

Cytometry inference through adaptive atomic deconvolution

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Abstract

In this paper we consider a statistical estimation problem known as atomic deconvolution. Introduced in reliability, we show that this model has a direct application when considering biological data produced by flow cytometers. In cytometer experiments, biologists measure the fluorescence emission of treated cells and compare them with their natural emission to study the presence of specific molecules on the cells' surface. They observe a signal which is composed of a noise (the natural fluorescence) plus some additional signal related to the quantity of molecule present on the surface if any. From a statistical point of view, we aim at inferring the percentage of cells expressing the selected molecule and the probability distribution function associated with its fluorescence emission. We propose here an adaptive estimation procedure based on a previous deconvolution procedure introduced by van Es, Gugushvili, and Spreij (2008) and Gugushvili, van Es, and Spreij (2011). For both estimating the mixing parameter and the mixing density automatically, we use the Lepskii method based on the optimal choice of a bandwidth parameter using a bias-variance decomposition. We then derive some concentration inequalities for our estimators and obtain some optimal rates of convergence, that are shown to be minimax optimal (up to some log terms) in Sobolev classes. Finally, we apply our algorithm on simulated and real biological datasets.

Keywords: Mixture models, Atomic deconvolution, Adaptive kernel estimators, Inverse problems .

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1 Introduction

1.1 Motivation

This work is motivated by an issue commonly encountered by biologist researchers when using flow cytometry measurements: a flow cytometer is an important technical instrument that makes it possible to assess the efficiency of a drug treatment. Such an electronic machine analyze a large number of cells and produce cell engineering results such as counting, sorting or bio-marker detection.

This technology is commonly used for measuring the expression levels of proteins on the cells' surface. To this aim, the biologist use fluorescent antibodies which bind with specific proteins on the surface of cells. The flow cytometer then measures on each cells the fluorescent intensity, which is a reflection of the quantity of marker expressed by the cell. More precisely, these measurements derive from a standard procedure in flow cytometry :

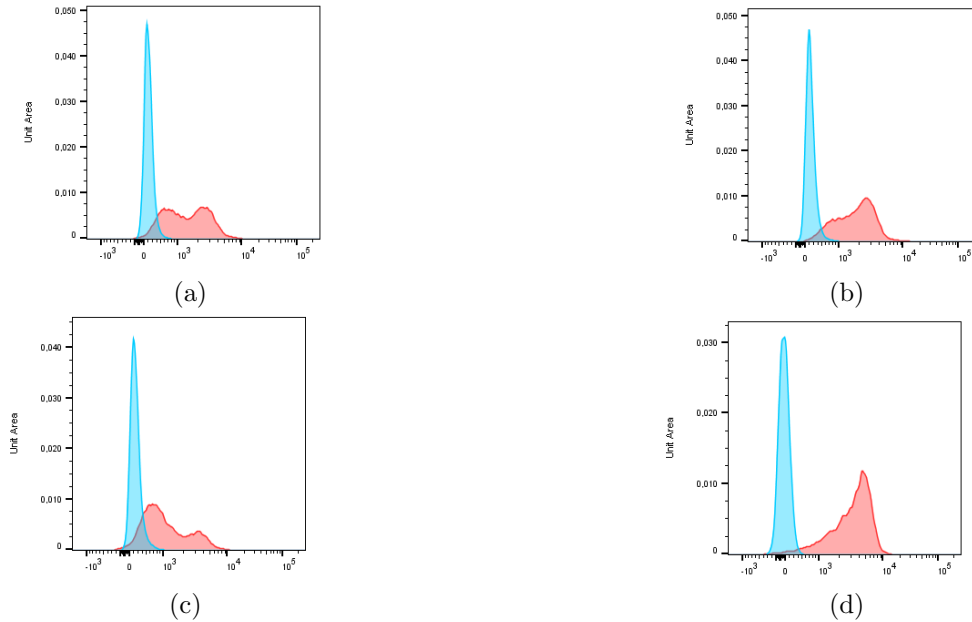


Figure 1: Effect of binding of an antibody specific for the CD27 protein expressed at the cell surface of lymphocytes extracted from the blood of an healthy individual (four different samples). The blue histograms represent the distribution of the baseline photo emission of untreated cells while the red histograms gives the fluorescent distribution of treated cells. We refer to Section 4.2 for more details on the samples.

- First the biologist performs a *calibration* that corresponds to a preliminary estimation of the baseline population of cells without any treatment: a large number of cells is placed in the cytometer and a fluorescent intensity is measured on each cell. This calibration ends with a baseline estimation of the the baseline photon emission by a population of cells and yields for example blue curves in Figure 1.
- Then the biologist exposes the cells to a fluorescent antibody, which binds with the marker of interest on the surface of cells. The new fluorescence empirical distribution is built by the cytometer. In particular, this step mimics the administration of a drug (related to a specific antibody) to a patient and is therefore useful for the understanding of the effect of this drug. This step allows to obtain the red curves in Figure 1.
- The expertise of the biologist is at last used to calibrate a qualitative analysis to decide if there is an effect (or not) of the treatment and what is its mean efficiency. We should mention that this human expertise may also be assisted by some statistical computational analyses and have resulted in an open project called *FlowCAP* that makes it possible to produce standard statistical analysis (essentially data clustering). However, the effect of the treatment is yet estimated visually by the biologist researcher.

Figure 1 is an illustration of the empirical distribution of fluorescence measured by a cytometer before and after a treatment on different kind of cells with a logarithmic scale (before treatment in blue, after treatment in red).

Our objective here is to enrich the possibilities for biologists by deriving an additional and new signal processing step for flow cytometers datasets. We aim to develop an automatic estimation of drug treatments in order to retrieve both the proportion of positive cells (*i.e.* the cells that express a given marker) and the distribution of marker in a population of cells. At the moment (one of the author of this paper is a biologist and intensive user of flow cytometers), there is no statistical tool developed to solve this issue and the analysis of the proportion of reacting cells is estimated empirically with a visual inspection of blue and red curves derived from the flow cytometers (see Figure 1). As we will see below, our approach will be to develop a nonparametric statistical deconvolution problem.

1.2 Model

1.2.1 Atomic deconvolution

The phenomenon described above can be described as an atomic deconvolution problem. Before treatment, the fluorescence of a cell possesses a natural fluorescence that is distributed randomly (as illustrated by the blue curves in Figure 1). This baseline fluorescence is described by a random variable denoted by e^U (the cytometer produces logarithmic scales so that some realisations of U are directly observed). Then the treatment is performed by biologists on a large set of cells and it is observed empirically that sometimes the treatment has unfortunately no effect on the cells, while the other times when the treatment works, it applies a random multiplicative factor denoted by e^X on the fluorescence of the cells.

Therefore, we can reasonably assume that we observe on the second step after treatment among the set of cells treated in the cytometer some i.i.d. realizations $(Z_i)_{1 \leq i \leq n}$ of a model given by:

$$Z = U + AX, \quad (1)$$

where U , A and X are jointly independent random variables. In Equation (1), the random variable Z is therefore the measurement of the flow cytometer of the fluorescent intensity on a given cell. The random variable U represents the natural fluorescence of the cell: it is its baseline reaction regardless of the impact of drug administrated during the experiment. The random variable X stands for the quantitative effect of the reagent when the cell actively reacts to the treatment. At last, A represents the effectiveness of the treatment on the cell and belongs to $\{0, 1\}$. We can therefore list below our definitions related to Equation (1):

- U represents the baseline “noise” on the observations, leading to the convolution model. We assume that the distribution of U is absolutely continuous with respect to the Lebesgue measure on \mathbb{R} and the density of U is assumed to be **known** and is denoted by g . This assumption for example translates a preliminary step of calibrating the cytometer before any treatment with a nonparametric estimation of the density g .
- A is distributed according to a Bernoulli random variable $A \sim \mathcal{B}(1 - p)$ and p is an **unknown** parameter. We use the convention $\mathbb{P}[A = 0] = p$ (the cell does not react to the administrated reagent) while $\mathbb{P}[A = 1] = 1 - p$ (the reagent induces an effect on the cell). We are interested in the estimation of the success of the treatment, *i.e.* in the estimation of $1 - p$.
- X is the quantitative effect of the treatment when the cell is reacting to the treatment and we assume that the distribution of X is also absolutely continuous with respect to the Lebesgue measure on \mathbb{R} , with an **unknown** density f . Of course, biologists are interested in estimating the quantitative effect of the treatment, so that we also need to produce an estimation of f .

For example, in each situation illustrated in Figure 1, we can see that the baseline density g is represented by the blue area while the “mixture” distribution obtained after the statistical contamination of U by AX is represented by the red area.

1.2.2 State-of-the-art and related works

Deconvolution Even though relatively simple in appearance, we will see below that Equation (1) deserves both theoretical development and numerical efforts to obtain a practical and theoretically supported method of estimation for p and f . For example, we could think of the use of a moment estimation strategy to obtain information on p . Unfortunately, we can rapidly compute that

$$\mathbb{E}(Z) = \mathbb{E}(U) + (1 - p)\mathbb{E}(X) \quad \text{and} \quad \mathbb{V}(Z) = \mathbb{V}(U) + p(1 - p)\mathbb{E}(X)^2 + (1 - p)\mathbb{V}(X),$$

leading to a non trivial relationship between the known moments of the distribution of U , the Bernoulli parameter p , and the moments of the distribution of X .

This problem has been recently introduced in van Es et al. (2008) and was motivated by a reliability problem estimation, which is an application completely different from our. To bypass the difficulty raised by atomic deconvolution, these authors introduced a nonparametric deconvolution strategy (see van Es et al. (2008) and Gugushvili et al. (2011)) adapted to the situation of atomic deconvolution.

We emphasize at this stage that if the baseline density g is assumed to be known, nothing else can be expected for the knowledge of the model. In particular, the parameter of the Bernoulli trials has to be estimated and there is no reason to assume a specific parametric form for the density f : in the red curves of Figure 1, it seems difficult to decipher a typical structure of the convolution kernel. Hence, the use of nonparametric statistics is here unavoidable for a practical feasible estimation of the density f .

The estimator of van Es et al. (2008) is inspired from the seminal contributions of Carroll and Hall (1988) and Fan (1991) and uses kernel density estimation associated with Fourier analysis. In particular, van Es et al. (2008) builds a consistent estimator and proves a central limit theorem related to this estimator while Gugushvili et al. (2011) studies the associated convergence rates and their minimax properties. The former works are related to the statistical deconvolution inverse problems (see Meister (2009), Cavalier et al. (2002) and the references therein) and a cornerstone of this kind of statistical issues is the crucial importance of the smoothness of the object to be recovered. A well known feature of nonparametric density estimation is the great influence of the “number of existing derivatives” of f on the statistical ability of any estimator (for example regarding the \mathbb{L}^2 -consistency rate of convergence). In particular, an imprecise knowledge on the smoothness parameter generally lead to inappropriate estimators.

If van Es et al. (2008) and Gugushvili et al. (2011) develop a nonparametric solution for estimating f and p , both works unfortunately assume the knowledge of the smoothness of f , which is clearly unrealistic in practice. Moreover, we have pointed out above that this smoothness has a crucial impact on the ability of the estimators. In particular, to obtain a tractable efficient statistical procedure, there is an important need to estimate also a smoothness class $\mathbf{H}_s(R)$ where the unknown density f is living (see below for details on the functional space $\mathbf{H}_s(R)$). Since the choice of the “good” smoothness parameter s may be a real issue from a practical point of view, there is an important need to obtain an estimator regardless the value of s is: we refer to *adaptive* estimators when they are able to adapt themselves to the unknown value of this regularity parameter. In the general settings of nonparametric regression and density estimation, such adaptive property may be attained by different strategies that are detailed below.

Adaptation In nonparametric statistics, estimators are importantly affected by a wrong choice of the parameters that are ideally smoothness-dependent. This is in particular the case for the bandwidth of a kernel deconvolution (see Fan (1991)) or when a frequency threshold in a Tychonov-type regularization is used (see Tsybakov (2009)) for example. Different strategies have been designed these last decades to attain *ad-hoc* adaptivity of estimators.

Among them, model selection theory introduced initially in Birgé and Massart (1998) produced a lot of interesting derivations in nonparametric and high-dimensional statistics, while the cornerstone of a tractable use of such an approach relies on a suitable minimal penalty calibration. Nevertheless, this calibration is sometimes difficult both from a theoretical and from a practical point of view and other methods can also be used to produce efficient adaptive estimators. For example, resampling methods with an additional cross-validation procedure (see *e.g.* Picard and Cook (1984)) is a popular method but we should notice that deriving theoretical results with this simple method requires a significant amount of work (see the recent contributions of Arlot and Lerasle (2016) for density estimation and of Arlot (2009) on general resampling methods).

In statistical signal processing, adaptivity of estimation may also be achieved with the help of a suitable decomposition basis and a thresholding strategy. This is what is done with wavelets on Besov balls on specific situations: introduced for density estimation in Donoho, Johnstone, Kerkyacharian, and Picard (1996) and used in different type of inverse problems in Bigot and Gadat (2010) and Bigot, Gadat, Klein, and Marteau (2013) for example.

Nevertheless, wavelets are not so easy to use with mixture problems and we propose to follow another possible guideline. We will use the so-called Lepskii method to derive adaptive estimators. This method introduced in Lepskii (1992) has been successfully applied in many nonparametric estimation problems and we refer to Goldenshluger and Lepskii (2011) and Lepskii (2015) for recent contributions using this method. We also refer to Chichignoud (2010) for an introduction of this method in different frameworks. The success of this method relies on a good bias-variance trade-off in the estimation procedure, and does not require a too much involved parameters tuning step.

Inverse problem with noisy operator At last, we point out that we have chosen to only address the situation where the density g is assumed to be known exactly. Such assumption may be seen as reasonable because it translates the fact that a biologist may repeat some fluorescence measurements on untreated cells a larger amount of time comparing to measurement on treated cells. Below, it will be translated by the use of the characteristic function of the random variable U denoted by Φ_U .

However, from a theoretical point of view, the effect of a preliminary estimation of the density g on the estimation of p and f will be certainly of interest, and is an open problem, left for a future subject of investigation in our current paper. When the operator described by Φ_U is unknown and has to be estimated from the data, then the problem falls into the framework of deconvolution with noise in the operator (see Cavalier (2011)). In such a case, it is highly suspected that the attainable rates may be damaged by the preliminary estimation of Φ_U . We refer to Cavalier and Hengartner (2005), Cavalier and Raimondo (2007) and Johannes (2009) and the references therein.

1.3 Main assumptions

Below, we will use i to refer to the complex number such that $i^2 = -1$ and the notation “:=” will introduce the definition of a mathematical object. We assume that the random variables U and X have a bounded second moment. The characteristic function of any random variable W will be denoted by Φ_W . Therefore, we will frequently use the following notations:

$$\forall t \in \mathbb{R} \quad \Phi_Z(t) := \mathbb{E}[e^{itZ}] \quad \text{and} \quad \Phi_U(t) := \mathbb{E}[e^{itU}] \quad \text{and} \quad \Phi_X(t) := \mathbb{E}[e^{itX}],$$

where (Z, U, X) are the random variables involved in Equation (1). We assume that U (resp. X) has a known density g (resp. unknown density f) with respect to the Lebesgue measure on \mathbb{R} . We will use for the sake of convenience the notation Φ_h as the Fourier transform of any density h :

$$\Phi_h(t) = \int_{\mathbb{R}} e^{itx} h(x) dx.$$

The notation \lesssim refers to an inequality up to a multiplicative constant independent of n .

It is well known that the Fourier transform of Z and its empirical counterpart may be used to obtain reliable estimations of Φ_X when dealing with a standard convolution inverse problem (see Fan (1991)) at the price of an assumption on the deconvolution operator translated in Φ_U . Moreover, we shall notice that our model includes the situation where $p = 0$, which turns our atomic deconvolution problem into the standard deconvolution problem. Therefore, it is expected that our estimation problem can be solved at the minimal price of some standard smoothness and invertibility conditions on the Fourier transform of U and some smoothness assumptions on f . Indeed, these kind of assumptions are well known in the inverse problem literature and commonly used to derive convergence rates of estimators (see among many references the work *e.g.* Fan and Koo (2002) where this assumption is used in its great generality, or Cavalier and Koo (2002) where this assumption is specified in the Poisson inverse problem situation). We also refer to Bigot and Gadat (2010); Bigot et al. (2012) for other applications of this kind of assumptions with different “deconvolution + mixture problems”. We are therefore driven to introduce two sets of densities $\mathbf{H}_s(R)$ and \mathbf{H}_ν^+ :

- The set $\mathbf{H}_s(R)$ denotes the set of densities f that belong to the Sobolev space of regularity s (and radius R) described with the help of the associated characteristic functions Φ_f such that:

$$\mathbf{H}_s(R) := \left\{ h \in \mathbb{L}^2(\mathbb{R}) : \int_{\mathbb{R}} |\Phi_h(t)|^2 (1 + |t|^{2s}) dt \leq R^2 \right\}$$

Below, the density of X denoted by f is assumed to belong to $\mathbf{H}_s(R)$ with an *unknown* parameter s , meaning that we assume that $\int_{\mathbb{R}} |\Phi_X(t)|^2 (1 + |t|^{2s}) dt \leq R^2$.

- The set \mathbf{H}_ν^+ denotes the set of densities such that the Fourier transform satisfies the smoothness and “invertibility” condition : $\exists (d_0, d_1) : 0 < d_0 < d_1$ and $d_0 |t|^{-\nu} \leq |\Phi(t)| \leq d_1 |t|^{-\nu}$ as $|t| \rightarrow +\infty$.

$$\mathbf{H}_\nu^+ := \{h \in \mathbb{L}^2(\mathbb{R}) : \exists 0 < d_0 < d_1 \quad d_0|t|^{-\nu} \leq |\Phi_h(t)| \leq d_1|t|^{-\nu}\}.$$

Below, we will assume that the density g of the random variable U belongs to \mathbf{H}_ν^+ .

Let us briefly comment on this last assumption. In Fan (1991), \mathbf{H}_ν^+ refers to the set of ordinary smooth densities of order ν , which includes many distributions as gamma, double exponential distributions. We also point out that it would be possible to address the super-smooth case with an exponential decrease of the Fourier transform to handle Gaussian or Cauchy densities. Nevertheless, we have chosen in this work to restrict our study to the ordinary smooth case for the sake of brevity. As pointed in Section 1.2, we assume that g is known, meaning that Φ_U is known on \mathbb{R} . Such an assumption is reasonable regarding the practical example we want to handle where we can repeat the experiments for the calibration of the cytometer many times.

The estimators introduced in van Es et al. (2008) exploit the knowledge of \mathbf{H}_ν^+ and of $\mathbf{H}_s(R)$ (which is more annoying for practical purpose) to obtain an optimal estimator with the important assumption of the knowledge of s (see Section 2.1).

1.4 Main results

In this article we introduce an adaptive procedure to estimate the unknown parameters p and f . We first consider the estimators \hat{p}_n and \hat{f}_n proposed in van Es et al. (2008) based on a kernel estimation using the relationship between the Fourier transform of each random variable because Equation (1) yields:

$$\forall t \in \mathbb{R} \quad \Phi_Z(t) = \Phi_U(t) [p + (1-p)\Phi_X(t)]. \quad (2)$$

We detail in Sections 2.1 and 3.1 the non-adaptive construction proposed by van Es et al. (2008). We then develop a well-designed strategy (see *e.g.* Lepskii (1992)) to obtain an adaptive estimator of p and f . This method **consists in** choosing among a grid of regularity, the associated bandwidth parameter that realizes the bias-variance tradeoff. The precise constructions of the adaptive estimator \hat{p}_{n,\hat{j}_n^p} and \hat{f}_{n,\hat{j}_n^f} are given in Section 2.2 and in Section 3.2. The aim of this article is to establish the consistency rate of our *adaptive* strategy and in particular its optimality (among a functional space for f). To assess this optimality, we handle the minimax risk paradigm and define the minimax risk for the estimation of p by:

$$r_n(p) := \inf_{\hat{p}_n} \sup_{p \in]0,1[, f \in \mathbf{H}_s(R)} \mathbb{E}_{\mathbb{P}_{p,f}} (\hat{p}_n - p)^2, \quad (3)$$

where the above infimum is taken over the set of all possible estimators constructed from (Z_1, \dots, Z_n) and where $\mathbb{E}_{\mathbb{P}_{p,f}}$ refers to the expectation over the n -observations i.i.d. according to our atomic deconvolution model described in Equation (1). For the sake of convenience, we will omit the subscripts $\mathbb{P}_{p,f}$ in the notation $\mathbb{E}_{\mathbb{P}_{p,f}}$ and simply denote by \mathbb{E} this expectation.

In the same time, we also introduce the minimax risk for the estimation of f defined by:

$$r_n(f) := \inf_{\hat{f}_n} \sup_{p \in]0,1[, f \in \mathbf{H}_s(R)} \mathbb{E}_{\mathbb{P}_{p,f}} \|\hat{f}_n - f\|_2^2, \quad (4)$$

The minimax risk $r_n(p)$ (resp. $r_n(f)$) represents the optimal rate of convergence that may be attained by any estimator of p (resp. of f) in the quadratic sense. We emphasize that the estimators proposed in van Es et al. (2008), Gugushvili et al. (2011) are minimax optimal and *not adaptive to the smoothness parameter s* (see Theorem 3 and Theorem 7).

Below, we will prove the following results. The first result concerns the estimation of p .

Theorem 1 (Minimax adaptivity for the estimation of p) *Assume that f belongs to $\mathbf{H}_s(R)$ and g to \mathbf{H}_ν^+ with $\nu > 1$. If $p \in (0, 1)$, then the estimator \hat{p}_{n,\hat{j}_n^p} defined in (12) and (15) satisfies:*

$$r_n(p) \lesssim \mathbb{E} \left(\hat{p}_{n,\hat{j}_n^p} - p \right)^2 \lesssim (\log n) r_n(p),$$

where $r_n(p) = n^{-(2s+1)/(2s+2\nu)}$.

The proof of Theorem 1 follows a multiple-testing strategy jointly used with a concentration inequality. The additional logarithmic term (regarding the minimax rate $n^{-(2s+1)/(2s+2\nu)}$) involved in our upper bound may be seen as the price to pay for using a multiple testing strategy and identifying the smoothness parameter.

Then we use a plug-in strategy to estimate the function f from its Fourier transform.

Theorem 2 (Minimax adaptivity for the estimation of f) *For any $p \in (0, 1)$, assume that Z has a finite variance and assume that $f \in \mathbf{H}_s(\mathbb{R})$ and $g \in \mathbf{H}_\nu^+$ with $\nu > 1$ and select \hat{j}_n^f as in Equation (31). Then, the estimator \hat{f}_{n, \hat{j}_n^f} defined in (32) satisfies:*

$$r_n(f) \lesssim \mathbb{E} \|\hat{f}_{n, \hat{j}_n^f} - f\|_2 \lesssim (\log n)^{a+1/2} r_n(f),$$

where $r_n(f) = n^{-s/(2s+2\nu+1)}$ and $a > 0$ is defined in Equation (28).

Hence, Theorem 1 and Theorem 2 show that our estimation procedures derived from (15) and (32) lead to optimal rates of convergence, that are of polynomial order with the sample size. This rate deteriorates as long as the degree of ill-posedness ν increases. We recover a standard behaviour in deconvolution situation. We also emphasize that our estimators for p and f , induced by the Lepskii rule, are non-asymptotic and fully-adaptive with respect to the smoothness parameter s . Therefore, our results produce an adaptive minimax optimal upper bound up to some logarithmic term. Finally, note that the lower bounds presented in Theorems 1 and 2 are already established in van Es et al. (2008) and Gugushvili et al. (2011) and are indicated here to assess the statistical efficiency of our adaptive estimation. Consequently, our theoretical contribution will be to provide the proof of the upper bounds involved in Theorems 1 and 2.

The article is organized as follows: Section 2 details the non-adaptive estimator of van Es et al. (2008) and then our adaptive estimation of the contamination parameter p while Section 3 proposes to solve the similar estimation problem for f . We present numerical simulations in Section 4 on both simulated and real datasets and the interest of our results for biological purposes. Finally, we gather some technical results in Appendix A, especially for the adaptive result on the estimation of f (see Subsection A.3).

2 Estimation of the contamination rate $1 - p$

We first consider the problem of estimating the contamination rate $1 - p$. We recall the non-adaptive results obtained by van Es et al. (2008) and expose our adaptive strategy afterwards.

2.1 Non adaptive estimation of p

We describe below the estimators proposed by van Es et al. (2008), that will be used to obtain our adaptive procedure. A key observation relies on Equation (2). This last equation makes it possible, from n independent observations $(Z_i)_{1 \leq i \leq n}$, to obtain an estimation of Φ_X and p . Given this set of observations, we first introduce the empirical estimator of the Fourier transform of Z :

$$\forall t \in \mathbb{R} \quad \hat{\Phi}_{Z,n}(t) := \frac{1}{n} \sum_{i=1}^n e^{itZ_i}.$$

Since f belongs to $\mathbf{H}_s(\mathbb{R})$, then $\lim_{|t| \rightarrow +\infty} \Phi_X(t) = 0$. Therefore, we expect to recover p by using the information brought by Φ_Z and Φ_U at large frequencies. In particular, Equation (2) implies the identifiability of the model and provides some insights for an estimation strategy. Then the knowledge of Φ_U and in particular the fact that the density of U belongs to \mathbf{H}_ν^+ , entails some lower bounds of Φ_U for large frequencies.

We use the construction of Gugushvili et al. (2011) and introduce a smooth real valued kernel Φ_K in the Fourier domain that satisfies:

$$\Phi_K(t) \neq 0 \iff t \in [-1; 1] \quad \text{and} \quad \int_{-1}^1 \Phi_K(t) dt = 2, \quad (5)$$

and a flatness condition on the neighborhood of 0 (\mathcal{V}_k below denotes an open neighborhood of 0):

$$\forall k \in \mathbb{R}_+, \quad \exists C_k > 0, \quad \exists \mathcal{V}_k \quad \forall t \in \mathcal{V}_k : \quad |\Phi_K(t)| \leq C_k |t|^k.$$

This last condition is not restrictive and is satisfied when Φ_K is chosen for example as:

$$\Phi_K(t) := \frac{e^{-a|t|^{-m}}}{C_{a,m}} \mathbf{1}_{|t| \leq 1},$$

for any $a > 0$, $m \geq 1$ ($C_{a,m}$ is the normalizing constant associated to Equation (5)). Note that this local condition may be replaced by a global one since Φ_K is bounded on $[-1; 1]$. Therefore, we keep the notation C_k and assume that:

$$\forall k \in \mathbb{R}_+, \quad \exists C_k > 0 \quad \forall t \in [-1; 1] : \quad |\Phi_K(t)| \leq C_k |t|^k. \quad (6)$$

Following the works of Fan (1991) and Gugushvili et al. (2011), we use the kernel Φ_K on the Fourier transform $p + (1-p)\Phi_X$ and obtain that:

$$\frac{h}{2} \int_{\mathbb{R}} \Phi_K(ht) \frac{\Phi_Z(t)}{\Phi_U(t)} dt = \frac{h}{2} \int_{\mathbb{R}} \Phi_K(ht) [p + (1-p)\Phi_X(t)] dt = p + \frac{h(1-p)}{2} \int_{-h^{-1}}^{h^{-1}} \Phi_K(ht) \Phi_X(t) dt.$$

The last term of the r.h.s. vanishes when $h \rightarrow 0$. Moreover, it is possible to obtain a tight upper bound in terms of h of this *bias* term:

$$\begin{aligned} \left| h \int_{-h^{-1}}^{h^{-1}} \Phi_K(ht) \Phi_X(t) dt \right| &= \left| h \int_{-h^{-1}}^{h^{-1}} \Phi_K(ht) |t|^{-s} \Phi_X(t) |t|^s dt \right| \\ &\leq h \sqrt{\int_{-h^{-1}}^{h^{-1}} |\Phi_K(ht)|^2 |t|^{-2s} dt \int_{\mathbb{R}} |\Phi_X(t)|^2 |t|^{2s} dt} \leq \sqrt{2} C_s R h^{s+1/2}, \end{aligned} \quad (7)$$

where we applied successively the Cauchy-Schwarz inequality, Inequality (6) with $k = s$ and the fact that $f \in \mathbf{H}_s(R)$. Therefore, we can deduce that:

$$\lim_{h \rightarrow 0} \frac{h}{2} \int_{\mathbb{R}} \Phi_K(ht) \frac{\Phi_Z(t)}{\Phi_U(t)} dt = p.$$

We finally plug Equation (2) in the above limit and define a natural estimator of p (that depends on a small bandwidth parameter h_n):

$$\hat{p}_n := \frac{h_n}{2} \int_{-h_n^{-1}}^{h_n^{-1}} \Phi_K(h_n t) \frac{\hat{\Phi}_{Z,n}(t)}{\Phi_U(t)} dt. \quad (8)$$

According to (7), we compute an upper-bound of the bias of \hat{p}_n as:

$$[\mathbb{E}\hat{p}_n - p]^2 \leq 2C_s^2 R^2 h_n^{2s+1}.$$

The variance of \hat{p}_n is handled in a standard way following the arguments of Fan (1991) with $\nu > 1$:

$$\mathbb{V}(\hat{p}_n) \lesssim \frac{h_n^2}{n} \int_{\mathbb{R}} \left| \int_{\mathbb{R}} e^{itz} \frac{\Phi_K(h_n t)}{\Phi_U(t)} dt \right|^2 dz \lesssim \frac{h_n}{n} \int_{\mathbb{R}} \left| \frac{\Phi_K(t)}{\Phi_U(th_n^{-1})} \right|^2 dt \lesssim \frac{1}{nh_n^{2\nu-1}}. \quad (9)$$

Now, a classical optimization of the bias-variance tradeoff yields the optimal (non-adaptive) choice for the bandwidth parameter $h_n^* := n^{-\frac{1}{2s+2\nu}}$. It can be shown that this choice leads to the minimax optimal consistency rate $n^{-(2s+1)/(2s+2\nu)}$. In other words, we have obtained the following result:

Theorem 3 (Theorem 1(i) and 5 in Gugushvili et al. (2011)) *Assume that $f \in \mathbf{H}_s(R)$ and $g \in \mathbf{H}_\nu^+$ with $\nu > 1$, then the choice $h_n = n^{-\frac{1}{2s+2\nu}}$ in (8) leads to an estimator \hat{p}_n that satisfies the consistency rate:*

$$\mathbb{E}[|\hat{p}_n - p|^2] \leq C(s, R) n^{-\frac{(2s+1)}{2s+2\nu}},$$

where $C(s, R)$ is a positive constant that continuously depend of s and R . Moreover, this estimator is minimax optimal under the additional assumption that $|\phi'(t)|(1 + |t|^\nu) \leq d_2$ for all $t \in \mathbb{R}$ and $d_2 > 0$.

This result highly depends on the knowledge of s and a bad strategy for the choice of s and h_n will significantly arms the estimation procedure. Nevertheless it is our starting point to produce an adaptive estimator of p .

2.2 Adaptive estimation of p

We propose to adapt the Lepskii strategy (see *e.g.* Lepskii (1992)) to obtain an adaptive estimator of p .

Grid on s We define a grid \mathcal{S}_n and will estimate s with an exploration of the possible values in \mathcal{S}_n . We consider an interval $[0, s_0]$ where we assume that $s \in [0, s_0]$ and define a regularly spaced sequence

$$s_{max} = s_0 > s_1 > \dots > s_{k_n} \geq 0,$$

such that for every $0 \leq j \leq k_n$,

$$s_j := s_0 - j \frac{\varepsilon}{\log n}. \quad (10)$$

In the last formula, $\varepsilon > 0$ is a fixed parameter. This parameter will be calibrated later on and will permit to obtain some good estimation properties. For each $0 \leq j \leq k_n$, we associate the bandwidth parameter $h_{n,j}$ that corresponds to the optimal bandwidth parameter chosen above when the smoothness of f is known and equal to s_j . This rule yields:

$$h_{n,j} := n^{-1/(2s_j+2\nu)}. \quad (11)$$

The corresponding estimator is then denoted by:

$$\hat{p}_{n,j} := 0 \vee \left(\frac{h_{n,j}}{2} \int_{-h_{n,j}^{-1}}^{h_{n,j}^{-1}} \Phi_K(h_n t) \frac{\hat{\Phi}_{Z,n}(t)}{\Phi_U(t)} dt \right) \wedge 1. \quad (12)$$

We choose to constraint the estimator to obtain a value in $[0, 1]$, which will not damage the properties of the estimator since $p \in [0, 1]$.

The associated minimax risk in the smoothness class $\mathcal{H}_{s_j}(R)$ is of the order $n^{-(s_j+1/2)/(2s_j+2\nu)}$ (see Theorem 3). Given a positive parameter β , we introduce a *penalty* term defined by this minimax risk up to a log term:

$$\kappa_{n,j} := \beta \sqrt{\log nn}^{-(s_j+1/2)/(2s_j+2\nu)}, \quad (13)$$

We should notice the important monotonic variations of the quantities defined above: when $j > l$ then $s_j < s_l$ and the bandwidth and penalty parameters satisfy $h_{n,j} < h_{n,l}$ and $\kappa_{n,j} > \kappa_{n,l}$. We introduce the notation s^* for the closest element to the regularity s from below in the grid \mathcal{G}_n :

$$s^* := \sup \{s_j \in \mathcal{G}_n : s_j \leq s\}. \quad (14)$$

For the sake of convenience, we denote by j^* the integer in $\{0, \dots, k_n\}$ such that $s^* = s_{j^*}$. This integer deterministically depends on n and ε , but we omit this dependence to alleviate the notations. The calibration of the grid yields

$$0 \leq s - s^* \leq \frac{\varepsilon}{\log n}.$$

Model selection with the Lepskii rule To construct the adaptive estimator, we use the following decision rule driven by the bias variance decomposition of each estimator of $\hat{p}_{n,j}$:

$$\hat{j}_n^p := \inf \{0 \leq j \leq k_n, \quad \forall l > j : |\hat{p}_{n,j} - \hat{p}_{n,l}| < \kappa_{n,l}\}. \quad (15)$$

The decision rule defined by (15) is costly from a computational point of view. However, some alternative coding strategies can be derived from Katkovnik (1999) and can significantly improve the efficiency of the computation of (15). These strategies rely on the construction of confidence intervals for the sequence $(\hat{p}_{n,j})_j$. We will see in Section 4 that if such a strategy works for the computation of a scalar parameter as p ; it is unfortunately no more possible for the estimation of a density.

The objective of the next paragraphs is to establish the consistency rate of this adaptive strategy, whose performance is described by Theorem 1. We first detail the Lepskii strategy, then give a concentration result and conclude.

2.2.1 Analysis of the Lepskii rule

We begin by the statement of two preliminary results. The first one links the average performance of \hat{p}_{n,\hat{j}_n^p} with some deviations inequalities.

Proposition 4 *Assume that $f \in \mathbf{H}_s(R)$ and g is a known density in \mathbf{H}_ν^+ . Then the estimator \hat{p}_{n,\hat{j}_n^p} defined with (12) and (15) satisfies*

$$\mathbb{E}|\hat{p}_{n,\hat{j}_n^p} - p|^2 \leq (\beta^2 \log(n) + C) e^{2\varepsilon/(2s+2\nu)} n^{-(2s+1)/(2s+2\nu)} + \sum_{l \geq j^*} \mathbb{P}\left(|\hat{p}_{n,l} - p| \geq \frac{\kappa_{n,l}}{2}\right).$$

Proof

We can decompose the estimation error as:

$$\mathbb{E}\left(\left|\hat{p}_{n,\hat{j}_n^p} - p\right|^2\right) = \mathbb{E}\left(\left|\hat{p}_{n,\hat{j}_n^p} - p\right|^2 \mathbf{1}_{\hat{j}_n^p \leq j^*} + \left|\hat{p}_{n,\hat{j}_n^p} - p\right|^2 \mathbf{1}_{\hat{j}_n^p > j^*}\right)$$

- We first consider the event $\{\hat{j}_n^p \leq j^*\}$ and apply the inequality $(a + b)^2 \leq 2a^2 + 2b^2$ to obtain:

$$\left|\hat{p}_{n,\hat{j}_n^p} - p\right|^2 \leq 2\left|\hat{p}_{n,\hat{j}_n^p} - \hat{p}_{n,j^*}\right|^2 + 2\left|\hat{p}_{n,j^*} - p\right|^2. \quad (16)$$

We study the first term of the r.h.s. of Equation (16). On the event $\{\hat{j}_n^p < j^*\}$, the definition of \hat{j}_n^p brought by Equation (15) yields:

$$\left|\hat{p}_{n,\hat{j}_n^p} - \hat{p}_{n,j^*}\right| < \kappa_{n,j^*},$$

and obviously when $\hat{j}_n^p = j^*$ the upper bound above also holds. Hence, we deduce that:

$$\mathbf{1}_{\hat{j}_n^p \leq j^*} \left|\hat{p}_{n,\hat{j}_n^p} - \hat{p}_{n,j^*}\right|^2 < \{\kappa_{n,j^*}\}^2.$$

Moreover, writing $s^* = s + (s^* - s)$ in the definition of the penalty (13) leads to:

$$\begin{aligned} \kappa_{n,j^*} &= \beta \sqrt{\log(n)} n^{-(s^*+1/2)/(2s^*+2\nu)} \\ &= \beta \sqrt{\log(n)} n^{-(s+1/2)/(2s+2\nu)} \exp\left[\left(\frac{s+1/2}{2s+2\nu} - \frac{s^*+1/2}{2s^*+2\nu}\right) \log n\right] \\ &\leq \beta \sqrt{\log(n)} n^{-(s+1/2)/(2s+2\nu)} \exp\left[\frac{2\nu \varepsilon \{\log n\}^{-1}}{(2s^*+2\nu)(2s+2\nu)} \log n\right] \\ &\leq \beta \exp\left[\frac{\varepsilon}{2s+2\nu}\right] \sqrt{\log(n)} n^{-(s+1/2)/(2s+2\nu)}. \end{aligned}$$

We can handle the second term of the r.h.s. of Equation (16) easily: the nonadaptive estimator obtained with j^* satisfies the upper bound obtained in Theorem 3, so that:

$$\begin{aligned} \mathbb{E}\left(\left|\hat{p}_{n,j^*} - p\right|^2\right) &\leq C n^{-(2s^*+1)/(2s^*+2\nu)} \\ &\leq C \exp\left(\frac{2\varepsilon}{2s+2\nu}\right) n^{-(2s+1)/(2s+2\nu)}, \end{aligned}$$

where C is a positive constant independent from s , n , and ε (the constant C only depends on R and s_0). Then,

$$\mathbb{E}\left(\mathbf{1}_{\hat{j}_n^p \leq j^*} \left|\hat{p}_{n,\hat{j}_n^p} - p\right|^2\right) \leq (\beta^2 \log(n) + C) \exp\left(\frac{2\varepsilon}{2s+2\nu}\right) n^{-(2s+1)/(2s+2\nu)}. \quad (17)$$

- We now consider the complementary event $\{\hat{j}_n^p > j^*\}$ and remark that

$$\hat{j}_n^p > j^* \iff \forall j \leq j^* \exists l > j : |\hat{p}_{n,j} - \hat{p}_{n,l}| \geq \kappa_{n,l}.$$

Therefore, we deduce that

$$\begin{aligned} \{\hat{j}_n^p > j^*\} &= \bigcap_{j \leq j^*} \bigcup_{l > j} \left\{ |\hat{p}_{n,j} - \hat{p}_{n,l}| \geq \kappa_{n,l} \right\} \\ &\subset \bigcup_{l > j^*} \left\{ |\hat{p}_{n,j^*} - \hat{p}_{n,l}| \geq \kappa_{n,l} \right\} \\ &\subset \bigcup_{l > j^*} \left\{ |\hat{p}_{n,l} - p| \geq \frac{\kappa_{n,l}}{2} \right\} \bigcup_{l > j^*} \left\{ |\hat{p}_{n,j^*} - p| \geq \frac{\kappa_{n,l}}{2} \right\}. \end{aligned} \quad (18)$$

where the second line comes from the inclusion $\bigcap_{j \leq j^*} \{\cdot\} \subset \bigcap_{j=j^*} \{\cdot\}$ and the last line from the triangle inequality. Since the map $l \mapsto \kappa_{n,l}$ is increasing then when $l > j^*$, we have $\kappa_{n,l} \geq \kappa_{n,j^*}$ so that

$$\bigcup_{l > j^*} \left\{ |\hat{p}_{n,j^*} - p| \geq \frac{\kappa_{n,l}}{2} \right\} \subset \left\{ |\hat{p}_{n,j^*} - p| \geq \frac{\kappa_{n,j^*}}{2} \right\}.$$

Using this last inclusion in the second union of the right hand side of (18), we finally deduce that

$$\{\hat{j}_n^p > j^*\} \subset \bigcup_{l \geq j^*} \left\{ |\hat{p}_{n,l} - p| \geq \frac{\kappa_{n,l}}{2} \right\}.$$

From the obvious upper-bound $|\hat{p}_{n,\hat{j}_n^p} - p|^2 \leq 1$, we conclude that:

$$\mathbb{E}(|\hat{p}_{n,\hat{j}_n^p} - p|^2 \mathbf{1}_{\hat{j}_n^p > j^*}) \leq \mathbb{E}(\mathbf{1}_{\hat{j}_n^p > j^*}) \leq \sum_{l \geq j^*} \mathbb{P}\left(|\hat{p}_{n,l} - p| \geq \frac{\kappa_{n,l}}{2}\right). \quad (19)$$

Gathering Equation (17) and Equation (19) leads to the desired conclusion. \square

2.2.2 Concentration inequality

The adaptivity property of the estimator \hat{p}_{n,\hat{j}_n^p} will be deduced from the multiple-testing strategy induced by Proposition 5. A baseline property to successively apply this approach will be a derivation of a concentration inequality on $\hat{p}_{n,l} - p$ for each fixed l . Next proposition states that such a concentration inequality holds.

Proposition 5 *Let $f \in \mathbf{H}_s(R)$, g a known density in \mathbf{H}_ν^+ with $\nu > 1/2$, and $l \geq j^*$, then*

$$\mathbb{P}\left(|\hat{p}_{n,l} - p| \geq \frac{\kappa_{n,l}}{2}\right) \leq 2n^{-\beta^2/64}.$$

Proof

The proof is divided into two steps.

Step 1: reduction to a concentration inequality. For any $l \geq j^*$ and $n \in \mathbb{N}$, we write

$$\begin{aligned} \hat{p}_{n,l} - p &= \frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \frac{\hat{\Phi}_{Z,n}(t)}{\Phi_U(t)} dt - p \\ &= \frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \frac{\hat{\Phi}_{Z,n}(t) - \Phi_Z(t)}{\Phi_U(t)} dt + \frac{h_{n,l}^{-1}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \frac{\Phi_Z(t)}{\Phi_U(t)} dt - p \\ &= \frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \frac{\hat{\Phi}_{Z,n}(t) - \Phi_Z(t)}{\Phi_U(t)} dt + \frac{(1-p)h_{n,l}^{-1}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \Phi_X(t) dt. \end{aligned}$$

The triangle inequality and the bias upper bound provided by Equation (7) yield:

$$|\hat{p}_{n,l} - p| \leq \left| \frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \frac{\hat{\Phi}_{Z,n}(t) - \Phi_Z(t)}{\Phi_U(t)} dt \right| + C_s R h_{n,l}^{s+1/2}.$$

The r.h.s. of the upper bound is of the order $h_{n,l}^{s+1/2} = n^{-(s+1/2)/(2s_l+2\nu)}$ while $\kappa_{n,l} = \beta \sqrt{\log nn}^{-(s_l+1/2)/(2s_l+2\nu)}$. From the definition of the grid on $s, l \geq j^*$ implies that $s_l \leq s_{j^*}$ so that $s_l \leq s$. Therefore, we can write:

$$h_{n,l}^{s+1/2} \{\kappa_{n,l}\}^{-1} \lesssim \{\log n\}^{-1/2} n^{(s_l-s)/(2s_l+2\nu)} \lesssim \frac{1}{\sqrt{\log n}}.$$

We then deduce that a sufficiently large n exists such that for $l \geq j^*$, we have $C_s R h_{n,l}^{s+1/2} < \kappa_{n,l}/4$ and

$$\begin{aligned} \mathbb{P}\left(|\hat{p}_{n,l} - p| \geq \frac{\kappa_{n,l}}{2}\right) &\leq \mathbb{P}\left(\left|\frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \frac{\hat{\Phi}_{Z,n}(t) - \Phi_Z(t)}{\Phi_U(t)} dt\right| \geq \frac{\kappa_{n,l}}{4}\right) \\ &\leq \mathbb{P}\left(\left|\frac{1}{n} \sum_{k=1}^n \underbrace{\frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \frac{e^{iZ_k t} - \Phi_Z(t)}{\Phi_U(t)} dt}_{:=\xi_{k,l}}\right| \geq \frac{\kappa_{n,l}}{4}\right). \end{aligned} \quad (20)$$

Step 2: application of the Bernstein concentration inequality. To handle the r.h.s. of Inequality (20), we will apply the Bernstein inequality to the random variables

$$\xi_{k,l} = \frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \Phi_K(h_{n,l}t) \frac{e^{iZ_k t} - \Phi_Z(t)}{\Phi_U(t)} dt,$$

(see Theorem 2.9 of Boucheron, Lugosi, and Massart (2013) and exercise 2.8 therein, a precise statement is given in Theorem 9 of our Appendix A). The i.i.d. random variables $(\xi_{k,l})_{1 \leq k \leq n}$ are centered and satisfy an almost sure bound:

$$\forall k \in \{1 \dots n\} \quad |\xi_{k,l}| \leq \frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \frac{|\Phi_K(h_{n,l}t)|}{|\Phi_U(t)|} dt.$$

We now use the assumption that $g \in \mathbf{H}_\nu^+$ to deduce that a d_0 exists such that $\Phi_U(t)(1+|t|)^\nu \geq d_0$ for all $t \in \mathbb{R}$. Therefore, a large enough constant C exists such that for any $k \in \{1 \dots n\}$:

$$\begin{aligned} |\xi_{k,l}| &\leq \frac{h_{n,l}}{2} \int_{-h_{n,l}^{-1}}^{h_{n,l}} \frac{|\Phi_K(h_{n,l}t)|(1+|t|)^\nu}{|\Phi_U(t)|(1+|t|)^\nu} dt \\ &\leq \frac{h_{n,l}}{2d_0} \int_{-h_{n,l}^{-1}}^{h_{n,l}} |\Phi_K(h_{n,l}t)|(1+|t|)^\nu dt \\ &\leq (1+|h_{n,l}^{-1}|)^\nu \frac{h_{n,l}}{2d_0} \int_{-h_{n,l}^{-1}}^{h_{n,l}} |\Phi_K(h_{n,l}t)| dt \leq C h_{n,l}^{-\nu}. \end{aligned}$$

We can also bound the variance of each $\xi_{k,l}$ and a computation similar to Equation (9) leads to $\mathbb{V}(\xi_{k,l}) \leq h_{n,l}^{-2\nu+1}$. We then apply Theorem 9 with $b = C h_{n,l}^{-\nu}$ and $v = n h_{n,l}^{-2\nu+1}$ and deduce that:

$$\mathbb{P}\left(\left|\frac{1}{n} \sum_{k=1}^n \xi_{k,l}\right| \geq \frac{\kappa_{n,l}}{4}\right) \leq 2 \exp\left(-\frac{n \kappa_{n,l}^2}{32 (h_{n,l}^{-2\nu+1} + C h_{n,l}^{-\nu} \kappa_{n,l}/3)}\right).$$

Now, remark that $h_{n,l}^{-2\nu+1} = n^{(2\nu-1)/(2\nu+2s_l)}$ while $h_{n,l}^{-\nu} \kappa_{n,l} = \beta n^\nu / (2\nu+2s_l) n^{-(s_l+1/2)/(2\nu+2s_l)} \sqrt{\log n}$. Since we assumed that $\nu > 1/2$, the main contribution in the denominator of the exponential bound above is brought by $h_{n,l}^{-2\nu+1}$. We therefore conclude that a large enough n exists such that:

$$\mathbb{P}\left(\left|\frac{1}{n} \sum_{k=1}^n \xi_{k,l}\right| \geq \frac{\kappa_{n,l}}{4}\right) \leq 2 \exp\left(-\frac{n \kappa_{n,l}^2}{64 h_{n,l}^{-2\nu+1}}\right) = 2n^{-\beta^2/64},$$

because the penalty term $\kappa_{n,l}$ has been designed so that $\frac{n\kappa_{n,l}^2}{h_n^{2\nu+1}} = \beta^2 \log(n)$. \square

2.2.3 Proof of Theorem 1

We can achieve the proof of Theorem 1 and establish the adaptation of our estimator \hat{p}_{n,\hat{j}_n^p} : Propositions 4 and 5 yield

$$\mathbb{E}|\hat{p}_{n,\hat{j}_n^p} - p|^2 \leq (\beta^2 \log(n) + C) \exp\left(\frac{2\varepsilon}{2s+2\nu}\right) n^{-(2s+1)/(2s+2\nu)} + 2 \sum_{l \geq j^*} n^{-\beta^2/64}.$$

The size of the grid is bounded from above by $\varepsilon^{-1} s_{max} \log n$. Therefore, for n large enough

$$\mathbb{E}|\hat{p}_{n,\hat{j}_n^p} - p|^2 \leq (\beta^2 + 1) e^{\frac{2\varepsilon}{2s+2\nu}} (\log n) n^{-(2s+1)/(2s+2\nu)} + \varepsilon^{-1} s_{max} \log n n^{-\beta^2/64}.$$

It remains to choose the constant ε and β . We first pick β such that

$$n^{-\beta^2/64} \leq n^{-(2s+1)/(2s+2\nu)}, \quad (21)$$

then we choose ε in order to minimize the constant, regardless the value of s ,

$$(\beta^2 + 1) e^{\frac{2\varepsilon}{2s+2\nu}} + s_{max} \varepsilon^{-1}. \quad (22)$$

These choices yield the desired adaptive property (optimal up to a log term):

$$\mathbb{E}|\hat{p}_{n,\hat{j}_n^p} - p|^2 \lesssim (\log n) n^{-(2s+1)/(2s+2\nu)}.$$

\square

Remark 6 For numerical experiments, the calibration of the parameters ε and β given by (21) and (22) is crucial. Since the value of s is unknown, these inequality have to be true for all $s \in [0, s_{max}] = [0, s_0]$. We are driven to an optimal choice of the form

$$\beta^* = 8 \sqrt{\frac{2s_{max} + 1}{2s_{max} + 2\nu}} \text{ and } \varepsilon^* = \operatorname{argmin} \{((\beta^*)^2 + 1) e^{\frac{\varepsilon}{\nu}} + s_{max} \varepsilon^{-1}\}.$$

3 Estimation of f

We first briefly describe the estimator proposed by Gugushvili et al. (2011) and then detail our adaptive procedure.

3.1 Non adaptive approach

The estimation of f is highly similar to the one of p and relies on the Fourier transform of U and of the empirical data. It uses the relationship:

$$\Phi_X(t) = \frac{\Phi_Z(t) - p\Phi_U(t)}{(1-p)\Phi_U(t)},$$

and therefore uses a plug-in estimator of p and of Φ_Z with another kernel smoothing strategy. Following Gugushvili et al. (2011), we introduce a second kernel Q whose Fourier transform is denoted by Φ_Q has its support in $[-1, 1]$ and satisfies the following assumptions:

$$\Phi_Q(0) = 1, \quad \forall t \in \mathbb{R} \quad |\Phi_Q(t) - 1| \leq M|t|^s, \quad \int_{-1}^1 |\Phi_Q(t)|^2 dt < \infty, \quad (23)$$

where $M > 0$. It is worth saying that the choice of the kernel used for this deconvolution is important regarding the numerical results as well as the theoretical ones. For our purpose, we only handle ordinary smooth inverse problem in Sobolev spaces. In that case, it will be enough to handle a very simple kernel Φ_Q , given by the sinc function:

$$\Phi_Q(t) := \mathbf{1}_{[-1;1]} \quad \text{so that} \quad \forall x \in \mathbb{R} \quad Q(x) = \frac{\sin x}{\pi x} \quad \text{with} \quad Q(0) = \frac{1}{\pi}.$$

It is immediate to check that this kernel automatically matches the requirements given in Equation (23) on the smoothness of Φ_Q around 0. We refer to Delaigle and Hall (2006) for a detailed discussion on the influence of the kernel choice from a numerical point of view, and to Comte and Lacour (2013) and Goldenshluger and Lepskii (2011) for deeper insights on functional spaces where the estimation is done (*e.g.* in anisotropic Nikol'skii classes). The plug-in strategy proposed in Gugushvili et al. (2011) reads as follows:

$$\hat{\Phi}_{X,n} = \frac{\hat{\Phi}_{Z,n}(t) - \check{p}_n \Phi_U(t)}{(1 - \check{p}_n) \Phi_U(t)} \Phi_Q(\delta_n t), \quad (24)$$

where \check{p}_n is a preliminary truncated estimation of p that is plugged into Equation 24, with the following rule:

$$\check{p}_n := \hat{p}_n \wedge 1 - \tau_n. \quad (25)$$

The truncation step under $1 - \tau_n$ makes it possible to avoid numerical instability when \hat{p}_n is estimated close to 1. It is important to remark that for a theoretical purpose, it is necessary to build a preliminary estimator of p with \check{p}_n independent from the estimated Φ_Z in (24). This can be achieved using $\lfloor n/2 \rfloor$ samples to compute \check{p}_n and the remaining ones to build $\hat{\Phi}_{X,n}$ according to our plug-in strategy. We can now state the minimax consistency result of Gugushvili et al. (2011), which is obtained with a similar bias-variance tradeoff.

Theorem 7 (Theorem 2(i) and 3(i) in Gugushvili et al. (2011)) *Assume $f \in \mathbf{H}_s(R)$ and $g \in \mathbf{H}_\nu^+$ with $\nu > 1$ and set $\tau_n = \log(3n)^{-1}$. Then the choice $h_n = n^{-\frac{1}{2s+2\nu}}$ in (8) and $\delta_n = n^{-1/(2s+2\nu+1)}$ in (24) leads to an estimator \hat{f}_n that satisfies the consistency rate:*

$$\mathbb{E}[\|\hat{f}_n - f\|^2] \lesssim n^{-\frac{(2s)}{2s+2\nu+1}}.$$

It is shown in Gugushvili et al. (2011) that such estimator is minimax optimal under the additional assumptions that $s \geq 1/2$ and $|\phi'(t)|(1 + |t|^\nu) \leq d_2$ for all $t \in \mathbb{R}$ and $d_2 > 0$. In particular, we note that the consistency rate corresponds to the standard rate of ordinary smooth deconvolution inverse problem (see *e.g.* Fan (1991)). Again, we stress the fact that the proposed estimator highly depends on s through $\delta_n = n^{-1/(2s+2\nu+1)}$, which is unknown in practice and should be estimated. The next paragraph describes our proposed strategy.

3.2 Adaptive estimation of f

3.2.1 Grid on s

The adaptation follows the same strategy as the one for p , even though significantly harder from a theoretical point of view. We still use the same grid \mathcal{S}_n : $s_0 > s_1 > \dots > s_{k_n} \geq 0$, such that for every $0 \leq j \leq k_n$, $s_j = s_0 - j \frac{\varepsilon}{\log n}$, and the notation $h_{n,j}$ for the bandwidth parameter associated to the optimal estimation of p in $\mathbf{H}_s(R)$ when $s = s_j$, *i.e.* we use $h_{n,j} := n^{-1/(2s_j+2\nu)}$. A second bandwidth parameter associated to the estimation of f is defined by $\delta_{n,j}$, $\forall 0 \leq j \leq k_n$:

$$\delta_{n,j} := n^{-1/(2s_j+2\nu+1)}. \quad (26)$$

With these two bandwidth parameters derived from a regularity s_j , we build a preliminary estimator of p with the observations $(Z_i)_{1 \leq i \leq \lfloor n/2 \rfloor}$:

$$\hat{p}_{n,j}^{(1)} = 0 \vee \left\{ \left\{ \frac{h_{n,j}}{2} \int_{-h_{n,j}^{-1}}^{h_{n,j}^{-1}} \Phi_K(h_{n,j}t) \frac{\hat{\Phi}_{Z,n}^{(1)}(t)}{\Phi_U(t)} dt \right\} \wedge 1 - \tau_n \right\} \quad (27)$$

where $\hat{\Phi}_{Z,n}^{(1)}(t)$ and $\hat{\Phi}_{Z,n}^{(2)}(t)$ are two unbiased *independent* estimates of $\Phi_Z(t)$:

$$\forall t \in \mathbb{R} \quad \hat{\Phi}_{Z,n}^{(1)}(t) := \frac{1}{\lfloor n/2 \rfloor} \sum_{k=1}^{\lfloor n/2 \rfloor} e^{itZ_k} \quad \text{and} \quad \hat{\Phi}_{Z,n}^{(2)}(t) := \frac{1}{n-1-\lfloor n/2 \rfloor} \sum_{k=\lfloor n/2 \rfloor+1}^n e^{itZ_k}.$$

Theorem 7 entails for a good choice of the truncation parameter introduced in Equation (25) and we choose:

$$\tau_n := \log(n)^{-a}, \quad (28)$$

for $a > 0$. Then, we use $\hat{p}_{n,j}^{(1)}$ to obtain an estimator of the empirical Fourier transforms of X by:

$$\hat{\Phi}_{X_{n,j}}^{(2)} := \frac{\hat{\Phi}_{Z,n}^{(2)}(t) - \hat{p}_{n,j}^{(1)}\Phi_U(t)}{(1 - \hat{p}_{n,j}^{(1)})\Phi_U(t)} \Phi_Q(\delta_{n,j}t) \quad (29)$$

The application of the Lepskii method relies on a penalty term, which is defined as:

$$\rho_{n,l} := \beta(\log n)^{a+1/2} n^{-s_l/(2s_l+2\nu+1)}. \quad (30)$$

This penalty is the minimax risk of estimation of f in $\mathbf{H}_s(R)$ when $s_l = s$, up to a log term (see Theorem 7). We recall that for the estimation of p , the penalty was associated to a supplementary $(\log n)^{1/2}$ term. However, such a penalty term is not strong enough to obtain a good concentration inequality for the estimator of f (see Proposition 13 below).

3.2.2 Lepskii's rule for the estimation of f

We use the penalization introduced in (30) to define:

$$\hat{j}_n^f := \inf\{0 \leq j \leq k_n, \quad \forall l > j : \|\hat{f}_{n,j} - \hat{f}_{n,l}\|_2 < \rho_{n,l}\}.$$

Thanks to the Parseval identity, this can be also formulated in terms of the Fourier transforms:

$$\hat{j}_n^f := \inf\{0 \leq j \leq k_n, \quad \forall l > j : \|\hat{\Phi}_{X_{n,j}}^{(2)} - \hat{\Phi}_{X_{n,l}}^{(2)}\|_2 < \rho_{n,l}\}. \quad (31)$$

The estimator derived from the penalization above is then written as $\hat{\Phi}_{X_{n,\hat{j}_n^f}}^{(2)}$ where we used the selection of \hat{j}_n^f in (31) with the definition of $\hat{\Phi}_{X_{n,\hat{j}_n^f}}^{(2)}$ given in (29). It leads to the estimation of f itself using the Fourier reconstruction formula:

$$\hat{f}_{n,\hat{j}_n^f}(x) := \frac{1}{2\pi} \int_{\mathbb{R}} e^{-itx} \hat{\Phi}_{X_{n,\hat{j}_n^f}}^{(2)}(t) dt. \quad (32)$$

Again, we start with the statement of a proposition that links the average performance of $\hat{\Phi}_{X_{n,\hat{j}_n^f}}^{(2)}$ with a family of deviation inequalities.

Proposition 8 *Let $f \in \mathbf{H}_s(R)$ and g a known density in \mathbf{H}_ν^+ , then the estimator \hat{f}_{n,\hat{j}_n^f} satisfies:*

$$\mathbb{E}\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \lesssim (\log n)^{a+1/2} n^{-s/(2s+2\nu+1)} + \sqrt{\sum_{l \geq j^*} \mathbb{P}\left(\|\hat{f}_{n,l} - f\|_2 \geq \frac{\rho_{n,l}}{2}\right)}.$$

Proof

The proof is close to the one of Proposition 4. We denote by $s^* = s_{j^*}$ the closest element from s from below in the grid \mathcal{S}_n (see Equation (14)). Following the same guidelines, we obtain that:

$$\mathbb{E}\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 = \mathbb{E}\left(\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \mathbf{1}_{\hat{j}_n^f \leq j^*} + \|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \mathbf{1}_{\hat{j}_n^f > j^*}\right).$$

On the event $\{\hat{j}_n^f \leq l^*\}$: we first apply the triangle inequality:

$$\mathbb{E}(\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \mathbf{1}_{\hat{j}_n^f \leq j^*}) \leq \mathbb{E}\left[\left(\|\hat{f}_{n,\hat{j}_n^f} - \hat{f}_{j^*}\|_2 + \|\hat{f}_{j^*} - f\|_2\right) \mathbf{1}_{\hat{j}_n^f \leq j^*}\right]$$

On $\hat{j}_n^f \leq j^*$, the first term is upper bounded by:

$$\begin{aligned} \rho_{n,j^*} &= \beta(\log n)^{a+1/2} n^{-s_{j^*}/(2s_{j^*}+2\nu+1)} \\ &= \beta(\log n)^{a+1/2} n^{-s/(2s+2\nu+1)} \exp\left[\left(\frac{s}{2s+2\nu+1} - \frac{s_{j^*}}{2s_{j^*}+2\nu+1}\right) \log n\right] \\ &\leq \beta(\log n)^{a+1/2} n^{-s/(2s+2\nu+1)} \exp\left(\frac{\varepsilon}{2s+2\nu+1}\right). \end{aligned}$$

For the second term, we apply the results of the non adaptive estimator in Theorem 7:

$$\mathbb{E}\left(\|\hat{f}_{j^*} - f\|_2 \mathbf{1}_{\hat{j}_n^f \leq j^*}\right) \leq C n^{-s_{j^*}/(2s_{j^*}+2\nu+1)},$$

and deduce that:

$$\mathbb{E}(\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \mathbf{1}_{\hat{j}_n^f \leq j^*}) \leq \left(\beta(\log n)^{a+1/2} + C\right) n^{-s/(2s+2\nu+1)} \exp\left(\frac{\varepsilon}{2s+2\nu+1}\right). \quad (33)$$

where C is a positive constant independent from β , ε and n .

On the event $\{\hat{j}_n^f > l^*\}$: On this event, the triangle inequality yields $\|\hat{f}_{n,l} - \hat{f}_{n,j}\|_2 \leq \|\hat{f}_{n,l} - f\|_2 + \|\hat{f}_{n,j} - f\|_2$, so that

$$\{\hat{j}_n^f > j^*\} \subset \bigcup_{l \geq j^*} \left\{ \|\hat{f}_{n,l} - f\|_2 \geq \frac{\rho_{n,l}}{2} \right\}.$$

Therefore, the Cauchy-Schwarz inequality and the inclusion above yields:

$$\begin{aligned} \mathbb{E}(\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \mathbf{1}_{\hat{j}_n^f > j^*}) &\leq \sqrt{\mathbb{E}(\|\hat{f}_{n,\hat{j}_n^f} - f\|_2^2)} \sqrt{\mathbb{E}(\mathbf{1}_{\hat{j}_n^f > j^*})} \\ &\leq \sqrt{\sum_{j=0}^{k_n} \mathbb{E}(\|\hat{f}_{n,j} - f\|_2^2)} \sqrt{\mathbb{E}\left(\mathbf{1}_{\bigcup_{l \geq j^*} \left\{ \|\hat{f}_{n,l} - f\|_2 \geq \frac{\rho_{n,l}}{2} \right\}}\right)}. \end{aligned}$$

We now use a refinement of Theorem 7 given by Proposition 10 in the appendix Section A to bound $\mathbb{E}(\|\hat{f}_{n,j} - f\|_2^2)$:

$$\mathbb{E}(\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \mathbf{1}_{\hat{j}_n^f > j^*}) \leq \sqrt{\sum_{j=0}^{k_n} \Phi(s, R) n^{-2 \min(s_j, s)/(2s_j+2\nu+1)}} \sqrt{\mathbb{E}\left(\mathbf{1}_{\bigcup_{l \geq j^*} \left\{ \|\hat{f}_{n,l} - f\|_2 \geq \frac{\rho_{n,l}}{2} \right\}}\right)}$$

Since $k_n \leq s_0 \varepsilon^{-1} \log(n)$ and Φ is continuous in s , then $C = \{\sup_{s \in [0, s_0]} \Phi(s, R)\}^{1/2} \sqrt{s_0 \varepsilon^{-1}}$ is a finite constant. Setting now $\alpha = s_0 \varepsilon^{-1}$, we shall now study the next sum:

$$\sum_{j=0}^{k_n} n^{-s_j/(s_j+\nu+1/2)} = \sum_{j=0}^{\alpha \log(n)} n^{-\frac{j\varepsilon/\log(n)}{j\varepsilon/\log(n)+\nu+1/2}}.$$

This sum may be divided into two parts: one with $s_j \leq 1$ and the other part with smoothness $s_j \geq 1$. If $j\varepsilon/\log(n) \geq 1$, then we use the rough bound $s_j/(s_j + \nu + 1/2) \geq 1/(\nu + 3/2)$ and obtain that:

$$\sum_{j=\varepsilon^{-1} \log(n)}^{k_n} n^{-s_j/(s_j+\nu+1/2)} \leq \alpha \log(n) n^{-1/(\nu+3/2)},$$

which is a bounded sequence. Now, if $j\epsilon/\log(n) \leq 1$, we have:

$$\sum_{j=0}^{\epsilon^{-1} \log(n)} n^{-\frac{j\epsilon/\log(n)}{j\epsilon/\log(n)+\nu+1/2}} \leq \sum_{j=0}^{\epsilon^{-1} \log(n)} e^{-\frac{j\epsilon}{j\epsilon/\log(n)+\nu+1/2}} \leq \sum_{j=0}^{\epsilon^{-1} \log(n)} e^{-\frac{\epsilon}{\nu+3/2}j} \leq \frac{1}{1 - e^{-\frac{\epsilon}{3/2+\nu}}}.$$

Therefore, a constant $C(\epsilon, s_0, R)$ exists such that:

$$\mathbb{E}(\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \mathbf{1}_{\hat{j}_n^f > j^*}) \leq C(\epsilon, s_0, R) \sqrt{\sum_{l \geq j^*} \mathbb{P}\left(\|\hat{f}_{n,l} - f\|_2 \geq \frac{\rho_{n,l}}{2}\right)}. \quad (34)$$

The conclusion follows from (33) and (34). \square

3.2.3 Proof of Theorem 2

Proposition 8 and Proposition 12 (see Appendix A, paragraph A.3) yield:

$$\begin{aligned} \mathbb{E}\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 &\lesssim (\log n)^{a+1/2} n^{-s/(2s+2\nu+1)} + \sqrt{\sum_{m \geq j^*} \mathbb{P}\left(\|\hat{f}_{n,m} - f\|_2 \geq \frac{\rho_{n,m}}{2}\right)} \\ &\lesssim (\log n)^{a+1/2} n^{-s/(2s+2\nu+1)} + \sqrt{\sum_{m \geq j^*} \mathbb{P}\left(\|\hat{\Phi}_{X_{n,l}}^{(2)} - \mathbb{E}\hat{\Phi}_{X_{n,l}}^{(2)}\|_2^2 \geq \frac{\rho_{n,l}^2}{8}\right)} \end{aligned}$$

From the definition of the grid \mathcal{S}_n (see the beginning of Section 3.2), it contains a number of order $s_0 \epsilon^{-1} \log(n)$ points and for all l , $n^{-\frac{3/2+s_l+\nu}{2s_l+2\nu+1}} \leq n^{-1/2}$. Then, we conclude from Proposition 13 (see Appendix A, paragraph A.3) that:

$$\begin{aligned} \mathbb{E}\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 &\lesssim \beta (\log n)^{a+1/2} n^{-s/(2s+2\nu+1)} \\ &+ \sqrt{s_0 \epsilon^{-1} \log n \left(n^{-\frac{\beta^2}{64}} + n^{-1} + (\log n)^{-1/2} n^{-\frac{\beta^2 d_2^2}{24 \|\Phi_Q\|_2^2} + \frac{3/2+\nu+s_l}{2\nu+2s_l+1}} \right)} \end{aligned}$$

Therefore, for β large enough:

$$\mathbb{E}\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \lesssim (\log n)^{a+1/2} n^{-s/(2s+2\nu+1)},$$

which ends the proof of the almost optimal (up to some log terms) adaptivity of our Lepskii's rule. \square

4 Numerical experiments

This section presents our numerical investigations for solving the adaptive atomic deconvolution problem modeled by Equation (1). We first detail in Section 4.1 some simulations on simulated synthetic dataset where both the targeted density f and the targeted proportion p are known. Then, Section 4.2 studies a real original dataset obtained through flow cytometer measurements and derive some statistical conclusions from our analysis.

Below, our nonparametric estimation of the density f will be obtained by considering the real positive part of \hat{f}_{n,\hat{j}_n^f} . In particular, we estimate f with

$$\widetilde{\hat{f}}_{n,\hat{j}_n^f} := \max\left(\Re e\left(\hat{f}_{n,\hat{j}_n^f}\right), 0\right).$$

We emphasize that $\widetilde{\hat{f}}_{n,\hat{j}_n^f}$ possesses a lower quadratic risk than \hat{f}_{n,\hat{j}_n^f} thanks to the following arguments:

$$\begin{aligned}
|\widetilde{\hat{f}}_{n,\hat{j}_n^f}(x) - f(x)| &= |\widetilde{\hat{f}}_{n,\hat{j}_n^f}(x) - f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) > 0} + |\widetilde{\hat{f}}_{n,\hat{j}_n^f}(x) - f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) < 0} \\
&= |\Re(\hat{f}_{n,\hat{j}_n^f})(x) - f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) > 0} + |f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) < 0} \\
&\leq |\hat{f}_{n,\hat{j}_n^f}(x) - f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) > 0} + |f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) < 0} \\
&\leq |\hat{f}_{n,\hat{j}_n^f}(x) - f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) > 0} + |\Re(\hat{f}_{n,\hat{j}_n^f})(x) - f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) < 0} \\
&\leq |\hat{f}_{n,\hat{j}_n^f}(x) - f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) > 0} + |\hat{f}_{n,\hat{j}_n^f}(x) - f(x)|\mathbf{1}_{\Re(\hat{f}_{n,\hat{j}_n^f}) < 0} \\
&= |\hat{f}_{n,\hat{j}_n^f}(x) - f(x)|.
\end{aligned}$$

Therefore, all our simulations will use the truncated estimator $\widetilde{\hat{f}}_{n,\hat{j}_n^f}$ that essentially possesses the same optimal statistical properties as those of \hat{f}_{n,\hat{j}_n^f} , and that remains a real non-negative function everywhere.

4.1 Synthetic experiments

4.1.1 Description of the statistical setup

We have first run some simulations with a synthetic example based on Gamma distributions. This family of distribution is parametrized by an integer γ and a scaling real number θ and is denoted by $Gamma(\gamma, \theta)$ below. We recall that the density with respect to the Lebesgue measure is given by:

$$\forall x \geq 0 \quad \Gamma_{\gamma, \theta}(x) = \frac{x^{\gamma-1} e^{-x/\theta}}{\Gamma(\gamma) \theta^\gamma}.$$

The main interest of the $Gamma(\gamma, \theta)$ distribution is that the smoothness parameter of this distribution is exactly known since its characteristic function $\Phi_{\gamma, \theta}(t)$ ($t \in \mathbb{R}$) is given by:

$$\Phi_{\gamma, \theta}(t) = (1 - \theta it)^{-\gamma}. \quad (35)$$

According to the definition of $\mathbf{H}_s(R)$ and \mathbf{H}_ν^+ , we deduce from Equation (35) that $\Gamma(\gamma, \theta)$ has a smoothness parameter s such that $s = \gamma - \frac{1}{2}$. Oppositely, the ill-posedness parameter is $\nu = \gamma$.

For the simulated random variables U, A, X , we then use the model $Z = U + AX$ and we considered several settings detailed below. We will pay a specific attention to the effect of the smoothness parameter s and to the probability of appearance of the effect p .

First dataset: effect of the smoothness parameter s of f

$$U \sim Gamma(2, 1), \quad A \sim Ber(0.4), \quad X \sim Gamma(\gamma, 1),$$

for $\gamma \in \{4, 6, 8\}$ (see Figure 2). Therefore the density of U is $g = \Gamma_{2,1}$, the density of Z is $f = \Gamma_{\gamma,1}$ and the smoothness parameter s of the unknown density f satisfies $s = \gamma$. In the first experiment, this parameter varies while the smoothness parameter of g , denoted by ν , is kept fixed to $\nu = 2$. As pointed out in Equation (35), we shall observe that the smoothness of f increases with γ , and translate a polynomial decay in the tail of the Fourier transform of f .

Second dataset: effect of the ill-posedness parameter ν of g

$$U \sim Gamma(\gamma, 1), \quad A \sim Ber(0.4), \quad X \sim Gamma(2, 1),$$

for $\gamma \in \{2, 4, 6, 8\}$. In that case, the density of U is $g = \Gamma_{\gamma,1}$, the density of Z is $f = \Gamma_{2,1}$ and the smoothness parameter s of the unknown density f satisfies $s = 2$. In the second experiment, the parameter $\nu = \gamma$ varies while s the smoothness parameter of f is kept fixed. In that situation, the degree of ill-posedness of the inverse problem increases with γ (see Equation (35)).

Below, we will investigate on these two datasets the influence of the smoothness parameters s and ν on the estimations of p and f .

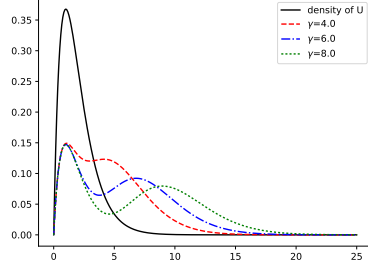


Figure 2: Density of the baseline perturbation U (in plain black) and of the observed signal Z for different values of γ (dotted curves).

Third dataset: effect of the contamination rate $1 - p$

$$U \sim \text{Gamma}(1, 1), \quad A \sim \text{Ber}(1 - p), \quad X \sim \text{Gamma}(2, 1),$$

for $1 - p \in \{0.01, 0.05, 0.1, 0.2\}$. This last dataset is used to assess the efficiency of our statistical procedure for the estimation of f when the contamination rate becomes smaller and smaller: the parameter $1 - p$ represents the probability to observe the event $U + X$ and it is expected that when $p \rightarrow 1$, the statistical estimation of f becomes harder and harder since the law of X is observed less frequently. We still use the densities $f = \Gamma_{2,1}$ and $g = \Gamma_{1,1}$.

4.1.2 Results

Results - Estimation of p We first focus on the estimation of p . For each dataset, we used some Monte Carlo replications (with 100 repetitions) in order to obtain an approximation of $\mathbb{E}\left(\|\hat{p}_{n, \hat{j}_n^p} - p\|_2^2\right)$ for different values of the sample size $n \in \{100, 200, 500, 700, 1000, 2000\}$. Let us remark that the Lepskii procedure is costly in terms of numerical simulations since the criterion for selection requires to compute all the estimators on a grid. An alternative procedure has thus been proposed by Katkovnik (1999) based on the construction of confidence intervals. However, in the setting of the estimation of a density, this construction is no longer possible. Therefore, our simulations are based on the exact Lepskii decision rule given in (15) for p and (31) for f . Figure 3 gives in a log-log plot that translates the relationship between the approximation of $\mathbb{E}\left(\|\hat{p}_{n, \hat{j}_n^p} - p\|_2^2\right)$ obtained by the Monte Carlo replications and the number of observations n used in the estimation procedure. The left hand side (resp. right hand side) of Figure 3 corresponds to the first dataset when the smoothness of f s varies (resp. the second dataset when the ill-posedness ν varies).

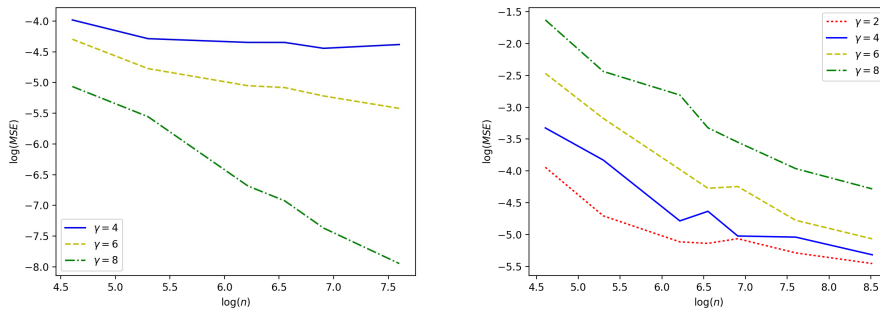


Figure 3: Log-Log plot of $n \mapsto \mathbb{E}\left(\|\hat{p}_{n, \hat{j}_n^p} - p\|_2^2\right)$ obtained by Monte Carlo replications. Left: first dataset for several values of $s = \gamma$. Right: second dataset for several values of $\nu = \gamma$. For these simulations, we choose $\varepsilon = 0.5$ and $\beta = 9.0$.

We observe that in both cases, the logarithm of the mean squared error decreases linearly with $\log(n)$, which is consistent with Theorem 1 and 3: this translates a polynomial decrease of the mean squared error in terms of the number of observations.

As shown in the left hand side of Figure 3, the estimation of p is easier for large values of s (large values of γ), while in the right hand side the estimation becomes more and more difficult when ν increases (large values of γ). This phenomenon is completely consistent with the rates derived in Theorem 1.

Finally, we notice that, for example, the theoretical slope for $\gamma = 8$ should be $-\frac{2(8-1/2)}{2(8-1/2)+2\nu} = -15/19 \simeq -0.8$ since $\nu = 2$ and we obtain an approximate slope of the order $-2.6/3$, which is rather close to the theoretical value.

Results - Estimation of f and $\mathbb{E}\left(\|\hat{f}_{n,\hat{j}_n^f} - f\|_2\right)$ In this experiment, we used the simulated dataset that corresponds to the first dataset with $\gamma = 4$.

For the non parametric estimation of f , we compared the estimation obtained with the Lepskii procedure proposed in this article with a similar one assuming that the mixing parameter p is known. In Figure 4, we draw the true value of the Fourier transform Φ_X and the estimators we obtained with the Lepskii procedure for different sample sizes $n \in \{100, 1000, 10000\}$. The left hand side corresponds to the full Lepskii procedure (see Equation (31)) when both p and f are estimated, while in the right hand side the parameter p is known in this simulation step and used in our adaptive estimation of f . On these graphs, we can first see that when the sample size increases, the bandwidth selected in the algorithm is improved since the support of $\hat{\Phi}_{X_n}$ widens.

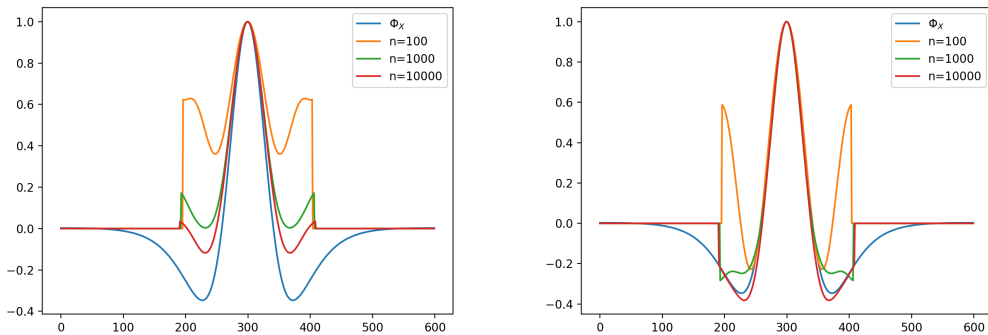


Figure 4: First dataset with $\gamma = 4$, $\varepsilon = 0.5$ and $\beta = 9$. True Fourier transform Φ_X (in blue) and its estimations counterpart for different sample sizes $n \in \{100, 1000, 10000\}$. Left hand side: Lepskii procedure described in Section 3.2 when p and f are unknown. Right hand side: the true value of the mixing parameter p is assumed to be known.

First, we shall observe that the algorithm leads to better results when the mixing parameter is known, which is not surprising since the estimation of p provides an additional noise in our statistical estimator.

We still use some Monte Carlo replications to obtain an approximation of the mean squared estimation error of f for different sample sizes (see Figure 5). Once again, we compared our algorithm (left hand side) with the case where p is known (right hand side). We find that the estimators are more accurate when p is known and that the regularity $s = \gamma$ of the function f improves the convergence speed.

Results - Impact of the contamination rate $1 - p$ Finally, in Figure 6 we investigate the effect of the size of $1 - p$ itself on the quality of our estimation of the unknown density f . As expected, the estimation becomes more and more difficult when p increases and becomes close to 1 since the number of observations of the perturbed signal $U + X$ is low. In particular, we observe a linear log-log plot when p is close to 1 ($\log(p)$ is close to 0). We therefore conjecture a polynomial dependency of

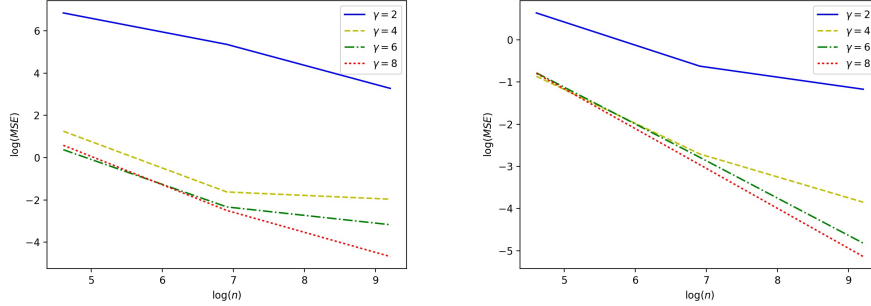


Figure 5: First dataset with several values of γ , $\beta = 9.0$ and $\varepsilon = 0.5$. $\log - \log$ plot between the approximation of $\mathbb{E}\|\hat{f}_{n, \hat{j}_n^f} - f\|_2$ obtained by Monte Carlo simulations and $n \in \{100, 1000, 10000\}$. Left hand side: p is unknown. Right hand side: known p .

the risk $\mathbb{E}\left(\|\hat{f}_{n, \hat{j}_n^f} - f\|_2\right)$ in $(1-p)^{-\alpha}$, where α is a constant related to the smoothness parameter s and the ill-posedness parameter ν . Identifying the true value of α is out of the scope of our paper, and deserves a careful inspection of the deconvolution step. We have decided to leave this issue opened in Theorems 1 and 2 and we refer to Gadat et al. (2018) for a related study in a contamination mixture model where the effect of the contamination rate is investigated in depth.

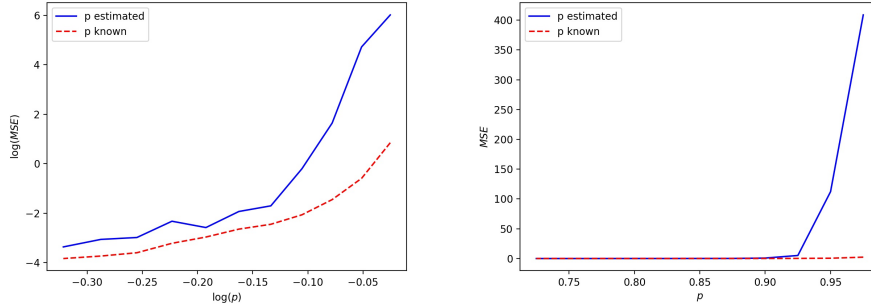


Figure 6: Third dataset: the parameters are set up to $\beta = 9.0$ and $\varepsilon = 0.5$. Approximation of $\mathbb{E}\|\hat{f}_{n, \hat{j}_n^f} - f\|_2^2$ obtained by Monte Carlo simulations (for $n = 5000$) for different values of p .

4.2 Real data

Description of the dataset The dataset used in this study is coming from fluorescence distribution measured using a flow cytometer instrument (BD Biosciences, LSRII FORTRESSA) on cells obtained from human blood. Lymphocytes are cells of the immune system useful for cancer cell destruction in immunotherapy (among other), they were extracted from the fresh blood of a healthy individual by the biologist and this cell suspension was then used for the experiment. Technically, the cell suspension was split into two parts. One half was left untouched and the baseline photo emission of untreated cells was recorded by the cytometer. The second half was mixed with a fluorescently labeled antibody (reagent) that specifically binds to the CD27 protein. Then, the photo emission by treated cells was again recorded by the cytometer. The amount of reagent binding is reflected by the fluorescence emitted by the cell that is coming only from the antibody treatment (see Figure 1). The biological experiment's goal was to assess the expression of the molecule CD27 by human lymphocytes. This protein is expressed at the surface of lymphocytes and reflect their activation status. It is therefore used to estimate the functionality of a lymphocyte population. Methods that would help to decipher the mixture of functionalities among a lymphocyte population is thus of major biological interest.

In this setting, we use the estimation procedure developed in our paper to infer the percentage of lymphocytes expressing CD27 on their surface and the conditional probability for a cell to express CD27 conditionally to its fluorescence intensity.

Preliminary estimation of the distribution of the noise U From the recorded fluorescence intensity of untreated cells, we infer the density g of U with a preliminary kernel density estimation using a Gaussian kernel through the *scipy.stats.gaussian_kde* Python software. We then use the recorded fluorescence intensity of treated cells as data for the analysis (see Figure 7(a)). In this case, the size of the recorded data is $n = 12645$.

Of course, our work only deals with the situation where the density g is known, which is unfortunately not possible in our biological situation. Therefore, we admit as a reasonable approximation the estimation of g provided by a preliminary kernel density estimation. We have not treated in our theoretical study the consequences of such a preliminary nonparametric estimation, and we leave this subject open as a future subject of investigation. Such a work would then fall into the field of statistical inverse problems with noise in the operator (see *e.g.* Cavalier and Hengartner (2005) and the references therein).

Estimation of p For the estimation of the proportion of cells expressing CD27 on their cells' surface, which corresponds to $1 - p$, we use the Lepskii procedure proposed in Section 2.2. To verify the convergence of our algorithm, we use a subsampling strategy of our data set and repeat it on permuted versions of the data.

We observe on Figure 7(b) the good behaviour of our algorithm: it produces a sharp estimation even using the half of the data and it converges to the value $\hat{p}_{n, \hat{j}_n^p} = 0.31$. We aim at comparing this value with the measurement classically used in cytometric analysis. Usually biologist use a quantity called *percentage of positive cells* which reflects the percentage of marked cells whose fluorescent intensity is higher than a given threshold. This threshold is computed as the 95th percentile of the density g (see Figure 7(a)). On this data, the percentage of positive cells is 0.85, which is much larger than the quantity we estimate $1 - \hat{p}_{n, \hat{j}_n^p} = 0.69$.

Estimation of f Next, we focus on the estimation of the density f , which represents here the distribution of fluorescence intensity due to the presence of CD27 on the cells' surface. We use two different estimation strategies:

- First, we apply the global Lepskii estimation procedure introduced in Section 3.2.
- Second, we use a similar approach but we consider that the value of p is known and given by the previously estimated $\hat{p}_{n, \hat{j}_n, p}$ above. We then use the Lepskii procedure with a fixed value of p as described in Section 3.2.

In both cases we then compute an approximation of the Fourier transform Φ_Z of the mixed signal and compared it with the empirical estimator $\hat{\Phi}_{Z, n}$ in Figure 7(c). In particular, in Figure 7(c), we show several colored curves:

- the red curve is the empirical estimator $\hat{\Phi}_{Z, n}$
- the blue dotted curve is the estimator derived from the global Lepskii procedure
- the green dotted curve corresponds to the modified Lepskii procedure for f when p is previously estimated).

We observe that the estimator of f computed with $p = 0.31$ seems to behave better, which is not surprising since the Lepskii procedure for f does not provide any theoretical guaranty for estimating p (we obtained $\hat{p}_{n, \hat{j}_n^f}^{(1)} = 0.081$, which seems very low and not realistic for our biological framework). Then we calculate the inverse transform of both estimators (see Figure 7(d)), compute the estimated density of Z and compare it with the observed histogram (Figure 7(e)). We observe first that our estimation can take negative values (due to the bandwidth parameter in the estimation) and secondly that it fails to capture the exact shape of the density, however it reproduces reasonably well the mean and the variance.

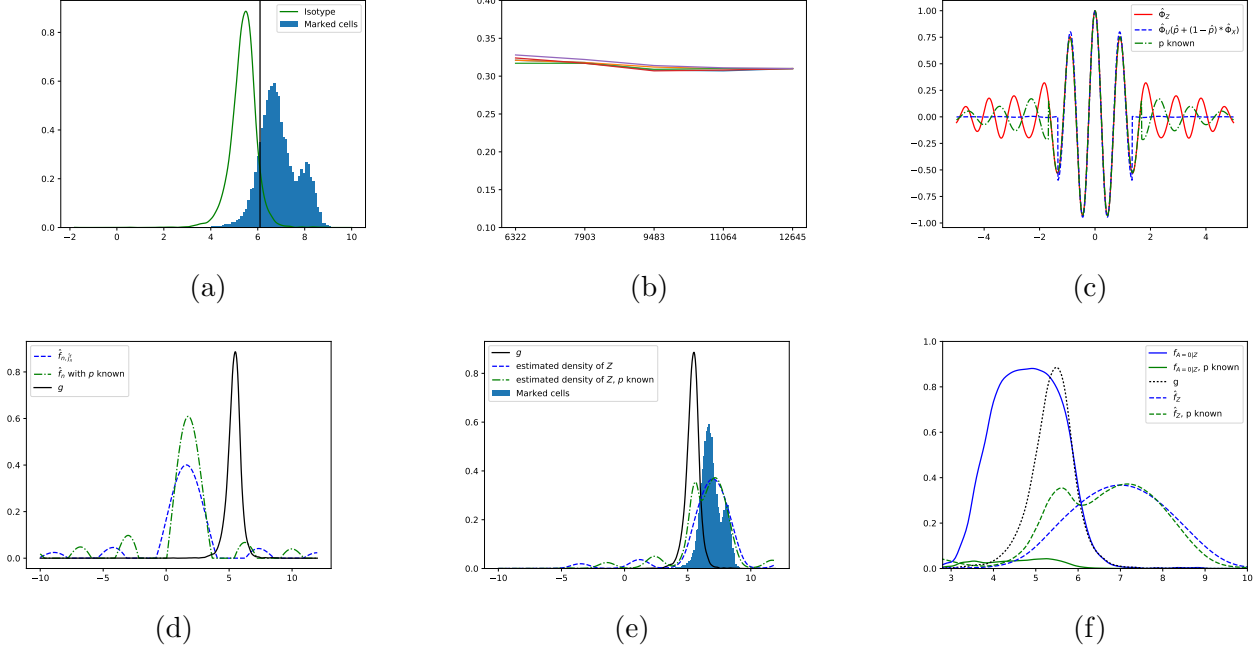


Figure 7: Results of the Lepskii procedure on real data. (a) Histogram of the fluorescence of treated cells and estimated probability distribution for the fluorescence of un-treated cells. (b) Estimation of p on an increasing sub-sample of the data (the data being permuted 5 times). (c) In red, empirical Fourier transform $\hat{\phi}_Z$ and two estimators computed respectively with the values \hat{f}_{n, \hat{j}_n^f} and $\hat{p}_{n, \hat{j}_n^f}^{(1)}$ obtained with the Lepskii procedure for f (in blue) and with the values obtained when p is first estimated and then used in the estimation of f (in green). (d) Comparison of the obtained estimators of f . (e) Estimated density of Z computed using the two different strategies for estimating f and comparison with the histogram of data. (f) Estimators of the conditional density $f_{A=0|Z}(z)$.

Finally, we study the probability that a cell expresses CD27 on its surface conditionally to the value of the observed fluorescence. We compute an estimator of this conditional density with the Bayes formula:

$$f_{A=0|Z}(z) = \frac{pg(z)}{pg(z) + (1-p)f * g(z)}$$

The results are drawn on Figure 7(f) for both estimators of f . We have not pushed further our investigations from a biological point of view since the goal of our paper is mainly theoretical.

A Technical results

A.1 Bernstein inequality

We first state a classical concentration inequality that may be found in Boucheron et al. (2013).

Theorem 9 (Bernstein inequality) *Let X_1, \dots, X_n be independent random variables with finite variance such that $|X_i| \leq b$ for some $b > 0$ almost surely for all $i \leq n$. Let $S = \sum_{i=1}^n X_i - \mathbb{E}[X_i]$ and $v = \sum_{i=1}^n \mathbb{E}[X_i^2]$. Then*

$$\mathbb{P}(|S| \geq t) \leq 2 \exp\left(-\frac{t^2}{2(v + bt/3)}\right).$$

A.2 Non adaptive estimation

In this section, we sketch the proof of Theorem Gugushvili et al. (2011) with $\tau_n = \log(n)^{-a}$ and $a > 0$ instead of the choice $\tau_n = \log(3n)^{-1}$ given in Gugushvili et al. (2011), in order to control

$$\mathbb{E}(\|\hat{f}_{n,\hat{z}_n^f} - f\|_2).$$

Proposition 10 *If $\mathbf{H}_s(R)$ and \mathbf{H}_ν^+ hold with $\nu > 1$, then the choices $h_{n,j} = n^{-\frac{1}{2s_j+2\nu}}$ in (8), $\tau_n = \log(n)^{-a}$ in (28) and $\delta_{n,j} = n^{-1/(2s_j+2\nu+1)}$ in (26) lead to an estimator $\hat{f}_{n,j}$ that satisfies the consistency rate:*

$$\mathbb{E}[\|\hat{f}_{n,j} - f\|_2^2] \leq C(s, R)n^{-\frac{2\min(s,s_j)}{2s_j+2\nu+1}},$$

where $C(s, R)$ is a positive constant depending continuously in s and R .

Proof

The bias-variance decomposition and Fubini's Theorem yield:

$$\mathbb{E}[\|\hat{f}_{n,j} - f\|_2^2] \leq \int_{-\infty}^{\infty} \left(\mathbb{E}(\hat{f}_{n,j}(t) - f(t)) \right)^2 dt + \int_{-\infty}^{\infty} \mathbb{V}(\hat{f}_{n,j}(t)) dt.$$

Integrated bias In this section, we introduce more explicit notations, we set $\hat{f}_{n,h_{n,j},\delta_{n,j}} = \hat{f}_{n,j}$ defined by (29) and set $\hat{f}_{n,\delta_{n,j}}$ the estimator of f when p is known :

$$\hat{f}_{n,\delta_{n,j}} = \frac{1}{2\pi} \int e^{-itx} \frac{\hat{\Phi}_{Z,n}^{(2)}(t) - p\Phi_U(t)}{(1-p)\Phi_U(t)} \Phi_Q(\delta_{n,j}t) dt.$$

We also define

$$\tilde{f}_{n,\delta_{n,j}} = \frac{1}{2\pi} \int e^{-itx} \frac{\hat{\Phi}_{Z,n}^{(2)}(t)}{\Phi_U(t)} \Phi_Q(\delta_{n,j}t) dt.$$

Then

$$\begin{aligned} \int \left(\mathbb{E}(\hat{f}_{n,j} - f) \right)^2 &= \int \left(\mathbb{E}(\hat{f}_{n,h_{n,j},\delta_{n,j}} - f) \right)^2 \\ &\leq 2 \int \left(\mathbb{E}(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}}) \right)^2 + 2 \int \left(\mathbb{E}(\hat{f}_{n,\delta_{n,j}} - f) \right)^2. \end{aligned}$$

Using the Parseval inequality, then assumptions (23) and $f \in \mathbf{H}_s(R)$ we deduce that the second term behaves like:

$$\begin{aligned} \int \left(\mathbb{E}(\hat{f}_{n,\delta_{n,j}} - f) \right)^2 &\leq \frac{1}{2\pi} \int_{-\infty}^{\infty} |\Phi_f(t)|^2 \frac{|\Phi_Q(\delta_{n,j}t) - 1|^2}{(\delta_{n,j}t)^{2s}} (\delta_{n,j}t)^{2s} dt, \\ &\leq MR\delta_{n,j}^{2s}. \end{aligned}$$

Then, for the first term we obtain that:

$$\begin{aligned} \int \left(\mathbb{E}(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}}) \right)^2 &\leq 2 \int \mathbb{E} \left[\tilde{f}_{n,\delta_{n,j}} \frac{(\hat{p}_{n,j}^{(1)} - p)}{(1-p)(1-\hat{p}_{n,j}^{(1)})} \right]^2 \\ &\quad + 2\mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)}{(1-p)(1-\hat{p}_{n,j}^{(1)})} \right]^2 \int \left(\frac{1}{\delta_{n,j}} Q(t\delta_{n,j}^{-1}) \right)^2 dt \\ &\leq 2\mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)}{(1-p)(1-\hat{p}_{n,j}^{(1)})} \right]^2 \int \mathbb{E} \left[\tilde{f}_{n,\delta_{n,j}} \right]^2 \\ &\quad + 2\mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)}{(1-p)(1-\hat{p}_{n,j}^{(1)})} \right]^2 \int \left(\frac{1}{\delta_{n,j}} Q(t\delta_{n,j}^{-1}) \right)^2 dt. \end{aligned}$$

Assumptions (23) on the kernel Q and the Parseval inequality lead to:

$$\int_{-\infty}^{\infty} \left(\frac{1}{\delta_{n,j}} Q(t\delta_{n,j}^{-1}) \right)^2 dt = \delta_{n,j}^{-1} \int_{-1}^1 Q(t)^2 dt < \infty.$$

Moreover, we use a concentration inequality similar to Proposition 5 stated in Lemma 11. It remains to compute:

$$\int \mathbb{E}[\tilde{f}_{n\delta_{n,j}}]^2 = \int \mathbb{V}(\tilde{f}_{n\delta_{n,j}}) + \int \mathbb{E}[\tilde{f}_{n\delta_{n,j}}^2].$$

From the definition of $\tilde{f}_{n\delta_{n,j}}$, we remark that:

$$\tilde{f}_{n\delta_{n,j}}(x) = \frac{1}{\delta_{n,j}(n-n')} \sum_{k=n'}^n W_{n,j} \left(\frac{x-Z_k}{\delta_{n,j}} \right),$$

where $W_{n,j}(x) = \int_{-1}^1 e^{-itx} \Phi_Q(t) \Phi_U(t\delta_{n,j}^{-1})^{-1} dt$. Then:

$$\int \mathbb{V}(\tilde{f}_{n\delta_{n,j}}) \leq \frac{1}{\delta_{n,j}^2(n-n')} \int_{-\infty}^{\infty} \mathbb{E} \left(W_{n,j} \left(\frac{x-Z}{\delta_{n,j}} \right)^2 \right) dx.$$

A straightforward computation using that $g \in \mathbf{H}_\nu^+$ leads to:

$$\int_{-\infty}^{\infty} \mathbb{E} \left(W_{n,j} \left(\frac{x-Z}{\delta_{n,j}} \right)^2 \right) dx \leq \frac{4\delta_{n,j}^{1-2\nu}}{d_2} \int_{-1}^1 |\Phi_Q|^2 (\delta_{n,j}^\nu + t^\nu)^2 dt.$$

Thus for some constant C :

$$\int \mathbb{V}(\tilde{f}_{n\delta_{n,j}}) \leq C\delta_{n,j}^{-1-2\nu}(n-n')^{-1},$$

which entails as $s_j > 0$ that $\mathbb{V}(\tilde{f}_{n\delta_{n,j}}) \rightarrow 0$.

It remains to bound $\int \mathbb{E}[\tilde{f}_{n,\delta_{n,j}}^2]$. The Parseval inequality and the fact that $|\Phi_X| \leq 1$ yields:

$$\int \mathbb{E}[\tilde{f}_{n,\delta_{n,j}}^2] \leq \delta_{n,j}^{-1} \int_{-1}^1 |\Phi_Q(t)|^2 dt.$$

Therefore, combining the previous inequality we deduce that:

$$\begin{aligned} \int \left(\mathbb{E}(\hat{f}_{n,j} - f) \right)^2 &\leq 2MR\delta_{n,j}^{2s} + 4C(s,R)(n-n')^{-\frac{2\min(s,s_j)+1}{2s_j+2\nu}} \\ &\quad \times \left(\delta_{n,j}^{-1} \int Q(t)^2 dt + C\delta_{n,j}^{-1-2\nu}(n-n')^{-1} + \delta_{n,j}^{-1} \int_{-1}^1 |\Phi_Q(t)|^2 dt \right). \end{aligned}$$

The second term of the r.h.s. is of order $h_{n,j}^{2\min(s,s_j)+1}\delta_{n,j}^{-1}$ which is negligible before $\delta_{n,j}^{2\min(s,s_j)}$. This entails that:

$$\int \left(\mathbb{E}(\hat{f}_{n,j} - f) \right)^2 \leq C(s,R)(n)n^{-\frac{2\min(s,s_j)}{2s_j+2\nu+1}}. \quad (36)$$

Integrated variance Let us first remark that:

$$\begin{aligned} \int_{-\infty}^{\infty} \mathbb{V}(\hat{f}_{n,j}) &= \int_{-\infty}^{\infty} \mathbb{V}(\hat{f}_{n,h_{n,j},\delta_{n,j}}) \\ &\leq 2 \int_{-\infty}^{\infty} \mathbb{V}(\hat{f}_{n,h_{n,j}}) + 2 \int_{-\infty}^{\infty} \mathbb{V}(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,h_{n,j}}). \end{aligned}$$

We have proved above that

$$2 \int_{-\infty}^{\infty} \mathbb{V}(\hat{f}_{n,h_{n,j}}) \leq C\delta_{n,j}^{-1-2\nu}(n-n')^{-1}.$$

It remains to consider the second term. We use once again an auxiliary sequence $\Psi_n \rightarrow 0$:

$$\begin{aligned}
\int_{-\infty}^{\infty} \mathbb{V}(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}}) &\leq \int_{-\infty}^{\infty} \mathbb{E}[(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}})^2] \\
&\leq \int_{-\infty}^{\infty} \mathbb{E}[(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}})^2 \mathbb{1}_{|\hat{p}_{n,j} - p| > \Psi_n}] \\
&\quad + \int_{-\infty}^{\infty} \mathbb{E}[(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}})^2 \mathbb{1}_{|\hat{p}_{n,j} - p| \leq \Psi_n}].
\end{aligned}$$

Similarly as for the integrated bias we obtain that:

$$\begin{aligned}
\int_{-\infty}^{\infty} \mathbb{E}[(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}})^2 \mathbb{1}_{|\hat{p}_{n,j} - p| > \Psi_n}] &\leq 2 \int \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \tilde{f}_{n,\delta_{n,j}}^2 \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| > \Psi_n} \right] \\
&\quad + 2\mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| > \Psi_n} \right] \int \left(\frac{1}{\delta_{n,j}} Q(t\delta_{n,j}^{-1}) \right)^2 dt.
\end{aligned}$$

The second term of the r.h.s. is bounded by:

$$2\delta_{n,j}^{-1} \frac{2(1-\tau_n)^2 + 2p^2}{\tau_n^2(1-p)} \left(\int_{-1}^1 Q(t) dt \right) \mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n).$$

We prove below in (37) that $\mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n)$ decreases exponentially fast to 0 when ψ_n is chosen as $\psi_n = n^{-1/(2(2s_{\max}+2\nu))}$. Thus, the previous term is $o\left(n^{-\frac{2\min(s,s_j)}{2s_j+2\nu+1}}\right)$.

Concerning the other term, we apply the Fubini theorem and obtain that:

$$\begin{aligned}
\int \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \tilde{f}_{n,\delta_{n,j}}^2 \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| > \Psi_n} \right] &= \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| > \Psi_n} \int \tilde{f}_{n,\delta_{n,j}}^2 \right] \\
&= \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| > \Psi_n} \int \frac{|\hat{\Phi}_{Z,n}(t)\Phi_Q(\delta_{n,j}t)|^2}{|\Phi_U(t)|^2} dt \right] \\
&\leq \frac{2+2p^2}{\tau_n^2(1-p)^2} \frac{4\delta_{n,j}^{-1-2\nu}}{d_2^2} \int_{-1}^1 |\Phi_Q(t)|^2 (1+t^\nu)^2 dt \mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n).
\end{aligned}$$

Hence, we deduce that similarly:

$$\int_{-\infty}^{\infty} \mathbb{E}[(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}})^2 \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| > \Psi_n}] \leq o\left(n^{-\frac{2\min(s,s_j)}{2s_j+2\nu+1}}\right).$$

We finally consider the event where $|\hat{p}_{n,j}^{(1)} - p| \leq \Psi_n$ and use the independence between $\hat{p}_{n,j}^{(1)}$ and $\tilde{f}_{n,\delta_{n,j}}$ to obtain:

$$\begin{aligned}
\int_{-\infty}^{\infty} \mathbb{E}[(\hat{f}_{n,h_{n,j},\delta_{n,j}} - \hat{f}_{n,\delta_{n,j}})^2 \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| \leq \Psi_n}] &\leq 2 \int \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| \leq \Psi_n} \right] \mathbb{E}[\tilde{f}_{n,\delta_{n,j}}^2] \\
&\quad + 2\mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \mathbb{1}_{|\hat{p}_{n,j}^{(1)} - p| \leq \Psi_n} \right] \int \left(\frac{1}{\delta_{n,j}} Q(t\delta_{n,j}^{-1}) \right)^2 dt.
\end{aligned}$$

From Lemma 11 and an argument similar to the one used for the integrated bias, we conclude that this last term is equal to $O\left(n^{-\frac{2\min(s,s_j)}{2s_j+2\nu+1}}\right)$, which concludes the proof of Proposition 10. \square

Lemma 11 Assume $s_j \geq 1/2$ and that $f \in \mathbf{H}_s(R)$, $g \in \mathbf{H}_\nu^+$, with the choices of $h_{n,j} = n^{-1/(2s_j+2\nu)}$ and $\tau_n = \log(n)^{-a}$ then

$$\mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)}{(1-p)(1-\hat{p}_{n,j}^{(1)})} \right]^2 \leq C(s, R) n^{-\frac{2 \min(s, s_j)+1}{2s_j+2\nu}}.$$

Proof

Let us first remark that using the Cauchy-Schwarz inequality and a truncation strategy, we have that:

$$\mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)}{(1-p)(1-\hat{p}_{n,j}^{(1)})} \right]^2 \leq \frac{1}{\tau_n^2(1-p)^2} \mathbb{E}[(\hat{p}_{n,j}^{(1)} - p)^2].$$

Moreover using (7) and (9)

$$\mathbb{E}[(\hat{p}_{n,j}^{(1)} - p)^2] \leq 2C_s R^2 h_{n,j}^{2s+1} + \frac{cte(s)}{nh_{n,j}^{2\nu-1}},$$

where $cte(s)$ is a constant that depends continuously on s . Therefore, with our choice of $h_{n,j}$ in (11) we deduce that there exist a constant $\phi(s, R)$ depending continuously on s and R such that

$$\mathbb{E}[(\hat{p}_{n,j}^{(1)} - p)^2] \leq \phi(s, R) n^{-\frac{2 \min(s, s_j)+1}{2s_j+2\nu}},$$

and thus

$$\mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)}{(1-p)(1-\hat{p}_{n,j}^{(1)})} \right]^2 \leq \frac{1}{\tau_n^2(1-p)^2} \phi(s, R) n^{-\frac{2 \min(s, s_j)+1}{2s_j+2\nu}}.$$

With this simple reasoning our upper bound depend on the truncation τ_n . Therefore, we will refine the result by using an auxiliary sequence $\Psi_n \rightarrow 0$ and split the events into two sub-cases:

$$\begin{aligned} \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)}{(1-p)(1-\hat{p}_{n,j}^{(1)})} \right]^2 &\leq \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \right] \\ &\leq \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \mathbf{1}_{|\hat{p}_{n,j}^{(1)} - p| \leq \Psi_n} \right] + \mathbb{E} \left[\frac{(\hat{p}_{n,j}^{(1)} - p)^2}{(1-p)^2(1-\hat{p}_{n,j}^{(1)})^2} \mathbf{1}_{|\hat{p}_{n,j}^{(1)} - p| > \Psi_n} \right] \\ &\leq \frac{1}{(1-p)^2(1-p-\Psi_n)^2} \phi(s, R) n^{-\frac{2 \min(s, s_j)+1}{2s_j+2\nu}} + \frac{1}{(1-p)^2 \tau_n^2} \mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n). \end{aligned}$$

In the following, our aim is to calibrate the sequence Ψ_n such that the second term is negligible compared to the first one. We have studied concentration inequalities in Section 2.2.2 for a non truncated version of $\hat{p}_{n,j}^{(1)}$, that we denote here $\tilde{p}_{n,j}^{(1)}$. Then

$$\begin{aligned} \mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n) &\leq \mathbb{P}(|\hat{p}_{n,j}^{(1)} - \tilde{p}_{n,j}^{(1)}| > \Psi_n) + \mathbb{P}(|\tilde{p}_{n,j}^{(1)} - p| > \Psi_n) \\ &\leq \mathbb{E}(\mathbf{1}_{\tau_n > \Psi_n/2} \mathbf{1}_{\tilde{p}_{n,j}^{(1)} > 1-\tau_n}) + \mathbb{P}(|\tilde{p}_{n,j}^{(1)} - p| > \Psi_n/2), \\ &\leq \mathbb{E}(\mathbf{1}_{\tau_n > \Psi_n/2} \mathbf{1}_{\tilde{p}_{n,j}^{(1)} > 1-\tau_n}) + \mathbb{P} \left(\left| \frac{1}{n} \sum_{k=1}^n \xi_{k,j} \right| > \Psi_n/4 \right) + \mathbb{P}(C_s R h_{n,l}^{s+1/2} > \Psi_n/4). \end{aligned}$$

where the random variables $\xi_{k,j}$ are defined in (20) Therefore

$$\begin{aligned} \mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n) &\leq \mathbb{P}(|\hat{p}_{n,j}^{(1)} - \tilde{p}_{n,j}^{(1)}| > \Psi_n) + \mathbb{P}(|\tilde{p}_{n,j}^{(1)} - p| > \Psi_n) \\ &\leq \mathbb{E}(\mathbf{1}_{\tau_n > \Psi_n/2} \mathbf{1}_{\tilde{p}_{n,j}^{(1)} > 1-\tau_n}) + \mathbb{P}(|\tilde{p}_{n,j}^{(1)} - p| > \Psi_n/2), \\ &\leq \mathbb{E}(\mathbf{1}_{\tau_n > \Psi_n/2} \mathbf{1}_{\tilde{p}_{n,j}^{(1)} > 1-\tau_n}) + \mathbb{P} \left(\left| \frac{1}{n-n'} \sum_{k=n'}^n \xi_{k,j} \right| > \Psi_n/4 \right) \\ &\quad + \mathbb{P}(C_s R h_{n,l}^{s+1/2} > \Psi_n/4). \end{aligned}$$

Using the Bernstein inequality, we then have:

$$\begin{aligned} \mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n) &\leq \mathbf{1}_{\tau_n > \Psi_n/2} \mathbb{P}(\tilde{p}_{n,j}^{(1)} > 1 - \tau_n) + \exp\left(-\frac{(n-n')\Psi_n^2}{8(C'h_{n,l}^{-2\nu+1} + Ch_{n,l}^{-\nu}\Psi_n/3)}\right) \\ &\quad + \mathbf{1}_{C_s R h_{n,l}^{s+1/2} > \Psi_n/4}. \end{aligned}$$

We choose $\Psi_n = n^{-1/2(2s_{max}+2\nu)}$ such that the last term of the r.h.s. is always null. Thus, it remains to consider $\mathbb{P}(\tilde{p}_{n,j}^{(1)} > 1 - \tau_n)$. Applying similarly Bernstein inequality leads to:

$$\mathbb{P}(\tilde{p}_{n,j}^{(1)} > 1 - \tau_n) \leq \mathbb{P}(|\tilde{p}_{n,j}^{(1)} - p| > 1 - \tau_n - p) \leq \exp\left(-\frac{(n-n')(1-\tau_n-p)^2}{8(C'h_{n,l}^{-2\nu+1} + Ch_{n,l}^{-\nu}(1-\tau_n-p)/3)}\right)$$

The two previous inequalities yield:

$$\begin{aligned} \mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n) &\leq \exp\left(-\frac{(n-n')(1-\tau_n-p)^2}{8(C'h_{n,l}^{-2\nu+1} + Ch_{n,l}^{-\nu}(1-\tau_n-p)/3)}\right) \\ &\quad + \exp\left(-\frac{(n-n')\Psi_n^2}{8(C'h_{n,l}^{-2\nu+1} + Ch_{n,l}^{-\nu}\Psi_n/3)}\right). \end{aligned} \tag{37}$$

It is clear that $\tau_n^{-2}\mathbb{P}(|\hat{p}_{n,j}^{(1)} - p| > \Psi_n) = o(n^{-1/2(2s_{max}+2\nu)})$, which concludes the proof. \square

A.3 Technical results for the proof of Theorem 2

Our starting point is Proposition 8:

$$\mathbb{E}\|\hat{f}_{n,\hat{j}_n^f} - f\|_2 \lesssim (\log n)^{a+1/2} n^{-s/(2s+2\nu+1)} + \sqrt{\sum_{l \geq j^*} \mathbb{P}\left(\|\hat{f}_{n,l} - f\|_2 \geq \frac{\rho_{n,l}}{2}\right)}.$$

A.3.1 Bias upper bound

Once again, the main difficulty of the proof is to obtain a deviation inequality on each event:

$$\Omega_{n,l} := \left\{ \|\hat{f}_{n,l} - f\|_2 \geq \frac{\rho_{n,l}}{2} \right\} = \left\{ \|\hat{\Phi}_{X_{n,l}}^{(2)} - \Phi_X\|_2^2 \geq \frac{\rho_{n,l}^2}{4} \right\}.$$

where the last equality comes from the Parseval identity. We begin by the following statement:

Proposition 12 *Assume $l \geq j^*$, for n large enough one has*

$$\Omega_{n,l} \subset \left\{ \|\hat{\Phi}_{X_{n,l}}^{(2)} - \mathbb{E}\hat{\Phi}_{X_{n,l}}^{(2)}\|_2^2 \geq \frac{\rho_{n,l}^2}{8} \right\}.$$

Proof

The main ingredient of the proof uses the bias-variance decomposition of the squared \mathbb{L}^2 norm. Hence, we define:

$$T_1 := \|\mathbb{E}\hat{\Phi}_{X_{n,l}}^{(2)} - \Phi_X\|_2^2 \quad \text{and} \quad T_2 := \mathbb{V}(\hat{\Phi}_{X_{n,l}}^{(2)}) = \mathbb{E}\|\mathbb{E}(\hat{\Phi}_{X_{n,l}}^{(2)}) - \hat{\Phi}_{X_{n,l}}^{(2)}\|_2^2. \tag{38}$$

A key remark comes from the fact that $\hat{\Phi}_{Z,n}^{(2)}(t)$ is built with a sample of $n - \lfloor n/2 \rfloor$ observations, that are independent of the one used to estimate p with $\hat{p}_{n,l}^{(1)}$. We can write that:

$$\begin{aligned}\mathbb{E}\hat{\Phi}_{X_{n,l}}^{(2)}(t) &= \mathbb{E}\left[\frac{p\Phi_U(t) + (1-p)\Phi_U(t)\Phi_X(t) - \hat{p}_{n,l}^{(1)}\Phi_U(t)}{(1 - \hat{p}_{n,l}^{(1)})\Phi_U(t)}\Phi_Q(\delta_{n,l}t)\right] \\ &= \mathbb{E}\left[\frac{p - \hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}}\right]\Phi_Q(\delta_{n,l}t) + \left[\mathbb{E}\left[\frac{1-p}{1 - \hat{p}_{n,l}^{(1)}}\right]\Phi_Q(\delta_{n,l}t) - 1\right]\Phi_X(t) + \Phi_X(t) \\ &= \mathbb{E}\left[\frac{p - \hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}}\right]\Phi_Q(\delta_{n,l}t)(1 - \Phi_X(t)) + [\Phi_Q(\delta_{n,l}t) - 1]\Phi_X(t) + \Phi_X(t).\end{aligned}$$

Using $(a+b)^2 \leq 2a^2 + 2b^2$, the bias term T_1 defined in (38) is upper bounded by:

$$T_1 \leq 2 \underbrace{\left\{\mathbb{E}\left[\frac{p - \hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}}\right]\right\}^2 \int_{\mathbb{R}} |\Phi_Q(\delta_{n,l}t)(1 - \Phi_X(t))|^2 dt}_{:=T_{1,1}} + 2 \underbrace{\int_{\mathbb{R}} |[\Phi_Q(\delta_{n,l}t) - 1]\Phi_X(t)|^2 dt}_{:=T_{1,2}},$$

We may apply the Cauchy-Schwarz inequality and Lemma 11 of the appendix, Section A and obtain:

$$\left\{\mathbb{E}\left[\frac{p - \hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}}\right]\right\}^2 \leq \mathbb{E}\left[\frac{p - \hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}}\right]^2 \lesssim h_{n,l}^{2s_l+1}.$$

The function $|\Phi_X|$ is bounded by 1 so that:

$$T_{1,1} \lesssim h_{n,l}^{2s_l+1} \int_{\mathbb{R}} |\Phi_Q(\delta_{n,l}t)(1 - \Phi_X(t))|^2 dt \lesssim h_{n,l}^{2s_l+1} \int_{\mathbb{R}} |\Phi_Q(\delta_{n,l}t)|^2 dt \lesssim h_{n,l}^{2s_l+1} \delta_{n,l}^{-1} \|\Phi_Q\|_2^2.$$

Moreover, for a given smoothness parameter s_l , the estimation of p is ‘‘easier’’ (faster) than the estimation of f , which is translated by $h_{n,l} = o(\delta_{n,l})$. We conclude that:

$$T_{1,1} \lesssim h_{n,l}^{2s_l} \sim n^{-2s_l/(2s_l+2\nu)} = o(\rho_{n,l}^2).$$

The upper bound of the second term $T_{1,2}$ uses the smoothness assumption on $f \in \mathbf{H}_s(R)$ and the construction of the kernel Q : applying Equation (23) with $\alpha = s$, we obtain:

$$T_{1,2} \leq 2 \int_{\mathbb{R}} |\delta_{n,l}t|^{2s} |\Phi_X(t)|^2 dt \lesssim \delta_{n,l}^{2s} = \mathcal{O}\left(n^{-2s/(2s_l+2\nu+1)}\right).$$

Since $l \geq j^*$, we deduce that $s_l \leq s$ and the additional log term in the definition of $\rho_{n,l}$ in (30) permits to conclude that:

$$T_{1,2} \leq \delta_{n,l}^{2s_l} = o(\rho_{n,l}^2).$$

Therefore, T_1 is smaller than $\rho_{n,l}^2/8$ for n large enough.

We can conclude the proof using the triangle inequality:

$$\|\hat{\Phi}_{X_{n,l}}^{(2)} - \mathbb{E}\hat{\Phi}_{X_{n,l}}^{(2)}\|_2 \geq \|\hat{\Phi}_{X_{n,l}}^{(2)} - \Phi_X\|_2 - \|\mathbb{E}\hat{\Phi}_{X_{n,l}}^{(2)} - \Phi_X\|_2.$$

Hence:

$$\|\hat{\Phi}_{X_{n,l}}^{(2)} - \Phi_X\|_2 \geq \frac{\rho_{n,l}}{2} \implies \|\hat{\Phi}_{X_{n,l}}^{(2)} - \mathbb{E}\hat{\Phi}_{X_{n,l}}^{(2)}\|_2 \geq \frac{\rho_{n,l}}{2} - \sqrt{T_1} \geq \frac{\rho_{n,l}}{2\sqrt{2}}.$$

□

A.3.2 Concentration inequality

From Proposition 12, we can see that the Lepskii rule will perform well if we succeed in deriving an upper bound of the following probability:

$$\mathbb{P} \left(\left\| \hat{\Phi}_{X_{n,l}}^{(2)} - \mathbb{E} \hat{\Phi}_{X_{n,l}}^{(2)} \right\|_2^2 \geq \frac{\rho_{n,l}^2}{8} \right).$$

For a given $t \in \mathbb{R}$, we can rewrite this expression as:

$$\begin{aligned} \hat{\Phi}_{X_{n,l}}^{(2)}(t) - \mathbb{E} \hat{\Phi}_{X_{n,l}}^{(2)}(t) &= \Phi_Q(\delta_{n,l}t) \left[\frac{\hat{\Phi}_{Z,n}^{(2)}(t) - \hat{p}_{n,l}^{(1)} \Phi_U(t)}{(1 - \hat{p}_{n,l}^{(1)}) \Phi_U(t)} \right] - \Phi_Q(\delta_{n,l}t) \mathbb{E} \left[\frac{\hat{\Phi}_{Z,n}^{(2)}(t) - \hat{p}_{n,l}^{(1)} \Phi_U(t)}{(1 - \hat{p}_{n,l}^{(1)}) \Phi_U(t)} \right] \\ &= \frac{\Phi_Q(\delta_{n,l}t)}{\Phi_U(t)} \left[\frac{\hat{\Phi}_{Z,n}^{(2)}(t) - \hat{p}_{n,l}^{(1)} \Phi_U(t)}{1 - \hat{p}_{n,l}^{(1)}} \right] - \frac{\Phi_Q(\delta_{n,l}t)}{\Phi_U(t)} \Phi_Z(t) \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] + \Phi_Q(\delta_{n,l}t) \mathbb{E} \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} \right] \\ &= \frac{\Phi_Q(\delta_{n,l}t)}{\Phi_U(t)} \left[\frac{\hat{\Phi}_{Z,n}^{(2)}(t)}{1 - \hat{p}_{n,l}^{(1)}} - \Phi_Z(t) \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] \right] - \Phi_Q(\delta_{n,l}t) \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} \right] \right] \\ &= T_{2,1} + T_{2,2} - T_{2,3}, \end{aligned}$$

where

$$T_{2,1} := \frac{\Phi_Q(\delta_{n,l}t)}{\Phi_U(t)} \frac{\hat{\Phi}_{Z,n}^{(2)}(t) - \Phi_Z(t)}{1 - \hat{p}_{n,l}^{(1)}} \quad T_{2,2} := \frac{\Phi_Q(\delta_{n,l}t) \Phi_Z(t)}{\Phi_U(t)} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] \right],$$

and

$$T_{2,3} := \Phi_Q(\delta_{n,l}t) \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} \right] \right].$$

The terms $T_{2,2}$ and $T_{2,3}$ can be upper bounded easily while $T_{2,1}$ deserves a specific attention.

Study of $T_{2,2}$: We first remark that $\mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right]$ is close to $\frac{1}{1-p}$ using the upper bound:

$$\left| \frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \frac{1}{1-p} \right| \leq \frac{|p - \hat{p}_{n,l}^{(1)}|}{1-p} (\log n)^a, \quad (39)$$

where the inequality derives from $\hat{p}_{n,l}^{(1)} \leq 1 - \tau_n$ and (28). Therefore, applying Theorem 3 with the smoothness parameter s_l we deduce that:

$$\left| \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] - \frac{1}{1-p} \right| \lesssim (\log n)^a n^{-(s_l+1/2)/(2s_l+2\nu)}, \quad (40)$$

Consider a constant $c > 0$, the term $T_{2,2}$ is then handled as follows:

$$\begin{aligned} \mathbb{P}(\|T_{2,2}\|_2 \geq c\rho_{n,l}) &\leq \mathbb{P} \left(\left| \frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c\rho_{n,l}}{\|\Phi_Q(\delta_{n,l}t) \Phi_Z \Phi_U^{-1}\|_2} \right) \\ &\leq \mathbb{P} \left(\left| \frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c\rho_{n,l}}{\|\Phi_Q(\delta_{n,l}t) \Phi_Z \Phi_U^{-1}\|_2} \right) \\ &\leq \mathbb{P} \left(\left| \frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c}{2\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} \right), \end{aligned}$$

where we use the upper bound of the L_2 norm: $\|\Phi_Q(\delta_{n,l}t) \Phi_Z \Phi_U^{-1}\|_2^2 \leq 4 \int_{\mathbb{R}} |\Phi_Q(\delta_{n,l}t)|^2 dt \leq 4\delta_{n,l}^{-1} \|\Phi_Q\|_2^2$. We now use Equation (40) and obtain:

$$\begin{aligned}
\mathbb{P} \left(\left| \frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c}{2\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} \right) \\
\leq \mathbb{P} \left(\left| \frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \frac{1}{1-p} \right| + \left| \frac{1}{1-p} - \mathbb{E} \left[\frac{1}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c}{2\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} \right) \\
\leq \mathbb{P} \left(\left| \frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \frac{1}{1-p} \right| \geq \frac{c}{2\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} - C(\log n)^a n^{-(s_l+1/2)/(2s_l+2\nu)} \right) \\
\leq \mathbb{P} \left(\left| \frac{1}{1 - \hat{p}_{n,l}^{(1)}} - \frac{1}{1-p} \right| \geq \frac{c}{4\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} \right),
\end{aligned}$$

for n large enough because the approximation term involved in the upper bound of Equation (40) is negligible compared to $\rho_{n,l} \delta_{n,l}^{1/2}$. Using again (39), we obtain that for a sufficiently small constant η (independent on n and l):

$$\mathbb{P}(\|T_{2,2}\|_2 \geq c\rho_{n,l}) \leq \mathbb{P}(|p - \hat{p}_{n,l}^{(1)}| \geq \eta \rho_{n,l} \delta_{n,l}^{1/2}).$$

In order to apply the Bernstein inequality (see Theorem 9 in the appendix Section A), let us remark first that the penalty $\kappa_{n,l}$ defined by (13) satisfies:

$$\kappa_{n,l} = o\left(\rho_{n,l} \delta_{n,l}^{1/2}\right).$$

Since for n large enough we have $p \leq 1 - \tau_n$, the truncated estimator satisfies:

$$\|\hat{p}_{n,l}^{(1)} - p\| \leq \|\hat{p}_{n/2,l} - p\|$$

where $\hat{p}_{n,l}^{(1)}$ is given by (27). Thus we can conclude that:

$$\mathbb{P}(\|T_{2,2}\|_2 \geq c\rho_{n,l}) \lesssim n^{-\beta^2/64}. \quad (41)$$

Study of $T_{2,3}$: This study is of the same nature because $\mathbb{E} \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} \right]$ is close to $\frac{p}{1-p}$: Inequality (40) reads:

$$\left| \mathbb{E} \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} \right] - \frac{p}{1-p} \right| = \left| \mathbb{E} \left[\frac{p - \hat{p}_{n,l}^{(1)}}{(1-p)(1 - \hat{p}_{n,l}^{(1)})} \right] \right| \lesssim (\log n)^a n^{-(s_l+1/2)/(2s_l+2\nu)},$$

Hence

$$\begin{aligned}
\mathbb{P}(\|T_{2,3}\|_2 \geq c\rho_{n,l}) &\leq \mathbb{P} \left(\left| \frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{p}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c\rho_{n,l}}{2\|\Phi_Q(\delta_{n,l} \cdot)\|_2} \right) \\
&\leq \mathbb{P} \left(\left| \frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{p}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c}{2\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} \right),
\end{aligned}$$

We use again Equation (40) and obtain that:

$$\begin{aligned}
\mathbb{P} \left(\left| \frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} - \mathbb{E} \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c}{2\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} \right) \\
\leq \mathbb{P} \left(\left| \frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} - \frac{p}{1-p} \right| + \left| \frac{p}{1-p} - \mathbb{E} \left[\frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} \right] \right| \geq \frac{c}{2\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} \right) \\
\leq \mathbb{P} \left(\left| \frac{\hat{p}_{n,l}^{(1)}}{1 - \hat{p}_{n,l}^{(1)}} - \frac{p}{1-p} \right| \geq \frac{c}{4\|\Phi_Q\|_2} \rho_{n,l} \delta_{n,l}^{1/2} \right),
\end{aligned}$$

and similar arguments used for $T_{2,2}$ yields:

$$\mathbb{P}(\|T_{2,3}\|_2 \geq c\rho_{n,l}) \lesssim n^{-\beta^2/64}. \quad (42)$$

Study of $T_{2,1}$: We recall that the support of Φ_Q is $[-1; 1]$, which implies that

$$\|T_{2,1}\|_2^2 = \int_{-\delta_{n,l}^{-1}}^{\delta_{n,l}^{-1}} \frac{|\Phi_Q(\delta_{n,l}t)|^2}{|\Phi_U(t)|^2} \frac{|\hat{\Phi}_{Z,n}^{(2)}(t) - \Phi_Z(t)|^2}{(1 - \hat{\rho}_{n,l}^{(1)})^2} dt.$$

We now define $M_{n,l}$ as the supremum of the empirical process $\hat{\Phi}_{Z,n}^{(2)}(t) - \Phi_Z(t)$ when t varies between $-\delta_{n,l}^{-1}$ and $\delta_{n,l}^{-1}$:

$$M_{n,l} := \sup_{|t| \leq \delta_{n,l}^{-1}} \left| \hat{\Phi}_{Z,n}^{(2)}(t) - \Phi_Z(t) \right|,$$

and write that:

$$\|T_{2,1}\|_2 \leq M_{n,l} \times \frac{1}{(1 - \hat{\rho}_{n,l}^{(1)})} \times \left\| \frac{\Phi_Q(\delta_{n,l}t)}{\Phi_U(t)} \right\|_2.$$

We then deduce that:

$$\begin{aligned} \mathbb{P}(\|T_{2,1}\|_2 \geq \rho_{n,l}) &\leq \mathbb{P}\left(\left\| \frac{\Phi_Q(\delta_{n,l}t)}{\Phi_U(t)} \right\|_2 \frac{M_{n,l}}{1 - \hat{\rho}_{n,l}^{(1)}} \geq \rho_{n,l}\right) \\ &\leq \mathbb{P}\left(\left\| \Phi_Q(\delta_{n,l}\cdot) \Phi_U(\cdot)^{-1} \right\|_2 M_{n,l} \geq \tau_n \rho_{n,l}\right). \end{aligned}$$

To upper bound the L^2 norm of $\Phi_Q(\delta_{n,l}\cdot) \Phi_U^{-1}$, we follow a standard argument of Fan (1991) using that $g \in \mathbf{H}_\nu^+$ and deduce that:

$$\left\| \Phi_Q(\delta_{n,l}\cdot) \Phi_U(\cdot)^{-1} \right\|_2^2 = \int_{\mathbb{R}} \frac{|\Phi_Q(\delta_{n,l}t)|^2}{|\Phi_U(t)|^2} dt \leq \frac{1}{\delta_{n,l}} \int_{-1}^1 \frac{|\Phi_Q(\xi)|^2}{|\Phi_U(\xi \delta_{n,l}^{-1})|^2} d\xi \leq \frac{\|\Phi_Q\|_2^2}{d_2^{-2}} \delta_{n,l}^{-(2\nu+1)}.$$

This inequality yields:

$$\mathbb{P}(\|T_{2,1}\|_2 \geq \rho_{n,l}) \leq \mathbb{P}\left(M_{n,l} \geq \frac{d_2}{\|\Phi_Q\|_2} \tau_n \rho_{n,l} \delta_{n,l}^{1/2+\nu}\right). \quad (43)$$

It remains to obtain a deviation inequality on $M_{n,l}$, which is the supremum norm of $\hat{\Phi}_{Z,n}^{(2)} - \Phi_Z$ on the interval $[-\delta_{n,l}^{-1}, \delta_{n,l}^{-1}]$. For this purpose, we could try to use the roadmap of applying the Talagrand inequality associated to a chaining strategy (see Boucheron et al. (2013) for example) but these general theorems will not lead to a satisfactory deviation bound because of our assumption on U , which is not a sub-Gaussian random variable.

Nevertheless, it is possible to exploit the feature of the process $(\hat{\Phi}_{Z,n}^{(2)}(t) - \Phi_Z(t))_{-\delta_{n,l}^{-1} \leq t \leq \delta_{n,l}^{-1}}$, which is an average of centered bounded random processes. This original strategy is based on a discretization of the interval $[-\delta_{n,l}^{-1}, \delta_{n,l}^{-1}]$ and a careful understanding of the variations of the random process over this grid of discretization. In particular, it makes it possible to use a union bound that do not harms the efficiency of our concentration inequality. We define $t \mapsto W_n(t)$ by:

$$\forall t \in \mathbb{R} : \quad W_n(t) = \hat{\Phi}_{Z,n}^{(2)}(t) - \Phi_Z(t).$$

We introduce a parameter $\alpha > 0$ that will be chosen small enough below. The control of $M_{n,l}$ relies on the simple remark that:

$$W_n(t+u) - W_n(t) = \frac{1}{n - n' + 1} \sum_{j=n'}^n \left[e^{itZ_j} - e^{i(t+u)Z_j} \right] - \mathbb{E} \left[e^{itZ} - e^{i(t+u)Z} \right],$$

where we set $n' = \lceil n/2 \rceil + 1$. The elementary remark $|e^{ia} - e^{ib}| \leq |a - b|$ for all $(a, b) \in \mathbb{R}^2$ yields

$$|W_n(t+u) - W_n(t)| \leq |u| \left(\frac{1}{n - n' + 1} \sum_{j=n'}^n |Z_j| + \mathbb{E}|Z| \right). \quad (44)$$

In particular, we define the following event:

$$\Omega_n := \left\{ \frac{1}{n - n' + 1} \sum_{j=n'}^n |Z_j| < 2\mathbb{E}|Z| \right\}.$$

The Tchebychev inequality applied to the \mathbb{L}^2 -random variables $(Z_j)_{n' \leq j \leq n}$ shows that:

$$\mathbb{P}(\Omega_n^c) \leq \frac{\text{Var}(|Z|)}{n\mathbb{E}[|Z|]^2} \leq \frac{c}{n},$$

with $c = \text{Var}(|Z|) \{\mathbb{E}[|Z|]\}^{-2}$. Obviously, $\mathbb{E}[|Z|] > 0$ because $Z = U + AX$ and Z cannot be a.s. 0 because U is assumed to satisfy \mathbf{H}_ν^+ . The important point on Ω_n is that Inequality (44) does not depend on t , so that $\forall s > 0$:

$$\begin{aligned} \mathbb{P}\left(\sup_{|t| \leq \delta_{n,l}^{-1}, |u| \leq \alpha} |W_n(t+u) - W_n(t)| > s\right) &\leq \mathbb{P}\left(\alpha \left(\frac{1}{n - n'} \sum_{j=n'}^n |Z_j| + \mathbb{E}|Z|\right) > s \cap \Omega_n\right) + \mathbb{P}(\Omega_n^c) \\ &\leq \mathbf{1}_{3\alpha\mathbb{E}[|Z|] > s} + \frac{c}{n}. \end{aligned} \quad (45)$$

We now define the threshold $s_{n,l}$ and the window size $\alpha_{n,l}$ by:

$$s_{n,l} := \frac{1}{2} \frac{d_2}{\|\Phi_Q\|_2} \tau_n \rho_{n,l} \delta_{n,l}^{1/2+\nu} \quad \text{and} \quad \alpha_{n,l} := \frac{s_{n,l}}{3\mathbb{E}[|Z|]}. \quad (46)$$

We are naturally driven to handle a discrete grid $\mathcal{T}_{n,l}$ regularly spaced from $-\delta_{n,l}^{-1}$ to $\delta_{n,l}^{-1}$ whose step-size is $\alpha_{n,l}$. We have:

$$|\mathcal{T}_{n,l}| \leq \frac{2\delta_{n,l}^{-1}}{\alpha_{n,l}} = \frac{6\mathbb{E}[|Z|]\delta_{n,l}^{-1}}{s_{n,l}} \quad \text{and} \quad \forall t \in [-\delta_{n,l}^{-1}, \delta_{n,l}^{-1}], \quad \exists \tau_t \in \mathcal{T}_{n,l} : \quad |t - \tau_t| \leq \alpha_{n,l}.$$

These settings permit to deduce from (45) that:

$$\mathbb{P}\left(\sup_{|t| \leq \delta_{n,l}^{-1}, |u| \leq \alpha_{n,l}} |W_n(t+u) - W_n(t)| > s_{n,l}\right) \leq \frac{c}{n}. \quad (47)$$

Now, for each element of the grid $\tau \in \mathcal{T}_{n,l}$, $W_n(\tau)$ is a mean of n random variables that whose modulus are bounded by 1. The Hoeffding inequality implies that $\forall \tau \in \mathcal{T}_{n,l}$:

$$\begin{aligned} \mathbb{P}(|W_n(\tau)| > s_{n,l}) &\leq \mathbb{P}\left(\left|\frac{1}{n - n'} \sum_{j=n'}^n \cos(\tau Z_j) - \mathbb{E}(\cos(\tau Z))\right| < \frac{s_{n,l}}{2}\right) \\ &\quad + \mathbb{P}\left(\left|\frac{1}{n - n'} \sum_{j=n'}^n \sin(\tau Z_j) - \mathbb{E}(\sin(\tau Z))\right| < \frac{s_{n,l}}{2}\right) \\ &\leq 4 \exp\left(-\frac{(n - n')s_{n,l}^2}{8}\right). \end{aligned} \quad (48)$$

We now produce an upper bound of $\mathbb{P}(\|T_{2,1}\|_2 \geq \rho_{n,l})$. Equation (43) yields:

$$\begin{aligned}
\mathbb{P}(\|T_{2,1}\|_2 \geq \rho_{n,l}) &= \mathbb{P}\left(\sup_{|t| \leq \delta_{n,l}^{-1}} |W_n(t)| \geq 2s_{n,l}\right) \\
&\leq \mathbb{P}\left(\sup_{|t| \leq \delta_{n,l}^{-1}, |u| \leq h_{n,l}} |W_n(t+u) - W_n(t)| \geq s_{n,l} \quad \text{and} \quad \sup_{|t| \leq \delta_{n,l}^{-1}} |W_n(t)| \geq 2s_{n,l}\right) \\
&\quad + \mathbb{P}\left(\sup_{|t| \leq \delta_{n,l}^{-1}, |u| \leq h_{n,l}} |W_n(t+u) - W_n(t)| \leq s_{n,l} \quad \text{and} \quad \sup_{|t| \leq \delta_{n,l}^{-1}} |W_n(t)| \geq 2s_{n,l}\right) \\
&\leq \mathbb{P}\left(\sup_{|t| \leq \delta_{n,l}^{-1}, |u| \leq h_{n,l}} |W_n(t+u) - W_n(t)| \geq s_{n,l}\right) + \mathbb{P}\left(\sup_{\tau \in \mathcal{T}_{n,l}} |W_n(\tau)| \geq s_{n,l}\right) \\
&\leq \frac{c}{n} + \sum_{\tau \in \mathcal{T}_{n,l}} \mathbb{P}(|W_n(\tau)| \geq s_{n,l}) \\
&\leq \frac{c}{n} + \frac{2\delta_{n,l}^{-1}}{\alpha_{n,l}} \times 4 \exp\left(-\frac{(n-n')s_{n,l}^2}{8}\right),
\end{aligned}$$

where we successively applied the triangle inequality, a union bound over $\mathcal{T}_{n,l}$, and then inequalities (47) and (48). An immediate computation from (46) (28), (30) and (26) shows that:

$$\begin{aligned}
ns_{n,l}^2 &= n \frac{d_2^2}{4\|\Phi_Q\|_2^2} \tau_n^2 \rho_{n,l}^2 \delta_{n,l}^{1+2\nu} \\
&= n \frac{d_2^2}{4\|\Phi_Q\|_2^2} \times (\log n)^{-2a} \times \beta^2 (\log n)^{2a+1} n^{-2s_l/(2s_l+2\nu+1)} \times n^{-(1+2\nu)/(2s_l+2\nu+1)} \\
&= \frac{\beta^2 d_2^2}{4\|\Phi_Q\|_2^2} \log(n),
\end{aligned}$$

and

$$\delta_{n,l}^{-1} \alpha_{n,l}^{-1} = \frac{6\mathbb{E}(|Z|)\|\Phi_Q\|_2}{d_2} \beta \log(n)^{-1/2} n^{\frac{3/2+\nu+s_l}{2\nu+2s_l+1}}.$$

We therefore deduce that:

$$\mathbb{P}(\|T_{2,1}\|_2 \geq \rho_{n,l}) \leq \frac{c}{n} + \beta \log(n)^{-1/2} \frac{48\mathbb{E}(|Z|)\|\Phi_Q\|_2}{d_2} n^{-\frac{\beta^2 d_2^2}{24\|\Phi_Q\|_2^2} + \frac{3/2+\nu+s_l}{2\nu+2s_l+1}}. \quad (49)$$

Conclusion To conclude with a concentration estimate of $\hat{\Phi}_{X_{n,l}}^{(2)}$ it remains to combine (41), (42) and (49) to obtain the following result.

Proposition 13 *Let $f \in \mathbf{H}_s(R)$, g a known density in \mathbf{H}_ν^+ with $\nu > 1$, and $l \geq j^*$, then*

$$\mathbb{P}\left(\|\hat{\Phi}_{X_{n,l}}^{(2)} - \mathbb{E}\hat{\Phi}_{X_{n,l}}^{(2)}\|_2^2 \geq \frac{\rho_{n,l}^2}{8}\right) \lesssim n^{-\frac{\beta^2}{64}} + n^{-1} + \log(n)^{-1/2} n^{-\frac{\beta^2 d_2^2}{24\|\Phi_Q\|_2^2} + \frac{3/2+\nu+s_l}{2\nu+2s_l+1}}.$$

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