in [21]). The basic property on which these tests are based is that, for large $N$, the sample autocorrelation of an i.i.d. $N$-length sequence is i.i.d. with Gaussian distribution $N(0, 1/N)$. Experiments have been conducted using a Ljung and Box statistical test with significance level of 5%, and applied to 1000 realizations of the AM and FM test signals described in Sect. III-B. The hypothesis of i.i.d. phase sequence was accepted/rejected as reported in Table I, with respective acceptance/rejection rates that were observed over the 1000 realizations. Except in the FM case where the period is equal to the observation duration, the i.i.d. hypothesis is always accepted. It clearly appears that little information is revealed by this kind of tests, which does not involve global vs. local time-frequency features, compared with the proposed framework based on second-order time-varying spectra. Nevertheless, we can mention some interesting on-going work [22] based on a Portmanteau type test for stationarity: comparing it to the present work is beyond the scope of this paper but this would certainly deserve consideration in further studies.

Existing works on statistics of evolutionary spectra [13] raise the question of constructing test statistics to compare time-frequency characteristics of the signal under study with what is expected under the null hypothesis. This kind of strategy has been adopted in many situations, e.g., to detect time-frequency features of interest [23]. In [11], [12], the authors proposed a global test of stationarity from approximate statistics. It consists of a two-step analysis of variance induced by the chi-squared distribution. As most of works in the area, the assumption of independence of time-frequency bins that are used is necessary to derive closed-form tests. This leads to poorly informative statistics and, in the present case, to stationary decision errors. Our resampling method does not rely on this hypothesis, but considers the spectrograms of surrogate data globally to provide more informative tests.

IV. A NON-SUPERVISED CLASSIFICATION APPROACH

Besides the distance-based approach described above, we can adopt an alternative viewpoint rooted in statistical learning theory by considering the collection of surrogates as a learning set and using it to estimate the support of the distribution of stationarized data. Let us make this approach more precise.

A. An overview on one-class classification

In the context considered here, the classification task is fundamentally a one-class classification problem and differs from conventional two-class pattern recognition problems in the way how the classifier is trained. The latter uses only target data to perform outlier detection. This is often accomplished by estimating the probability density function of the target data, e.g., using a Parzen density estimator [24]. Density estimation methods however require huge amounts of data, especially in high dimensional spaces, which makes their use impractical. Boundary-based approaches attempt to estimate the quantile function defined by $Q(\alpha) := \inf\{\lambda(C) : P(C) := \int_{\omega \in C} h(\omega) \geq \alpha\} \leq 0 \leq \alpha \leq 1$, where $C$ denotes a subset of the signal space $S$ that is measurable with respect to the probability measure $\mu$, and $\lambda(C)$ its volume. Estimators $C_{\alpha}$ that reach this infimum, in the case where $P$ is the empirical distribution, are called minimum volume estimators. The first boundary-based approach was probably introduced in [25], where the authors consider a class of closed convex boundaries in $\mathbb{R}^2$. More sophisticated methods were described in [26], [27]. Nevertheless, they are based upon neural networks training and therefore suffer from the same drawbacks such as slow convergence and local minima. Inspired by support vector machines, the support vector data description algorithm proposed in [28] enforces target data in a minimum volume hypersphere. More flexible boundaries can be obtained by using kernel functions, that map the data into a high-dimensional feature space. In the case of normalized kernel functions, this approach is equivalent to the one-class support vector machines introduced in [29], which use a maximum margin hyperplane to separate data from the origin. The generalization performance of these algorithms were investigated in [29], [30], [31] via the derivation of

<table>
<thead>
<tr>
<th>Period</th>
<th>AM</th>
<th>FM</th>
</tr>
</thead>
<tbody>
<tr>
<td>$T_0 = T/20$</td>
<td>accepted (94%)</td>
<td>accepted (94%)</td>
</tr>
<tr>
<td>$T_0 = T$</td>
<td>accepted (92%)</td>
<td>rejected (54%)</td>
</tr>
<tr>
<td>$T_0 = 20T$</td>
<td>accepted (94%)</td>
<td>accepted (90%)</td>
</tr>
</tbody>
</table>

**TABLE I**

RESULT OF A PORTMANTEAU TEST ON THE PHASE OF THE FOURIER TRANSFORM OF $x(t)$, FOR THE AM AND FM PROCESSES OF SECT. III-B.

THE PARAMETERS ARE $N = T = 1600$, SNR = 100B, $m = 0.5$ (AM) AND 0.02 (FM). THE RESULTS REPORTED ARE THE AVERAGE OF 1000 REALIZATIONS.
bounds. In what follows, we shall focus on the support vector data description algorithm.

B. Support vector data description

Let us assume that we are given a training set \( \{z_1, \ldots, z_J\} \) (this may correspond either to the surrogates themselves or to some features derived from them). The center of the smallest enclosing hypersphere is the point \( a^* \) that minimizes the distance from the furthest training data, namely,

\[
a^* = \text{arg min}_a \max_{i=1,\ldots,J} \|z_i - a\|.
\]

We observe that the solution of this problem is highly sensitive to the location of just one point, which may result in a pattern analysis system that is not robust. This suggests us to consider hyperspheres that balance the loss incurred by missing a small fraction of target data with the reduction in radius that results. The following optimization problem implements this strategy

\[
\min_{a, r, \xi} \quad r^2 + \sum_{i=1}^J \xi_i / \nu J
\]

subject to

\[
\|z_i - a\|^2 \leq r^2 + \xi_i, \quad \xi_i \geq 0, \quad i = 1, \ldots, J
\]

with a parameter \( \nu \) in \([0, 1]\) to control the trade-off between minimizing the radius and controlling the slack variables defined as \( \xi_i = (\|a - z_i\|^2 - r^2)_+ \), see Fig. 6. The exact role of \( \nu \) will be discussed next.

We can solve this constrained optimization problem by defining a Lagrangian involving one Lagrange multiplier for each constraint:

\[
L(a, r, \xi; \alpha, \beta) = r^2 + \sum_{i=1}^J \xi_i / \nu J \]

\[
+ \sum_{i=1}^J \alpha_i (\|z_i - a\|^2 - r^2 - \xi_i) - \sum_{i=1}^J \beta_i \xi_i,
\]

with \( \alpha_i, \beta_i \geq 0 \). Setting to zero its derivatives with respect to the primal variables \( a, r \) and \( \xi \) gives

\[
a^* = \sum_{i=1}^J \alpha_i^* z_i
\]

(19)

where \( a^* \) is the solution of the optimization problem

\[
\max_{\alpha} \quad \sum_{i=1}^J \alpha_i (\|z_i - a\|^2 - \sum_{i,j=1}^J \alpha_i \alpha_j (z_i, z_j)) \]

subject to

\[
\sum_{i=1}^J \alpha_i = 1, \quad 0 \leq \alpha_i \leq 1 / \nu J, \quad i = 1, \ldots, J
\]

(20)

Depending on whether a training data \( z_i \) lies inside the hypersphere, on or outside, it can be shown that its Lagrange multiplier \( \alpha_i^* \) in (19) satisfies one of the three conditions: \( \alpha_i^* = 0 \) (inside), \( 0 < \alpha_i^* < 1 / \nu J \) (on), and \( \alpha_i^* = 1 / \nu J \) (outside). With condition \( \sum_{i=1}^J \alpha_i^* = 1 \) in (20), we know that there can be at most \( \nu J \) training points lying outside the hypersphere. Furthermore, with the upper bound \( 1 / \nu J \) on \( \alpha_i^* \), we observe that at least \( \nu J \) training data do not lie inside the hypersphere.

After discussing the role of the parameter \( \nu \), which allows the user some control over the fraction of points that are excluded from the hypersphere, we shall now derive the decision rule for novelty detection. The distance of any point \( x \) from the center \( a^* \) of the hypersphere can be shown to be

\[
\|x - a^*\| = \sqrt{\langle x, x \rangle - 2 \sum_{i=1}^J \alpha_i^* \langle x, z_i \rangle + \sum_{i,j=1}^J \alpha_i^* \alpha_j^* (z_i, z_j)}
\]

(21)

Equation (21) can be used to calculate the radius \( r^* \) of the optimal hypersphere, that is, \( r^* := \|z_{i_0} - a^*\| \) with \( i_0 \) the index of any training data such that \( 0 < \alpha_{i_0}^* < 1 / \nu J \). This allows us to compute the resulting slack values \( \xi_i \), see the definition below equation (17). Consider now the test statistics

\[
\Theta(x) := \|x - a^*\|^2 - (r^*)^2
\]

such that the decision function

\[
d(x) := \left\{ \begin{array}{ll}
1 & \text{if } \Theta(x) > \gamma : \text{"nonstationarity"}; \\
0 & \text{if } \Theta(x) < \gamma : \text{"stationarity"}
\end{array} \right.
\]

(22)

outputs 1 if the test point \( x \) lies outside the hypersphere of squared radius \( (r^*)^2 + \gamma \) and so is considered novel, and 0 otherwise. The threshold parameter \( \gamma \) has a direct influence upon the performance of the novelty detector. With probability greater than \( 1 - \delta \), we can bound the probability that function \( d \) outputs 1 on a test point drawn according to the original distribution by

\[
\frac{1}{\gamma J} \sum_{i=1}^J \xi_i + \frac{6R^2}{\gamma \sqrt{J}} + 3 \frac{\ln(2/\delta)}{2J}
\]

(23)

where \( R \) is the radius of a ball in feature space centered at the origin containing the support of the distribution. Such examples are false positives in the sense that they are normal data identified as novel by the decision function \( d \). Obviously, we cannot bound the rate of true negatives since we have no way of guaranteeing what output \( d \) returns for data drawn from a different distribution. The strategy underlying this approach is however the smaller the hypersphere, the more likely novel data will fall outside and be identified as outlier.

When a more sophisticated model than the hypersphere is required to accurately fit the quantile functions, a possible strategy is to replace each inner product \( \langle z_i, z_j \rangle \) in equations (20)–(22) by a kernel function \( \kappa(z_i, z_j) := \langle \phi(z_i), \phi(z_j) \rangle \). An ideal kernel would implicitly map the data via \( \phi(\cdot) \) into a bounded spherically-shaped area of a new feature space. Kernel functions can usually be computed more efficiently as a direct function of the input data, without explicitly evaluating the mapping \( \phi(\cdot) \). Classic examples of kernels are the radially Gaussian kernel \( \kappa(z_i, z_j) = \exp(-\|z_i - z_j\|^2 / 2\sigma_0^2) \), and the Laplacian kernel \( \kappa(z_i, z_j) = \exp(-\|z_i - z_j\| / \sigma_0) \), with \( \sigma_0 \geq 0 \) the kernel bandwidth.

Another example which deserves attention in signal processing is the \( q \)-th degree polynomial kernel defined as \( \kappa(z_i, z_j) = (1 + \langle z_i, z_j \rangle)^q \), with \( q \in \mathbb{N}^* \). In [32], we have extended the framework of the so-called kernel-based methods to time-frequency analysis, showing that some specific reproducing kernels allow these algorithms to operate in the time-frequency domain. This link offers new perspectives in the field of non-stationary signal analysis, which can benefit from the developments of pattern recognition and statistical learning theory.
C. Testing stationarity

We shall now use support vector data description to estimate the support of probability density functions of stationary surrogate signals. The resulting decision rule will allow us to distinguish between stationary and nonstationary processes.

Let us assume that we are given a training set \( \{s_1(t), \ldots, s_J(t)\} \) of surrogate signals generated from the signal \( x(t) \) under investigation. In all the experiments reported above, time-frequency features were extracted from the normalized multitaper spectrogram of each signal, defined at time \( t_n \) by

\[
S_n(f) := \frac{S_{x,K}(t_n, f)}{\sum_{n=1}^N S_{x,K}(t_n, f)} df
\]  

(24)

for \( n = 1, \ldots, N \) and \( 0 \leq f < 1/2 \). More precisely, the local power \( P_n \) of each signal and its local frequency content \( F_n \) summarized below were considered:

\[
P_n := \int_0^{1/2} S_n(f) df ; \quad F_n := \frac{1}{P_n} \int_0^{1/2} f S_n(f) df.
\]  

(25)

Finally, for a sake of clarity, only the following two features comparing local time-frequency behavior to global one were retained

\[
P := \text{std}(P_n)_{n=1,\ldots,N} ; \quad F := \text{std}(F_n)_{n=1,\ldots,N},
\]  

(26)

where \( \text{std}(\cdot) \) denotes the standard deviation. The first one is a measure of the fluctuations over time of the local power of the signal, whereas the second one operates the same way with respect to the local mean frequency. For each experiment reported in Fig. 7, a training set consisting of 200 surrogate signals was generated from the AM or FM signal \( x(t) \) to be tested. Features \( P \) and \( F \) were extracted from each signal. Next, data were mean-centered and normalized so that the variance of both features was one. Finally, the support vector data description algorithm was run using the basic linear kernel \( \kappa(z_i, z_j) = \langle z_i, z_j \rangle \) and \( \nu = 0.15 \). The results are displayed for \( T_0 = T/20, T \) and \( 20T \), allowing to consider stationarity relatively to the ratio between the observation time \( T \) and the modulation period \( T_0 \). In each figure, the surrogate signals are shown with dots and the signal to be tested with a black triangle. The optimum circle having center at \( e^* \) and radius \( r^* \) is shown in dashed line. The training data lying on or outside this circle, and thus associated with non-zero Lagrange multipliers in (19)-(21), are indicated by the circled dots. The thin circles represent the decision function (22) tuned to different false positive probabilities, fixed by \( \gamma \) via the relation (23). To calculate \( \gamma \), note that we have neglected the contribution of the last two terms of equation (23) since they decay to zero as \( J \) tends to infinity. Figs. 7(b) and 7(e) show that the test signals can be considered as nonstationary with a false positive probability lower than 0.05. In the other figures, they are clearly identified as stationary signals.

The findings reported in this learning-theory-based study are clearly consistent with what had been obtained previously with the distance-based approach. For a small modulation period or a large observation time, i.e., when \( T_0 \ll T \), the situation can be considered as stationary due to the observation of many similar oscillations over the observed time scale. This is reflected by the test signal which lies inside the region defined by the support vector data description algorithm for the stationary surrogates. For a medium observation time, i.e., \( T \approx T_0 \), the local evolution due to the modulation is prominent and the black triangle for the modulated signal is well outside the stationary region, in accordance with a situation that can be referred to as nonstationary. Finally, if \( T_0 \gg T \), the result turns back to stationarity in the AM case because no significative change in the amplitude is observed over the considered time scale. In the FM case, the situation is slightly different, with the black triangle lying at the border of the surrogates domain. For the FM case, especially Figs. 7(e) and 7(f), \( P \) is negative and lower than the values of \( P \) taken by the surrogates: this is characteristic of some regularity and constancy in the amplitude, which has less fluctuations than any corresponding stationary random process. Indeed, the way stationarity is tested cannot end up with a better configuration in this case since, by construction, surrogates of finite length pure tones undergo necessarily some amplitude fluctuations leading to a negative, non-zero \( P \) index for the (non-fluctuating) test signal. This can be viewed as a limitation of the method but it should rather be interpreted as a known bias when the test signal lies at the border of the surrogates domain and comes along with a value \( F \) close to 0. It could be used as an indication to discriminate random processes from more deterministic ones.

Interestingly, one can also remark that the location of the test signal in the \( (P, F) \) plane turns out to provide some information about the type of nonstationarity, if any: \( F \neq 0 \) is characteristic of some FM structure, \( P > 0 \) indicates some AM, \( P < 0 \) is associated to a constant (maybe deterministic) behavior for the amplitude (see [33] for preliminary results in this direction).

V. Conclusion

A new approach has been proposed for testing stationarity from a time-frequency viewpoint, relatively to a given observation scale. A key point of the method is that the null hypothesis
of stationarity (which corresponds to time-invariance in the time-frequency spectrum) is statistically characterized on the basis of a set of surrogates which all share the same average spectrum as the analyzed signal while being stationarized. Two possible ways of making use of surrogates have been discussed, based either on a distance-based approach or on a machine learning technique (one class-SVM). Both ways are complementary since the first one is parametric whereas the second one is not. As is usual, a parametric approach requires such assumptions, but it is more demanding in terms of data size and computation. Moreover, for the distance-based approach, some analytical studies can be conducted (possibly in asymptotic situations, see [34]), whereas the SVM approach allows for more versatile characterization of types of non-stationarity. As for comparing to early works using asymptotic analysis of evolutionary spectra [11, 12] to formulate a global test of stationarity, the assumption of independence between the time-frequency bins requires that many data in the time-frequency representations have to be discarded for the test. Our resampling method with surrogates allow for lightening this limitation and using all the bins, hence providing a method that uses all available information.

The basic principles of the method have been outlined, with a number of considerations related to its implementation, but it is clear that the proposed framework still leaves room for more thorough investigations as well as variations and/or extensions. In terms of time-frequency distributions for instance, one could imagine to go beyond spectrograms and take advantage of more recent advances [35]. Two-dimensional extensions can also be envisioned for testing stationarity in the sense of homogeneity of random fields, e.g., for texture analysis. Preliminary results in this direction are given in [36].

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Fig. 7. Time-frequency features \((P, F)\) in AM (left) and FM (right) situations. From top to bottom, \(\frac{\lambda_0}{\lambda} = T/20, T\) and \(20T\), with \(T = 1600\). In each case, the black triangle corresponds to the \((P, F)\) pair of one test signal used to derive the surrogates. The latter are plotted as dots which, with support vector data description, define the minimum-volume domain of stationarity represented here by the dashed curve. The training data lying on or outside this area are indicated by the circled dots. The thin curves represent the decision functions with false positive probabilities 0.05, 0.1 and 0.15. Other parameters are as follows — number of tapers: \(K = 5\), length of tapers: \(T_h = 387\), modulation indices: (AM) \(m = 0.5\) and (FM) \(m = 0.02\), central frequency \(f_0 = 0.25\), signal-to-noise ratio: \(\text{SNR} = 10\) dB.