Detecting changes in the mean of functional observations

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Abstract

Principal component analysis (PCA) has become a fundamental tool of functional data analysis. It represents the functional data as $X_i(t) = \mu(t) + \sum_{1 \leq \ell < \infty} \eta_{i,\ell} v_{\ell}(t)$, where $\mu$ is the common mean, $v_{\ell}$ are the eigenfunctions of the covariance operator, and the $\eta_{i,\ell}$ are the scores. Inferential procedures assume that the mean function $\mu(t)$ is the same for all values of $i$. If, in fact, the observations do not come from one population, but rather their mean changes at some point(s), the results of PCA are confounded by the change(s). It is therefore important to develop a methodology to test the assumption of a common functional mean. We develop such a test using quantities which can be readily computed in the R package fda. The null distribution of the test statistic is asymptotically pivotal with a well-known asymptotic distribution. The asymptotic test has excellent finite sample performance. Its application is illustrated on temperature data from Prague, England and Greenland.

Keywords: Change point detection; Functional data analysis; Mean of functional data; Significance test

Abbreviated Title: Changes in functional mean

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Introduction

Functional data analysis (FDA) has been enjoying increased popularity over the last decade due to its applicability to problems which are difficult to cast into a framework of scalar or vector observations. Even if such standard approaches are available, the functional approach often leads to a more natural and parsimonious description of the data, and to more accurate inference and prediction results. The monograph of Ramsay and Silverman (2005) has become a standard reference to the ideas and tools of FDA. To name a few recent applications of FDA which illustrate its advantages alluded to above, we cite Antoniadis and Sapatinas (2003), Fernández de Castro et al. (2005), Müller and Stadtmüller (2005), Yao et al. (2005), Glendinning and Fleet (2007).

A main tool of FDA is principal component analysis (PCA). It represents the functional observations $X_i(t), t \in T, i = 1, 2, \ldots, N$, in the form $X_i(t) = \mu(t) + \sum_{1 \leq \ell < \infty} \eta_{i,\ell} v_\ell(t)$, where $\mu$ is the mean, $v_\ell$ are the eigenfunctions of the covariance operator, and the $\eta_{i,\ell}$ are the scores. The set $T$ can be interpreted as a time or a spatial domain, the methodology we develop requiring merely that it be a compact subset of a Euclidean space. To perform the functional PCA, the functional mean $\mu(t)$, approximated by the sample mean of the $X_i(t)$, is first subtracted from the data. The first principal component $v_1(t)$ is then interpreted as the main pattern of deviation of the observations from the mean $\mu(t)$, or equivalently, as the direction in a function space of the largest variability away from the mean function. The subsequent eigenfunction define analogous directions orthogonal to the previous eigenfunctions. This interpretation and the inferential procedures based on it assume that the mean function $\mu(t)$ is the same for all values of $i$. If, in fact, the mean changes at some index(es) $i$, the results of PCA are confounded by the change(s). Issues of this type are most likely to emerge if the data are collected sequentially over time. Applications we have in mind abound in climatology, environmental science and economics; detecting and locating changes in mean can be interpreted, for example, as climate shifts, a baseline change in a pollution level, or a shift in a long-term rate of growth.

It is thus useful to develop a methodology for the detection of changes in the mean of functional observations that is both easy to apply and justified by a clear large sample argument. We propose a significance test for testing the null hypothesis of a constant functional mean against the alternative of a changing mean. We also show how to locate the change points if the null hypothesis is rejected. Our methodology is readily implemented using the R package fda, R Development Core Team (2008). The null distribution of the test statistic is asymptotically pivotal with a well-known asymptotic distribution going back to the work of Kiefer (1959).

The problem of detecting a change in the mean of a sequence Banach–space valued random elements has recently been approached from a theoretical angle by Rackauskas
and Suquet (2006). Motivated by detecting an epidemic change (the mean changes and then returns to its original value), Rackauskas and Suquet (2006) proposed an interesting statistic based on increasingly fine dyadic partitions of the index interval, and derived its limit, which is nonstandard.

The change point problem has been extensively studied in the multivariate setting starting with Srivastava and Worsley (1986), while the work of Horváth et al. (1999) is most closely related to the present paper. Different multivariate settings with further references are discussed in Lavielle and Teyssiére (2006), Zamba and Hawkins (2006) and Qu and Perron (2007), among others.

Returning to the functional setting, a somewhat related problem has recently been studied by Benko et al. (2009) who considered two populations, admitting the PCA’s:

\[ X_{i,p}(t) = \mu_p(t) + \sum_{1 \leq \ell < \infty} \eta_{i,p,\ell} v_{p,\ell}(t), \quad p = 1, 2. \]

Benko et al. (2009) developed a bootstrap test for checking if the elements of the two decompositions (including the means) are the same. Earlier, Laukaitis and Račkauskas (2005) considered the model

\[ X_{i,g}(t) = \mu_g(t) + \varepsilon_{i,g}(t), \quad g = 1, 2, \ldots, G, \]

with innovations \( \varepsilon_{i,g} \) and group means \( \mu_g \), and tested \( H_0 : \mu_1(t) = \ldots = \mu_G(t) \). Other contributions in this direction include Cuevas et al. (2004), Delicado (2007) and Ferraty et al. (2007). In these settings, it is known which population or group each observation belongs to. In our setting, we do not have any partition of the data into several sets with possibly different means. The change can occur at any point, and we want to test if it occurs or not.

The paper is organized as follows. In Section 2, we introduce the required notation and assumptions, and recall several results which will be used in the following sections. Section 3 describes the proposed methodology and contains theorems which provide its asymptotic justification. The finite sample performance is investigated in Section 4, which also contains an illustrative application to the detection of changes in mean patterns of annual temperatures. The proofs of the Theorems of Section 3 are presented in the Appendix.

2 Notation and assumptions

We consider functional observations \( X_i(t), \ t \in \mathcal{T}, \ i = 1, 2, \ldots, N \), defined over a compact set \( \mathcal{T} \). We assume that the \( X_i \) are independent, and we want to test if their mean remains constant in \( i \). Thus we test the null hypothesis

\[ H_0 : \ E X_1(t) = E X_2(t) = \ldots = E X_N(t), \ t \in \mathcal{T}. \]
Note that under $H_0$, we do not specify the value of the common mean.

Under the alternative, $H_0$ does not hold. The test we construct has particularly good power against the alternative in which the data can be divided into several consecutive segments, and the mean is constant within each segment, but changes from segment to segment. The simplest case of only two segments (one change point) is specified in Assumption 2.4.

Under the null hypothesis, we can represent each functional observation as

$$X_i(t) = \mu(t) + Y_i(t), \quad EY_i(t) = 0.$$  

(2.1)

The following assumption specifies conditions on $\mu(\cdot)$ and the errors $Y_i(\cdot)$ needed to establish the asymptotic distribution of the test statistic.

In the following, unless indicated otherwise, all integrals denote integration over the set $T$.

**Assumption 2.1** The mean $\mu(\cdot)$ is in $L^2(T)$. The errors $Y_i(\cdot)$ are iid mean zero random elements of $L^2(T)$ which satisfy

$$E||Y_i||^2 = \int EY_i^2(t)dt < \infty.$$  

(2.2)

Their covariance function

$$c(t, s) = E[Y_i(t)Y_j(s)], \quad t, s \in T$$  

(2.3)

is square integrable, i.e. is in $L^2(T \times T)$.  

Assumption 2.1 implies the following expansions, see e.g. Chapter 4 of Indritz (1963):

$$c(t, s) = \sum_{1 \leq k < \infty} \lambda_k v_k(t)v_k(s)$$  

(2.4)

and

$$Y_i(t) = \sum_{1 \leq \ell < \infty} \lambda_\ell^{1/2} \xi_i,\ell v_\ell(t),$$  

(2.5)

where $\lambda_k$ and $v_k$ are the eigenvalues and eigenfunctions, respectively, of the covariance operator, i.e. they are defined by

$$\int c(t, s)v_\ell(s)ds = \lambda_\ell v_\ell(t), \quad \ell = 1, 2, \ldots$$  

(2.6)

The sequences $\{\xi_i,\ell, \ell = 1, 2, \ldots\}$ are independent, and within each sequence the $\xi_i,\ell$ are uncorrelated with mean zero and unit variance. The infinite sum in (2.5) converges in $L^2(T)$ with probability one. Recall also that $v_\ell, \ell = 1, 2, \ldots,$ form an orthonormal basis in $L^2(T)$, and all $\lambda_\ell$ are nonnegative.
In practice, we work with estimated eigenelements defined by
\[
\int \hat{c}(t, s) \hat{v}_\ell(s) ds = \hat{\lambda}_\ell \hat{v}_\ell(t), \quad \ell = 1, 2, \ldots,
\]
where
\[
\hat{c}(t, s) = \frac{1}{N} \sum_{1 \leq i \leq N} \left( X_i(t) - \bar{X}_N(t) \right) \left( X_i(s) - \bar{X}_N(s) \right)
\]
and \( \bar{X}_N(t) = \frac{1}{N} \sum_{1 \leq i \leq N} X_i(t) \).

To control the distance between the estimated and the population eigenelements, we need the following assumptions:

**Assumption 2.2** The eigenvalues \( \lambda_\ell \) satisfy, for some \( d > 0 \)
\[
\lambda_1 > \lambda_2 > \ldots > \lambda_d > \lambda_{d+1}.
\]

**Assumption 2.3** The \( Y_i \) in (2.1) satisfy
\[
\mathbb{E}[||Y_i||^4] = \int \mathbb{E}Y_i^4(t) dt < \infty.
\]

The results of Dauxois et al. (1982) and Bosq (2000) then imply that for each \( k \leq d \):
\[
\limsup_{N \to \infty} N \mathbb{E} \left[ ||\hat{c}_k v_k - \hat{v}_k||^2 \right] < \infty, \quad \limsup_{N \to \infty} N \mathbb{E} \left[ ||\lambda_k - \hat{\lambda}_k||^2 \right] < \infty,
\]
where \( \hat{c}_k = \text{sign} \int_T v_k(t) \hat{v}_k(t) dt \). The random sign \( \hat{c}_k \) is included because the \( v_k \) and \( \hat{v}_k \) are defined up to a sign, and since \( v_k \) is unknown, it is impossible to ensure that \( \int_T v_k(t) \hat{v}_k(t) dt \geq 0 \).

We establish the consistency of the test under the alternative of one change point formalized in Assumption 2.4. A similar argument can be developed if there are several change points, but the technical complications then obscure the main idea explained in Sections 3 and A.2 (in particular the functions (2.10) and (3.7) would need to be modified). The more general case is studied empirically in Section 4.

**Assumption 2.4** The observations follow the model
\[
X_i(t) = \begin{cases} 
\mu_1(t) + Y_i(t), & 1 \leq i \leq k^*, \\
\mu_2(t) + Y_i(t), & k^* < i \leq N,
\end{cases}
\]
in which the \( Y_i \) satisfy Assumption 2.1, the mean functions \( \mu_1 \) and \( \mu_2 \) are in \( L^2(T) \), and
\[
k^* = \lfloor n\theta \rfloor \quad \text{for some} \quad 0 < \theta < 1.
\]
We will see in the proof of Theorem 3.2 that under Assumption 2.4 the sample covariances of the functional observations converge to the function

\begin{equation}
\bar{c}(t, s) = c(t, s) + \theta(1 - \theta)(\mu_1(t) - \mu_2(t))(\mu_1(s) - \mu_2(s)).
\end{equation}

This is a symmetric, square integrable function, and it is easy to see that for any \(x, y \in L^2(\mathcal{T})\),

\[
\int \int \bar{c}(t, s)x(t)x(s)dtds \geq 0,
\]

so \(\bar{c}(t, s)\) is a covariance function. Consequently, it has orthonormal eigenfunctions \(w_k\) and nonnegative eigenvalues \(\gamma_k\) satisfying

\begin{equation}
\int \bar{c}(t, s)w_k(s)ds = \gamma_kw_k(t).
\end{equation}

The quantities \(\bar{c}(t, s), w_k\) and \(\gamma_k\) are used in Section 3 to describe the distribution of the test statistic under the alternative of a single change point.

## 3 Detection procedure

To explain the idea of the test procedure, denote

\[
\hat{\mu}_k(t) = \frac{1}{k} \sum_{1 \leq i \leq k} X_i(t), \quad \overline{\mu}_k(t) = \frac{1}{N-k} \sum_{k < i \leq N} X_i(t).
\]

If the mean is constant, the difference \(\Delta_k(t) = \hat{\mu}_k(t) - \overline{\mu}_k(t)\) is small for all \(1 \leq k < N\) and all \(t \in \mathcal{T}\). However, \(\Delta_k(t)\) can become large due to chance variability if \(k\) is close to \(1\) or to \(N\). It is therefore usual to work with the sequence

\begin{equation}
P_k(t) = \sum_{1 \leq i \leq k} X_i(t) - \frac{k}{N} \sum_{1 \leq i \leq N} X_i(t) = \frac{k(N-k)}{N}[\hat{\mu}_k(t) - \overline{\mu}_k(t)]
\end{equation}

in which the variability at the end points is attenuated by a parabolic weight function. If the mean changes, the difference \(P_k(t)\) is large for some values of \(k\) and of \(t\). Since the observations are in an infinite dimensional domain, we work with the projections of the functions \(P_k(\cdot)\) on the principal components of the data. These projections can be expressed in terms of functional scores which can be easily computed using the \texttt{R} package \texttt{fda}.

Consider thus the scores corresponding to the largest \(d\) eigenvalues:

\[
\hat{\eta}_{i, \ell} = \int [X_i(t) - \bar{X}(t)]\hat{v}_\ell(t)dt, \quad i = 1, 2, \ldots, N, \quad \ell = 1, 2, \ldots, d.
\]
Observe that the value of $P_k(t)$ does not change if the $X_i(t)$ are replaced by $X_i(t) - \bar{X}_N(t)$.

Consequently, setting $\ell = \lfloor Nx \rfloor$, $x \in (0, 1)$, we obtain

$$\int \left\{ \sum_{1 \leq i \leq Nx} X_i(t) - \frac{N x}{N} \sum_{1 \leq i \leq N} X_i(t) \right\} \hat{v}_\ell(t) \, dt = \sum_{1 \leq i \leq N x} \hat{h}_{i, \ell} - \frac{N x}{N} \sum_{1 \leq i \leq N} \hat{h}_{i, \ell}. \tag{3.2}$$

Identity (3.2) shows that functional scores can be used for testing the constancy of the mean function.

The following theorem can be used to derive a number of test statistics. To state it, introduce the statistic

$$T_N(x) = \frac{1}{N} \sum_{\ell=1}^d \hat{\lambda}_\ell^{-1} \left( \sum_{1 \leq i \leq Nx} \hat{h}_{i, \ell} - x \sum_{1 \leq i \leq N} \hat{h}_{i, \ell} \right)^2 \tag{3.3}$$

and let $B_1(\cdot), \ldots, B_d(\cdot)$ denote independent standard Brownian bridges.

**Theorem 3.1** Suppose Assumptions 2.1, 2.2,2.3 hold. Then, under $H_0$,

$$T_N(x) \xrightarrow{d} \sum_{1 \leq \ell \leq d} B^2_\ell(x) \quad (0 \leq x \leq 1),$$

in the Skorokhod topology of $D[0,1]$.

Theorem 3.1 is proved in the Appendix.

By Theorem 3.1, $U(T_N) \xrightarrow{d} U(\sum_{1 \leq \ell \leq d} B^2_\ell(\cdot))$, for any continuous functional $U : D[0,1] \to R$. Applying integral or max functionals, or their weighted versions, leads to useful statistics. In this paper, we focus on the integral of the squared function, i.e. the Cramer-von-Mises functional, which is known to produce effective tests (this functional was also selected in a different context by Bugni et al. (2006)). Thus, we consider the convergence

$$\int_0^1 T_N(x) \, dx \xrightarrow{d} \int_0^1 \sum_{1 \leq \ell \leq d} B^2_\ell(x) \, dx,$$

which can be rewritten as

$$S_{N,d} := \frac{1}{N^2} \sum_{\ell=1}^d \hat{\lambda}_\ell^{-1} \sum_{k=1}^N \left( \sum_{1 \leq i \leq k} \hat{h}_{i, \ell} - \frac{k}{N} \sum_{1 \leq i \leq N} \hat{h}_{i, \ell} \right)^2 \xrightarrow{d} \int_0^1 \sum_{1 \leq \ell \leq d} B^2_\ell(x) \, dx. \tag{3.4}$$

The distribution of the random variable

$$K_d = \int_0^1 \sum_{1 \leq \ell \leq d} B^2_\ell(x) \, dx \tag{3.5}$$

was derived by Kiefer (1959). Denoting by $c_d(\alpha)$ its $(1 - \alpha)$th quantile, the test rejects $H_0$ if $S_{N,d} > c_d(\alpha)$. The critical values $c_d(\alpha)$ are presented in Table 4.1.

A multivariate analog of statistic (3.4) considered in Horváth et al. (1999) is

$$M_{N,d} = \frac{1}{N^2} \sum_{k=1}^N \left( \frac{k}{N} - \frac{k}{N} \right)^2 \Delta(k) \tilde{D}_d^{-1} \Delta^T(k), \tag{3.6}$$
where $\Delta(k)$ is the difference of the mean vectors (of dimension $d$) computed from the first $k$ and the last $N - k$ data vectors, and $D_d$ is the $d \times d$ matrix of estimated residual vectors. If $d$ is large, the inverse of $D_d$ is unstable. In statistic (3.4), this inverse is “replaced” by inverses of the $d$ largest eigenvalues $\hat{\lambda}_t$, and the whole statistic is properly “diagonalized” so that only the most important variability of the data is considered, while the high dimensional noise is ignored.

We now turn to the behavior of the test under the alternative. We will show that it is consistent, i.e. $S_{N,d} \overset{P}{\to} \infty$. In fact, we can obtain the rate of divergence: under $H_A$, $S_{n,d}$ grows linearly with $N$. We formulate these results under the assumption of one change point.

Under Assumption 2.4, for $1 \leq k \leq d$, introduce the functions

$$g_k(x) = \begin{cases} 
x(1 - \theta) \int (\mu_1(t) - \mu_2(t))w_k(t)dt, & 0 < x \leq \theta \\
\theta(1 - x) \int (\mu_1(t) - \mu_2(t))w_k(t)dt, & \theta < x < 1.
\end{cases}$$

(3.7)

**Theorem 3.2** Under Assumption 2.4,

$$\sup_{0 \leq x \leq 1} \left| N^{-1} T_N - g^T(x) \Sigma^* g(x) \right| = o_P(1),$$

where

$$g(x) = [g_1(x), \ldots, g_d(x)]^T; \quad \Sigma^* = \begin{bmatrix} 1/\gamma_1 & 0 & \cdots & 0 \\
0 & 1/\gamma_2 & \cdots & 0 \\
\vdots & \vdots & \ddots & \vdots \\
0 & 0 & \cdots & 1/\gamma_d \end{bmatrix}.$$

Theorem 3.2 is proved in the Appendix.

It follows that the test statistic (3.4) satisfies the law of large numbers under the alternative, i.e.

$$\frac{1}{N} S_{N,d} \overset{P}{\to} \sum_{1 \leq k \leq d} \frac{1}{\gamma_k} \int_0^1 g_k^2(x)dx.$$

If $\int_0^1 g_k^2(x)dx > 0$ for some $1 \leq k \leq d$, then $S_{N,d} \overset{P}{\to} \infty$.

To understand when the test is consistent, introduce the jump function $\Delta(t) = \mu_1(t) - \mu_2(t)$. By (3.7), the condition $\int_0^1 g_k^2(x)dx > 0$ is equivalent to $\int_0^1 \Delta(s)w_k(s)ds \neq 0$. Thus the test will have no power if

$$\int_0^1 \Delta(s)w_k(s)ds = 0, \quad \text{for all } 1 \leq k \leq d.$$
(3.9) \[ \int c(t, s)w_k(s)ds = \gamma_k w_k(t), \quad \text{for all } 1 \leq k \leq d. \]

Comparing to (2.6), we see that condition (3.8) means that, up to a sign, the \( w_k, \gamma_k \) are equal to \( v_k, \lambda_k \), for \( 1 \leq k \leq d \). This lead us to the following corollary.

**Corollary 3.1** If Assumption 2.4 holds, and the jump function \( \Delta(t) = \mu_1(t) - \mu_2(t) \) is not orthogonal to the subspace spanned by the first \( d \) eigenfunctions of the covariance kernel \( c(t, s) \) (2.3), then \( S_{N,d} \overset{P}{\to} \infty \), as \( N \to \infty \).

To estimate the change point, we plot the function \( T_N(x) \) (3.3) against \( 0 \leq x \leq 1 \), and estimate \( \theta \) by the value of \( x \) which maximizes \( T_N(x) \). The intuition behind this estimator is clear from (3.3) and (3.2). To ensure uniqueness, we formally define this estimator as

(3.10) \[ \hat{\theta}_N = \inf \left\{ x : T_N(x) = \sup_{0 \leq y \leq 1} T_N(y) \right\}. \]

Its weak consistency is established in the following proposition

**Proposition 3.1** If the assumptions of Corollary 3.1 hold, then \( \hat{\theta}_N \overset{P}{\to} \theta \).

**Proof:** The argument \( x \) maximizing \( T_N(x) \) clearly maximizes \( A_N(x) = N^{-1}T_N(x) \). Theorem 3.2 states that \( \sup_{0 \leq x \leq 1} |A_N(x) - A(x)| \overset{P}{\to} 0 \), where

\[
A(x) = g^T(x) \Sigma^* g(x) = \begin{cases} 
  x(1 - \theta)A, & 0 \leq x \leq \theta \\
  \theta(1 - x)A, & \theta < x < 1,
\end{cases}
\]

with

\[
A = \sum_{1 \leq \ell \leq d} \frac{1}{\gamma_\ell} \left( \int \Delta(t)w_\ell(t)dt \right)^2.
\]

Under the assumptions of Corollary 3.1, \( A > 0 \), and it is easy to verify that \( A(x) \) has then a unique maximum at \( x = \theta \).

An important aspect of the procedure is the choice of the number \( d \) of the eigenfunctions \( v_k \). This issue is common to all FDA procedures using functional PCA, and several approaches have been proposed. These include an adaptation of the *scree plot* of Cattell (1966), see Kokoszka et al. (2008), the *cumulative percentage variance* approach used in Section 4.2, the *pseudo AIC* and the *cross-validation*, see Yao et al. (2005). All these methods are implemented in the MATLAB PACE package developed at the University of California at Davis. A general recommendation for the cumulative percentage variance method is to use \( d \) which explains 85% of the variance. This choice is suitable in the setting of Section 4.2, where \( d = 8 \) explains 84% of the variance.
### Table 4.1 Simulated critical values of the distribution of $K_d$.

| Nominal size | $d$ | 1   | 2   | 3   | 4   | 5   | 6   | 7   | 8   | 9   | 10  | 11  | 12  | 13  | 14  | 15  | 16  | 17  | 18  | 19  | 20  | 21  | 22  | 23  | 24  | 25  | 26  | 27  | 28  | 29  | 30  |
|--------------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| 10%          |     | 0.345165 | 0.606783 | 0.842567 | 1.065349 | 1.279713 | 1.485200 | 1.690773 | 1.897365 | 2.096615 | 2.288572 | 2.496635 | 2.686238 | 2.884214 | 3.066906 | 3.268958 | 3.462039 | 3.650724 | 3.837678 | 4.024313 | 4.214800 | 4.404677 | 4.591972 | 4.778715 | 4.965613 | 5.159057 | 5.346543 | 5.521107 | 5.714145 | 5.885108 | 6.083306 |

### 4 Finite sample performance and application to temperature data

In this section, we report the results of a simulation study that examines the finite sample performance of the test. Recall that the test rejects if $S_{N,d}$ (3.4) exceeds the $(1 - \alpha)$th quantile of $K_d$ (3.5). For $d \leq 5$, these quantiles were computed by Kiefer (1959) using a series expansion of the CDF of $K_d$. Horváth et al. (1999) used these expansions to find the critical values for $d = 12$ and noticed that the critical values obtained by simulating $K_d$ by discretizing the integral are slightly different, but actually lead to more accurate tests. To cover a fuller range of the $d$ values, Table 4.1 gives simulated critical values for $d = 1, \ldots, 30$, computed by discretizing the integral over 1,000 points and running 100,000 replications.

The simulation study consists of two parts. First we use standard Gaussian processes as the errors $Y_i$ and a number of rather arbitrary mean functions $\mu$. This part assesses...
the test in some generic cases analogous to assuming a normal distribution of scalar observations. In the second part, we use mean functions and errors derived from monthly temperature data. No assumptions on the marginal distribution of the \( Y_i \)'s or the shape of the \( \mu \)'s are made. This part assesses the test in a specific, practically relevant setting.

### 4.1 Gaussian processes

To investigate the empirical size, without loss of generality, \( \mu(t) \) was chosen to be equal to zero and two different cases of \( Y_i(t) \) were considered, namely the trajectories of the standard Brownian motion (BM), and the Brownian bridge (BB). These processes were generated by transforming cumulative sums of independent normal variables computed on a grid of \( 10^3 \) equispaced points in \([0, 1]\). Following Ramsay and Silverman (2005) (Chapter 3) discrete trajectories were converted to functional observations (functional objects in \( R \)) using B-spline and Fourier bases and various numbers of basis functions. No systematic dependence either on the type of the basis or on the number of basis functions was found.

The results reported in this section were obtained using B-spline basis with 800 basis functions. We used a wide spectrum of \( N \) and \( d \), but to conserve space, we present the results for \( N = 50, 150, 200, 300, 500 \) and \( d = 1, 2, 3, 4 \). All empirical rejection rates are based on 1,000 replications.

Table 4.2 shows the empirical sizes based on critical values reported in Table 4.1. The empirical sizes are fairly stable. Except for a very few cases of small sample sizes, all deviations from the nominal significance levels do not exceed two standard errors computed using the normal approximation \( \sqrt{p(1-p)/R} \), where \( p \) is a nominal level and \( R \) the number of repetitions. Table 4.2 shows that for these Gaussian processes, the empirical size does not depend appreciably either on \( n \) or on \( d \).

In the power study, several cases that violate the null were considered. We report the power for \( k^* = [N/2] \). Several other values of \( k^* \) were also considered, and only a small loss of power was observed for \( N/4 < k^* \leq 3N/4 \). A few different mean functions \( \mu \) before and after change were used, namely \( \mu_i(t) = 0, t, t^2, \sqrt{t}, e^t, \sin(t), \cos(t), i = 1, 2 \), for instance \( \mu_1(t) = t \) and \( \mu_2(t) = \cos(t) \), etc.

Table 4.3 presents selected results of the power study. It shows that the test has overall good power. For small samples, \( N \leq 100 \), in cases where the BB was used the power is slightly higher than for those with the BM. Nonetheless, for \( N \geq 150 \) the power approaches 100% for both processes and all choices of other parameters. The power decreases as the number of principal components \( d \) increases. This can be explained as follows: the critical values of \( S_{N,d} \) increase with \( d \), but the change point is mainly captured by a few initial leading principal components explaining the major part of the variance.
Table 4.2 Empirical size (in percent) of the test using the B-spline basis.

<table>
<thead>
<tr>
<th>Process</th>
<th>$d=1$</th>
<th>$d=2$</th>
<th>$d=3$</th>
<th>$d=4$</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>10%</td>
<td>5%</td>
<td>1%</td>
<td>10%</td>
</tr>
<tr>
<td>BM</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>10.3</td>
<td>4.6</td>
<td>0.1</td>
<td>9.9</td>
</tr>
<tr>
<td></td>
<td>11.2</td>
<td>5.5</td>
<td>0.8</td>
<td>10.6</td>
</tr>
<tr>
<td></td>
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<td></td>
<td></td>
</tr>
<tr>
<td>BB</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>12.2</td>
<td>5.6</td>
<td>1.3</td>
<td>9.8</td>
</tr>
<tr>
<td></td>
<td>12.4</td>
<td>5.7</td>
<td>0.7</td>
<td>10.2</td>
</tr>
<tr>
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</tr>
<tr>
<td>BM</td>
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<td></td>
</tr>
<tr>
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<td>10.8</td>
<td>5.7</td>
<td>1.3</td>
<td>9.7</td>
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<td>10.5</td>
<td>5.0</td>
<td>1.2</td>
<td>9.8</td>
</tr>
<tr>
<td></td>
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</tr>
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<td></td>
<td>9.7</td>
<td>5.4</td>
<td>0.8</td>
<td>9.2</td>
</tr>
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<td>5.1</td>
<td>0.8</td>
<td>10.8</td>
</tr>
<tr>
<td></td>
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<td></td>
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<td></td>
</tr>
<tr>
<td>BM</td>
<td></td>
<td></td>
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<td></td>
</tr>
<tr>
<td></td>
<td>10.3</td>
<td>5.2</td>
<td>1.5</td>
<td>11.1</td>
</tr>
<tr>
<td></td>
<td>10.4</td>
<td>5.6</td>
<td>1.1</td>
<td>9.4</td>
</tr>
<tr>
<td></td>
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<td></td>
</tr>
<tr>
<td>BB</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>11.6</td>
<td>6.3</td>
<td>1.3</td>
<td>10.6</td>
</tr>
<tr>
<td></td>
<td>11.7</td>
<td>5.1</td>
<td>1.3</td>
<td>9.7</td>
</tr>
</tbody>
</table>

$N = 50$

$N = 100$

$N = 150$

$N = 200$

$N = 300$

$N = 500$
Table 4.3 Empirical power (in percent) of the test using B-spline basis. Change point at $k^* = \lceil n/2 \rceil$.

<table>
<thead>
<tr>
<th>Process</th>
<th>$d=1$</th>
<th></th>
<th></th>
<th>$d=2$</th>
<th></th>
<th></th>
<th>$d=3$</th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>10%</td>
<td>5%</td>
<td>1%</td>
<td>10%</td>
<td>5%</td>
<td>1%</td>
<td>10%</td>
<td>5%</td>
<td>1%</td>
</tr>
<tr>
<td>$N = 50$</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>BM; BM + sin(t)</td>
<td>81.5</td>
<td>70.8</td>
<td>43.7</td>
<td>72.6</td>
<td>60.0</td>
<td>33.2</td>
<td>67.7</td>
<td>54.9</td>
<td>27.3</td>
</tr>
<tr>
<td>BM; BM + t</td>
<td>88.4</td>
<td>78.0</td>
<td>54.1</td>
<td>84.7</td>
<td>74.0</td>
<td>45.4</td>
<td>77.5</td>
<td>64.3</td>
<td>36.0</td>
</tr>
<tr>
<td>BB; BB + sin(t)</td>
<td>99.8</td>
<td>99.4</td>
<td>97.4</td>
<td>100</td>
<td>100</td>
<td>99.9</td>
<td>100</td>
<td>100</td>
<td>100</td>
</tr>
<tr>
<td>BB; BB + t</td>
<td>99.9</td>
<td>99.8</td>
<td>98.9</td>
<td>100</td>
<td>100</td>
<td>99.9</td>
<td>100</td>
<td>100</td>
<td>100</td>
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<tr>
<td>$N = 100$</td>
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<td></td>
<td></td>
</tr>
<tr>
<td>BM; BM + sin(t)</td>
<td>97.4</td>
<td>95.3</td>
<td>86.3</td>
<td>96.4</td>
<td>91.0</td>
<td>76.5</td>
<td>93.5</td>
<td>88.0</td>
<td>68.7</td>
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<tr>
<td>BM; BM + t</td>
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<td>97.5</td>
<td>91.2</td>
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<td>97.5</td>
<td>94.9</td>
<td>83.8</td>
</tr>
<tr>
<td>BB; BB + sin(t)</td>
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<tr>
<td>BB; BB + t</td>
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<td>100</td>
<td>100</td>
<td>100</td>
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<td></td>
<td></td>
</tr>
<tr>
<td>BM; BM + sin(t)</td>
<td>99.9</td>
<td>99.5</td>
<td>96.6</td>
<td>99.6</td>
<td>98.6</td>
<td>95.1</td>
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<td>99.7</td>
<td>98.8</td>
<td>99.9</td>
<td>99.7</td>
<td>97.8</td>
</tr>
<tr>
<td>BB; BB + sin(t)</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
</tr>
<tr>
<td>BB; BB + t</td>
<td>100</td>
<td>100</td>
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<td>100</td>
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<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>BM; BM + sin(t)</td>
<td>100</td>
<td>99.9</td>
<td>99.1</td>
<td>100</td>
<td>99.8</td>
<td>99.0</td>
<td>99.9</td>
<td>99.7</td>
<td>98.2</td>
</tr>
<tr>
<td>BM; BM + t</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>99.9</td>
<td>100</td>
<td>100</td>
<td>99.3</td>
</tr>
<tr>
<td>BB; BB + sin(t)</td>
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<td>100</td>
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<tr>
<td>BB; BB + t</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
</tr>
</tbody>
</table>
4.2 Analysis of central England temperatures

The goal of this section is twofold: to investigate the performance of the test in a real world setting, and to demonstrate the advantages of the functional approach for high-dimensional data.

The data consists of 228 years (1780 to 2007) of average daily temperatures in central England. The original data can thus be viewed as 228 curves with 365 measurements on each curve. These data were converted to functional objects in R using 12 B-spline basis functions. Multivariate observations were obtained as in Horváth et al. (1999) by computing monthly averages resulting in 228 vectors of dimension $d = 12$. (We could not even compute statistics (3.6) for vectors of dimension 365 because R reported that $\hat{D}$ was singular.) These two procedures are illustrated in Figure 4.1. Even though we used 12 B-splines and 12 averages, the resulting data look quite different, especially in spring and fall, when the temperatures change most rapidly. Gregorian months form a somewhat arbitrary fixed partition of the data, while the splines adapt to their shapes which differ from year to year.

To compute statistic (3.4), we used $d = 8$ eigenfunctions which explain 84% of variability. If the test indicates a change, we estimate it by the estimator $\hat{\theta}_N$ (3.10). This divides the data set into two subsets. The procedure is then repeated for each subset until periods of constant mean functions are obtained. We proceed in exactly the same
Table 4.4 Segmentation procedure of the data into periods with constant mean function.

<table>
<thead>
<tr>
<th>Iteration</th>
<th>Segment</th>
<th>Decision</th>
<th>$S_{N,d}$</th>
<th>$M_{N,d}$</th>
<th>P-value</th>
<th>Estimated change point</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1780 - 2007</td>
<td>Reject</td>
<td>8.020593</td>
<td>1926</td>
<td></td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>1780 - 1925</td>
<td>Reject</td>
<td>3.252796</td>
<td>1808</td>
<td></td>
<td></td>
</tr>
<tr>
<td>3</td>
<td>1780 - 1807</td>
<td>Accept</td>
<td>0.888690</td>
<td>1850</td>
<td></td>
<td></td>
</tr>
<tr>
<td>4</td>
<td>1808 - 1925</td>
<td>Reject</td>
<td>2.351132</td>
<td>-</td>
<td></td>
<td></td>
</tr>
<tr>
<td>5</td>
<td>1808 - 1849</td>
<td>Accept</td>
<td>0.890845</td>
<td>-</td>
<td></td>
<td></td>
</tr>
<tr>
<td>6</td>
<td>1850 - 1925</td>
<td>Accept</td>
<td>1.364934</td>
<td>-</td>
<td></td>
<td></td>
</tr>
<tr>
<td>7</td>
<td>1926 - 2007</td>
<td>Reject</td>
<td>2.311151</td>
<td>1993</td>
<td></td>
<td></td>
</tr>
<tr>
<td>8</td>
<td>1926 - 1992</td>
<td>Accept</td>
<td>0.927639</td>
<td>-</td>
<td></td>
<td></td>
</tr>
<tr>
<td>9</td>
<td>1993 - 2007</td>
<td>Accept</td>
<td>1.626515</td>
<td>-</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

England temperatures ($d = 8$) (FDA approach)

<table>
<thead>
<tr>
<th>Iteration</th>
<th>Segment</th>
<th>Decision</th>
<th>$S_{N,d}$</th>
<th>$M_{N,d}$</th>
<th>P-value</th>
<th>Estimated change point</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>1780 - 2007</td>
<td>Reject</td>
<td>7.971031</td>
<td>1926</td>
<td></td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>1780 - 1925</td>
<td>Reject</td>
<td>3.576543</td>
<td>1815</td>
<td></td>
<td></td>
</tr>
<tr>
<td>3</td>
<td>1780 - 1814</td>
<td>Accept</td>
<td>1.534223</td>
<td>-</td>
<td></td>
<td></td>
</tr>
<tr>
<td>4</td>
<td>1815 - 1925</td>
<td>Accept</td>
<td>2.813596</td>
<td>-</td>
<td></td>
<td></td>
</tr>
<tr>
<td>5</td>
<td>1926 - 2007</td>
<td>Accept</td>
<td>2.744801</td>
<td>-</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

England temperatures ($d = 12$) (MDA approach)

Table 4.5 Summary and comparison of segmentation. Beginning and end of data period in bold.

<table>
<thead>
<tr>
<th>Approach</th>
<th>Change points</th>
</tr>
</thead>
<tbody>
<tr>
<td>FDA</td>
<td>1780 1808 1850 1926 1992 2007</td>
</tr>
<tr>
<td>MDA</td>
<td>1780 1815 1926 2007</td>
</tr>
</tbody>
</table>
manner using statistic (3.6). We refer to these procedures, respectively, as FDA and MDA approaches. The resulting segmentations are shown in Tables 4.4 and 4.5.

The functional approach identified two more change point, 1850 and 1992, which roughly correspond to the beginning of mass industrialization and the advent of rapid global warming. The multivariate approach “almost” identified these change points with the P–values in iterations 4 and 5 being just above the significance level of 5%. This may indicate that the functional method has better power, perhaps due to its greater flexibility in capturing the shape of the data. This conjecture is investigated below. Figure 4.2 shows average temperatures in the last four segments, and clearly illustrates the warming trend.

The analysis presented above assumes a simple functional change point model for the daily temperatures. Obviously, one cannot realistically believe that the mean curves
change abruptly in one year, this is merely a modeling assumption useful in identifying patterns of change in mean temperature curves. Well-established alternative modeling approaches have been used to study the variability of temperatures. For example, Hosking (1984) fitted a fractionally differenced ARMA(1,1) model to the series of annual average temperatures in central England in 1659–1976. It is generally very difficult to determine on purely statistical grounds if a change-point or a long-range dependent model is more suitable for any particular finite length record, see Berkes et al. (2006) and Jach and Kokoszka (2008) for recent methodology, discussion and references. It is often more useful to choose a modeling methodology which depends on specific goals, and this is the approach we use. One way of checking an approximate adequacy of our model is to check if the residuals obtained after subtracting the mean in each segment are approximately independent and identically distributed. This can be done by applying the test developed by Gabrys and Kokoszka (2007) which is a functional analog of the well-known test of Hosking (1980) and Li and McLeod (1981) (see also Hosking (1981, 1989). The P-value of 8% indicates the acceptance of the hypothesis that the residuals are iid.

Keeping these caveats in mind, we use the partitions obtained above to generate realistic synthetic data with and without change-points. We use them to evaluate and compare the size and power properties of the FDA and MDA tests, and to validate our findings. We compute the residuals of every observation in a constant mean segment by subtracting the average of the segment, i.e. \( \hat{Y}_{is} = X_{is} - \hat{\mu}_s \), where \( s = 1, \ldots, S \) denotes the segment, and \( i = 1, \ldots, I_s \) indexes observations in the \( s \)th segment. The \( \hat{Y}_{is} \) are functional residuals, and their average in each segment is clearly the zero function.

To assess the empirical size, we simulate “temperature-like” data by considering two cases. Case I: for every constant mean segment \( s \), we produce synthetic observations by adding to its mean function \( \hat{\mu}_s \) errors drawn from the empirical distribution of the residuals of that segment, i.e. synthetic (bootstrap) observations in the \( s \)th segment are generated via \( X_{is}^* = \hat{\mu}_s + \hat{Y}_{i^*_s} \), where \( i^* \) indicates that \( \hat{Y}_{i^*_s} \) is obtained by drawing with replacement from \( \{ \hat{Y}_{is}, i = 1, \ldots, I_s \} \). Case II: We compute residuals in each segment and pool them together. We use this larger set of residuals to create new observations by adding to the average of a segment the errors drawn with replacement from that pool of residuals. For each segment, we generate 1000 of these bootstrap sequences. Table 4.6 shows the the resulting empirical sizes. As the sample size increases, the FDA rejection rates approach nominal sizes, while the MDA test is much more conservative. For the 1993–2007 segment, the size is not reported because the matrix \( D \) was (numerically) singular for most bootstrap replications.

We next investigate the power. Three cases are considered. Case I: For each segment, we produce synthetic observations using the bootstrap procedure and sampling residuals from a corresponding period. This means that the errors in each segment come from possibly different distributions. Case II: We pool together two, three, four, or five sets
of residuals (depending on how many constant mean segments we consider) and sample from that pool to produce new observations. This means that the errors in each segment come from the same distribution. Case III: We slightly modify Case II by combining all residuals from all segments into one population and use it to produce new observations. In both Case II and Case III, the theoretical assumptions of Section 2 are satisfied, cf. Assumption 2.4, i.e. the means change, but the errors come from the same population. Table 4.7 shows the power of the test for FDA approach and Table 4.8 presents results of discrete MDA method. As seen in Table 4.7, the differences between the three cases are of the order of the chance error. Table 4.7 shows that the test has excellent power, even in small samples, both for single and multiple change points. As for the Gaussian processes, power is slightly higher if there is a change point around the middle of the sample. Comparing Tables 4.7 and 4.8, it is seen that in FDA approach dominates the MDA approach. There are a handful of cases, indicated with *, when MDA performed better, but their frequency and the difference size suggests that this may be attributable to the chance error.

Acknowledgements

István Berkes was partially supported by OTKA grants T43037, K61052 and K67961. Lajos Horváth was partially supported by NSF grant DMS-0604670. Robertas Gabrys and Piotr Kokoszka were partially supported by NSF grants DMS-0413653 and DMS-0804165. The temperature data used in Section 4 were published by the British Atmospheric Data Centre, and compiled by Nick Humphreys.
Table 4.7 Empirical power of the test for change-point models derived from temperature data (FDA approach).

<table>
<thead>
<tr>
<th>Segment</th>
<th>Sample size</th>
<th>Change point(s) $\theta$</th>
<th>Nominal level</th>
<th>Case I</th>
<th>Case II</th>
<th>Case III</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>10%</td>
<td>5%</td>
<td>1%</td>
</tr>
<tr>
<td><strong>England</strong> ($d = 8$) (FDA approach)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>$\Delta_1, \Delta_2$</td>
<td>70</td>
<td>.41</td>
<td>85.6</td>
<td>76.8</td>
<td>49.7</td>
<td>86.4</td>
</tr>
<tr>
<td>$\Delta_1, \Delta_3$</td>
<td>104</td>
<td>.28</td>
<td>86.2</td>
<td>75.8</td>
<td>47.4</td>
<td>88.6</td>
</tr>
<tr>
<td>$\Delta_1, \Delta_4$</td>
<td>94</td>
<td>.31</td>
<td>100</td>
<td>99.7</td>
<td>98.7</td>
<td>100</td>
</tr>
<tr>
<td>$\Delta_1, \Delta_5$</td>
<td>44</td>
<td>.66</td>
<td>100</td>
<td>99.9</td>
<td>93.4</td>
<td>100</td>
</tr>
<tr>
<td>$\Delta_2, \Delta_3$</td>
<td>118</td>
<td>.36</td>
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<td>99.5$^*$</td>
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Table 4.8 Empirical power of the test for change-point models derived from temperature data (MDA approach).

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<th>Segment</th>
<th>Sample size</th>
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<th>Nominal level</th>
<th>Case I</th>
<th>Case II</th>
<th>Case III</th>
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<td></td>
<td>78.4</td>
<td>63.1</td>
<td>28.0</td>
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</table>

| $\Delta_1$, $\Delta_2$, $\Delta_3$ | 146         | .20 .49                  |               | 97.5   | 93.2    | 76.9     | 97.7   | 93.1    | 77.9     | 97.4   | 94.9    | 80.2     |
| $\Delta_1$, $\Delta_2$, $\Delta_4$ | 136         | .21 .52                  |               | 100    | 100     | 100      | 100    | 100     | 99.9     | 100    | 100     | 99.9     |
| $\Delta_1$, $\Delta_2$, $\Delta_5$ | 86          | .34 .83                  |               | 100    | 99.8    | 96.2     | 99.9   | 99.7    | 95.7     | 100    | 99.8    | 97.4     |
| $\Delta_2$, $\Delta_3$, $\Delta_4$ | 184         | .23 .65                  |               | 100    | 100     | 99.1     | 100    | 99.9    | 98.7     | 100    | 100     | 99.5     |
| $\Delta_2$, $\Delta_3$, $\Delta_5$ | 134         | .32 .89                  |               | 99.8   | 99.4    | 93.7     | 99.6   | 99.3    | 93.8     | 99.7   | 98.6    | 92.1     |
| $\Delta_3$, $\Delta_4$, $\Delta_5$ | 158         | .49 .91                  |               | 100    | 100     | 100      | 100    | 100     | 100      | 100    | 100     | 100      |
| $\Delta_1$, $\Delta_2$, $\Delta_3$, $\Delta_4$ | 212         | .14 .33 .69              |               | 100    | 100     | 99.9     | 100    | 100     | 100      | 100    | 100     | 100      |
| $\Delta_1$, $\Delta_2$, $\Delta_3$, $\Delta_5$ | 162         | .18 .44 .91              |               | 100    | 100     | 99.1     | 100    | 99.9    | 99.1     | 100    | 100     | 98.9     |
| $\Delta_2$, $\Delta_3$, $\Delta_4$, $\Delta_5$ | 200         | .22 .60 .93              |               | 100    | 100     | 100      | 100    | 100     | 100      | 100    | 100     | 100      |
| $\Delta_1$, $\Delta_2$, $\Delta_3$, $\Delta_4$, $\Delta_5$ | 228         | .13 .31 .64 .93          |               | 100    | 100     | 100      | 100    | 100     | 100      | 100    | 100     | 100      |
A Proof of Theorems 3.1 and 3.2

A key element of the proofs is the bound (A.5), which follows from a functional central limit theorem in a Hilbert space. A result of this type is needed because the observations $X_i(\cdot)$ are elements of a Hilbert space, and to detect a change point, we must monitor the growth of the partial sums $\sum_{1 \leq i \leq N} X_i(t)$ which are a function of $0 < x < 1$ (and of $t \in T$).

Lemma A.1 is particularly noteworthy because it shows that the eigenvalues and the eigenfunctions also converge under the alternative.

A.1 Proof of Theorems 3.1

We will work with the unobservable projections

$$\tilde{\beta}_{i,k} = \int Y_i(t) \hat{v}_k(t) dt, \quad \beta_{i,k} = \int Y_i(t) v_k(t) dt, \quad \beta_{i,k}^* = \hat{c}_k \beta_{i,k}$$

and the vectors

$$\beta_i = [\beta_{i,1}, \ldots, \beta_{i,d}]^T, \quad \beta_i^* = [\beta_{i,1}^*, \ldots, \beta_{i,d}^*]^T, \quad 1 \leq i \leq N.$$

Since the $Y_i$ are iid functions with mean zero, the $\beta_i$ are iid mean zero vectors in $\mathbb{R}^d$. A simple calculation using the orthonormality of the $v_k$ shows that each $\beta_i$ has a diagonal covariance matrix

$$\Sigma_d = \begin{bmatrix} \lambda_1 & 0 & \cdots & 0 \\ 0 & \lambda_2 & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & \lambda_d \end{bmatrix}$$

The functional central limit theorem, thus implies that

$$(A.1) \quad N^{-1/2} \sum_{1 \leq i \leq N} \beta_i \overset{d}{\rightarrow} \Delta_d(x) \quad (0 \leq x \leq 1),$$

where the convergence is in the Skorokhod space $D^d[0, 1]$. The process $\{\Delta_d(x), 0 \leq x \leq 1\}$ takes values in $\mathbb{R}^d$, has zero mean and covariance matrix $\Sigma_d$. Convergence (A.1) implies in turn that

$$(A.2) \quad \frac{1}{N} \left[ \sum_{1 \leq i \leq N} \beta_i - x \sum_{1 \leq i \leq N} \beta_i \right]^T \Sigma_d^{-1} \left[ \sum_{1 \leq i \leq N} \beta_i - x \sum_{1 \leq i \leq N} \beta_i \right] \overset{d}{\rightarrow} \sum_{1 \leq i \leq d} B_i^2(x)$$

in the Skorokhod space $D[0, 1]$.

The matrix $\Sigma_d$ is estimated by $\hat{\Sigma}_d$. By (2.8) and Assumption 2.2, $\hat{\Sigma}_d^{-1} \overset{P}{\rightarrow} \Sigma_d^{-1}$, so (A.2) yields

$$(A.3) \quad \frac{1}{N} \left[ \sum_{1 \leq i \leq N} \beta_i - x \sum_{1 \leq i \leq N} \beta_i \right]^T \hat{\Sigma}_d^{-1} \left[ \sum_{1 \leq i \leq N} \beta_i - x \sum_{1 \leq i \leq N} \beta_i \right] \overset{d}{\rightarrow} \sum_{1 \leq i \leq d} B_i^2(x).$$
Note that
\[
\sum_{1 \leq i \leq Nx} \beta_{i,k}^* - x \sum_{1 \leq i \leq N} \beta_{i,k}^* = \hat{c}_k \left( \sum_{1 \leq i \leq Nx} \beta_{i,k} - x \sum_{1 \leq i \leq N} \beta_{i,k} \right).
\]

Since \( \hat{c}_k^2 = 1 \), we can replace the \( \beta_i \) in (A.3) by the \( \beta_i^* \), and obtain
\[
(A.4) \quad \frac{1}{N} \left[ \sum_{1 \leq i \leq Nx} \beta_i^* - x \sum_{1 \leq i \leq N} \beta_i^* \right]^T \tilde{S}_d^{-1} \left[ \sum_{1 \leq i \leq Nx} \beta_i^* - x \sum_{1 \leq i \leq N} \beta_i^* \right] \to \sum_{1 \leq i \leq d} B_i^2(x).
\]

We now turn to the effect of replacing the \( \beta_{i,k}^* \) by \( \tilde{\beta}_{i,k} \). Observe that
\[
\sup_{0 < x < 1} \left| \left( \sum_{1 \leq i \leq Nx} \beta_i^* - x \sum_{1 \leq i \leq N} \beta_i^* \right) - \sum_{1 \leq i \leq Nx} \tilde{\beta}_{i,k} \right| = \sup_{0 < x < 1} \left| \int \left( N^{-1/2} \sum_{1 \leq i \leq Nx} Y_i(t) \right) \left( \hat{c}_k v_k(t) - \hat{\nu}_k(t) \right) dt \right|
\]
\[
\leq \sup_{0 < x < 1} \left[ \int \left( N^{-1/2} \sum_{1 \leq i \leq Nx} Y_i(t) \right)^2 dt \right]^{1/2} \left[ \int \left( \hat{c}_k v_k(t) - \hat{\nu}_k(t) \right)^2 dt \right]^{1/2}.
\]

The first factor is bounded in probability, i.e.
\[
(A.5) \quad \sup_{0 < x < 1} \int \left( N^{-1/2} \sum_{1 \leq i \leq Nx} Y_i(t) \right)^2 dt = O_P(1).
\]

Relation (A.5) follows from the weak convergence in \( D([0,1], L^2(T)) \) of the partial sum process \( \sum_{1 \leq i \leq Nx} Y_i, x \in [0,1] \), see e.g. Kuelbs (1973).

Combining (A.5) and (2.8), we obtain
\[
\sup_{0 < x < 1} \left| \left( N^{-1/2} \sum_{1 \leq i \leq Nx} \beta_i^* - x \sum_{1 \leq i \leq N} \beta_i^* \right) - \sum_{1 \leq i \leq Nx} \tilde{\beta}_{i,k} \right| \to 0,
\]
which in turn implies that
\[
(A.6) \quad \left\| \left[ \sum_{1 \leq i \leq Nx} \beta_i^* - x \sum_{1 \leq i \leq N} \beta_i^* \right] - \left[ \sum_{1 \leq i \leq Nx} \tilde{\beta}_{i,k} - x \sum_{1 \leq i \leq N} \tilde{\beta}_{i,k} \right] \right\| = o_P(N^{-1/2}),
\]
where the norm is the Euclidean norm in \( R^d \). Relations (A.4) and (A.6) yield the claim in Theorem 3.1.
A.2 Proof of Theorem 3.2

Theorem 3.2 follows from relation (A.10) and Lemma A.2. To establish them, we need the following Lemma.

**Lemma A.1** Under assumption 2.4, for every $1 \leq k \leq d$, as $N \to \infty$,

(A.7) \[ \lambda_k \xrightarrow{P} \gamma_k, \]

(A.8) \[ \int [\dot{v}_k(t) - \dot{c}_k w_k(t)]^2 dt \xrightarrow{P} 0, \]

where $\dot{v}_k$, $\dot{\lambda}_k$ are defined by (2.7), $w_k, \gamma_k$ by (2.11) and $\dot{c}_k = \text{sign} \int_T v_k(t) \dot{v}_k(t)dt$.

**Proof:** It is easy to see that

$$X_N(t) = \bar{Y}_N(t) + \frac{k^*}{N} \mu_1(t) + \frac{N - k^*}{N} \mu_2(t)$$

and, denoting $\Delta(t) = \mu_1(t) - \mu_2(t)$,

$$\dot{c}_N(t, s) = \frac{1}{N} \left( \sum_{1 \leq i \leq k^*} + \sum_{k^* < i \leq N} \right) (X_i(t) - \bar{X}_N(t))(X_i(s) - \bar{X}_N(s))$$

$$= \frac{1}{N} \sum_{1 \leq i \leq k^*} \left( Y_i(t) - \bar{Y}_N(t) + \mu_1(t) - \frac{k^*}{N} \mu_1(t) - \frac{N - k^*}{N} \mu_2(t) \right)$$

$$\times \left( Y_i(s) - \bar{Y}_N(s) + \mu_1(s) - \frac{k^*}{N} \mu_1(s) - \frac{N - k^*}{N} \mu_2(s) \right)$$

$$+ \frac{1}{N} \sum_{k^* < i \leq N} \left( Y_i(t) - \bar{Y}_N(t) + \mu_2(t) - \frac{k^*}{N} \mu_1(t) - \frac{N - k^*}{N} \mu_2(t) \right)$$

$$\times \left( Y_i(s) - \bar{Y}_N(s) + \mu_2(s) - \frac{k^*}{N} \mu_1(s) - \frac{N - k^*}{N} \mu_2(s) \right)$$

$$= \frac{1}{N} \sum_{1 \leq i \leq k^*} \left( Y_i(t) - \bar{Y}_N(t) + \frac{N - k^*}{N} \Delta(t) \right) \left( Y_i(s) - \bar{Y}_N(s) + \frac{N - k^*}{N} \Delta(s) \right)$$

$$+ \frac{1}{N} \sum_{k^* < i \leq N} \left( Y_i(t) - \bar{Y}_N(t) - \frac{k^*}{N} \Delta(t) \right) \left( Y_i(s) - \bar{Y}_N(s) - \frac{k^*}{N} \Delta(s) \right).$$

Rearranging terms, we obtain

$$\dot{c}_N(t, s) = \frac{1}{N} \sum_{i=1}^{N} \left( Y_i(t) - \bar{Y}_N(t) \right) \left( Y_i(s) - \bar{Y}_N(s) \right) + \frac{k^*}{N} \left( 1 - \frac{k^*}{N} \right) \Delta(t) \Delta(s) + r_N(t, s),$$

where

$$r_N(t, s) = \left( 1 - \frac{k^*}{N} \right) \frac{1}{N} \sum_{1 \leq i \leq k^*} \left[ (Y_i(t) - \bar{Y}_N(t)) \Delta(s) + (Y_i(s) - \bar{Y}_N(s)) \Delta(t) \right].$$
\[ \sum_{k^* < i < N} \left[ (Y_i(t) - \bar{Y}_N(t)) \Delta(s) + (Y_i(s) - \bar{Y}_N(s)) \Delta(t) \right]. \]

Using the law of large numbers for independent, identically distributed Hilbert space valued random variables (see e.g. Theorem 2.4 of Bosq (2000)), we obtain \( \int_T \int_T \epsilon_N^2(t, s) dt ds \rightarrow 0 \) and
\[
(A.9) \quad \int \int [\hat{\epsilon}_N(t, s) - \check{\epsilon}_N(t, s)]^2 \rightarrow 0.
\]

Hence Lemmas 4.2 and 4.3 of Bosq (2000) imply, respectively, (A.7) and (A.8).

As an immediate corollary to (A.7), we obtain
\[
(A.10) \quad \frac{1}{N} \sum_{i,k} \hat{\eta}_{i,k} - x \sum_{1 \leq i \leq N} \hat{\eta}_{i,k} = o_P(1),
\]

with the functions \( g_k \) defined by (3.7).

**Proof:** Denote
\[
\hat{g}_k(x) = \frac{1}{N} \left[ \sum_{1 \leq i \leq N} \hat{\eta}_{i,k} - x \sum_{1 \leq i \leq N} \hat{\eta}_{i,k} \right], \quad x \in [0, 1],
\]

and observe that
\[
\hat{\eta}_{i,k} = \int Y_i(t) \hat{v}_k(t) dt + \int \mu_1(t) \hat{v}_k(t) dt - \int X_i(t) \hat{v}_k(t) dt, \quad \text{if } 1 \leq i \leq k^*
\]
and
\[
\hat{\eta}_{i,k} = \int Y_i(t) \hat{v}_k(t) dt + \int \mu_2(t) \hat{v}_k(t) dt - \int X_i(t) \hat{v}_k(t) dt, \quad \text{if } k^* < i \leq N.
\]

We will use the relation
\[
(A.11) \quad \sup_{0 < x < 1} \left| \sum_{1 \leq i \leq N} \int Y_i(t) \hat{v}_k(t) dt \right| = O_P(N^{1/2}),
\]

which follows from (A.5).

Suppose first that \( 0 < x \leq \theta \). Then, by (A.11) and (A.8), uniformly in \( x \in [0, 1] \),
\[
\hat{g}_k(x) = x(1 - \theta) \left[ \int \mu_1(t) \hat{v}_k(t) dt - \int \mu_2(t) \hat{v}_k(t) dt \right] + o_P(N^{-1/2})
\]
\[= x(1 - \theta) \hat{c}_k \left[ \int \mu_1(t) \hat{w}_k(t) dt - \int \mu_2(t) \hat{w}_k(t) dt \right] + o_P(1).\]

If \( x > \theta \), then, uniformly in \( x \in [0, 1] \),
\[
\hat{g}_k(x) = \theta(1 - x) \hat{c}_k \left[ \int \mu_1(t) \hat{w}_k(t) dt - \int \mu_2(t) \hat{w}_k(t) dt \right] + o_P(1).
\]
References


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