

# NONPARAMETRIC MULTIPLICATIVE DECONVOLUTION IN SURVIVAL ANALYSIS

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**Abstract.** We study the non-parametric estimation of an unknown survival function  $S$  with support on  $\mathbb{R}_+$  based on a sample with multiplicative measurement errors. The proposed fully-data driven procedure is based on estimation of the Mellin transform of the survival function and a regularisation of the inverse of the Mellin transform by a spectral cut-off. The upcoming bias-variance trade-off is handled by a data-driven choice of the cut-off parameter. For the analysis of the variance term, we consider the i.i.d. case and incorporate dependent observations in form of Bernoulli shift processes and  $\beta$ -mixing sequences.

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## 1 Data-driven survival function estimator under multiplicative measurement errors

### 1.1 The model

In this work we are interested in estimating the unknown survival function  $S : \mathbb{R}_+ \rightarrow \mathbb{R}_+$  of a positive random variable  $X$ , defined as

$$S : \mathbb{R}_+ \rightarrow [0, 1], x \mapsto \mathbb{P}(X > x),$$

given identically distributed copies of  $Y = XU$  where  $X$  and  $U$  are independent of each other and  $U$  has a known density  $g : \mathbb{R}_+ \rightarrow \mathbb{R}_+$ . In this setting the survival function  $S_Y : \mathbb{R}_+ \rightarrow \mathbb{R}_+$  of  $Y$  is given by

$$S_Y(y) := \int_0^\infty S(x)g(y/x)dx, \quad y \in \mathbb{R}_+.$$

The estimation of  $S$  using a sample  $Y_1, \dots, Y_n$  from  $Y$  is thus an inverse problem called multiplicative deconvolution. We will allow for certain dependency structures on the sample  $Y_1, \dots, Y_n$ . More precisely, we assume that  $X_1, \dots, X_n$  is a stationary process while the error terms  $U_1, \dots, U_n$  will be independent and identically distributed (i.i.d.).

## 1.2 Estimation strategy

To solve this particular multiplicative deconvolution problem, we use the rich theory of Mellin transforms, in analogy to Brenner Miguel et al (2021). In fact, for a positive random variable  $Z$  and  $c \in \mathbb{R}$  with  $\mathbb{E}(Z^{c-1}) < \infty$  we can define the Mellin transform of the distribution  $\mathbb{P}^Z$  as the function

$$\mathcal{M}_c[\mathbb{P}^Z] : \mathbb{R} \rightarrow \mathbb{C}, \quad t \mapsto \mathbb{E}(Z^{c-1+it}).$$

As a direct consequence of this definition, we get the convolution theorem, which states for two positive, independent random variable  $Z_1, Z_2$  with  $\mathbb{E}(Z_1^{c-1}), \mathbb{E}(Z_2^{c-1}) < \infty$ , we have  $\mathcal{M}_c[\mathbb{P}^{Z_1 Z_2}] = \mathcal{M}_c[\mathbb{P}^{Z_1}] \mathcal{M}_c[\mathbb{P}^{Z_2}]$ . More general, we define for any function  $h \in \mathbb{L}^1(\mathbb{R}_+, x^{c-1})$ , the space of all measure function with  $\int_0^\infty |h(x)| x^{c-1} dx < \infty$ , the Mellin transform  $\mathcal{M}_c[h] : \mathbb{R} \rightarrow \mathbb{C}$  by

$$\mathcal{M}_c[h](t) := \int_0^\infty h(x) x^{c-1+it} dx, \quad t \in \mathbb{R}.$$

Using this definition, we can state that, under the assumption  $\mathbb{E}(X^{1/2}) < \infty$ , the following calculation rule of the Mellin transform of a survival function holds true

$$\mathcal{M}_{1/2}[S](t) = (1/2 + it)^{-1} \mathcal{M}_{3/2}[\mathbb{P}^X](t), \quad t \in \mathbb{R}.$$

Additionally,  $\mathbb{E}(X^{1/2})$  implies that  $S \in \mathbb{L}^2(\mathbb{R}_+)$ , that is  $\|S\|^2 := \int_0^\infty S^2(x) dx < \infty$ , and

$$S(x) = \frac{1}{2\pi} \int_{-\infty}^\infty x^{-1/2-it} \frac{\mathcal{M}_{3/2}[\mathbb{P}^X](t)}{(1/2 + it)} dt, \quad x \in \mathbb{R}_+,$$

using the inverse Mellin transform, compare Brenner Miguel et Phandoidaen (2021). Thus, we propose the spectral-cut off estimator  $\widehat{S}_k$  for  $k \in \mathbb{R}_+$  by

$$\widehat{S}_k(x) := \frac{1}{2\pi} \int_{-k}^k x^{-1/2-it} \frac{\widehat{\mathcal{M}}(t)}{(1/2 + it) \mathcal{M}_{3/2}[g](t)} dt, \quad \text{with } \widehat{\mathcal{M}}(t) := n^{-1} \sum_{j=1}^n Y_j^{1/2+it}. \quad (1)$$

## 1.3 Upper bounds and data-driven method

Setting  $S_k := \mathbb{E}(\widehat{S}_k)$  and assuming that  $\mathbb{E}(Y) < \infty$  we deduce the usual squared bias variance decomposition

$$\mathbb{E}(\|\widehat{S}_k - S\|^2) = \|S - S_k\|^2 + \mathbb{E}(\|\widehat{S}_k - S_k\|^2),$$

where we can decompose the variance term in a term dependent on the underlying inverse problem and a term driven by the dependence structure, that is

$$\mathbb{E}(\|\widehat{S}_k - S_k\|^2) \leq \frac{\mathbb{E}(Y_1)}{2\pi n} \int_{-k}^k |(1/2 + it)\mathcal{M}_{3/2}[g](t)|^{-2} dt + \frac{1}{2\pi} \int_{-k}^k \text{Var}(\mathcal{M}_{1/2}[\widehat{S}_X](t)) dt$$

where  $\widehat{S}_X(x) := n^{-1} \sum_{j=1}^n \mathbf{1}_{(0, X_j)}(x)$  is the empirical survival function. In total, we can show for any  $k \in \mathbb{R}_+$ ,

$$\mathbb{E}(\|\widehat{S}_k - S\|^2) \leq \|S - S_k\|^2 + \mathbb{E}(Y_1) \frac{\Delta_g(k)}{n} + \frac{1}{2\pi} \int_{-k}^k \text{Var}(\mathcal{M}_{1/2}[\widehat{S}_X](t)) dt,$$

where  $\Delta_g(k) := (2\pi)^{-1} \int_{-k}^k |(1/2 + it)\mathcal{M}_{3/2}[g](t)|^{-2} dt$ . In other words, we are able to decompose the risk of our estimator into a squared bias term, an variance term which is driven by the inverse problem and a variance term which is dependent on the dependence structure of the sample  $X_1, \dots, X_n$ .

For the case of independent observations  $X_1, \dots, X_n$  we deduce that

$$\frac{1}{2\pi} \int_{-k}^k \text{Var}(\mathcal{M}_{1/2}[\widehat{S}_X](t)) dt \leq \frac{\mathbb{E}(X)}{n}$$

making it neglectable. For different dependency structures, for instance beta mixing or functional dependency measures, upper bounds for the second variance term are derived in Brenner Miguel et Phandoidaen (2021).

## 1.4 Data-driven choice of $\widehat{k}$

While the squared bias term  $\|S - S_k\|^2$  is decreasing for  $k$  increasing, the variance term  $\mathbb{E}(\|\widehat{S}_k - S_k\|^2)$  is increasing. This contrary behaviour of these terms implies that the choice of a suitable cut-off parameter  $k \in \mathbb{R}_+$  is non-trivial. To handle this bias-variance dilemma, we suggest a data-driven choice of the parameter  $k \in \mathbb{R}_+$  based on a penalized contrast approach, that is, for a  $\chi > 0$  we set

$$\widehat{k} := \arg \min_{k \in \mathcal{K}_n} -\|\widehat{S}_k\|^2 + 2\chi \widehat{\sigma}_Y \Delta_g(k) n^{-1},$$

where  $\mathcal{K}_n := \{k \in \{1, \dots, n\} : \Delta_g(k) \leq n\}$  and  $\widehat{\sigma}_Y := n^{-1} \sum_{j=1}^n Y_j^{1/2}$ . Then under regularity assumptions on  $g$ , and the moment assumption  $\mathbb{E}(Y_1^{5/2}) < \infty$ , we can state that for all  $\chi > 96$ ,

$$\begin{aligned} \mathbb{E}(\|S - \widehat{S}_{\widehat{k}}\|^2) &\leq 6 \inf_{k \in \mathcal{K}_n} \left( \|S - S_k\|^2 + 2\chi \mathbb{E}(Y_1) \frac{\Delta_g(k)}{n} \right) \\ &\quad + C(g, f) \left( \frac{1}{n} + \text{Var}(\widehat{\sigma}_X) + \int_{-n}^n \text{Var}(\mathcal{M}_{1/2}[\widehat{S}_X](t)) dt \right) \end{aligned}$$

where  $C(g, f) > 0$  is a constant depending on  $\chi$ , the error density  $g$ ,  $\mathbb{E}(X_1^{5/2})$ ,  $\sigma_X := \mathbb{E}(X_1)$  and  $\widehat{\sigma}_X := n^{-1} \sum_{j=1}^n X_j$ . Thus, we can say that the data-driven estimator  $\widehat{S}_{\widehat{k}}$  realises the optimal choice of  $k \in \mathbb{R}_+$  among the set  $\mathcal{K}_n$  which minimise the sum of the squared bias and variance term up to an access risk consisting of an neglectable  $n^{-1}$  term and a term driven by the dependency structure of the sample  $X_1, \dots, X_n$ . Again, in the case of independent observations this term is neglectable, too.

## 1.5 Graphiques

To visualise the finite sample behaviour of our estimator, we present an extract of the simulation study of Brenner Miguel et Phandoidaen (2021) where we visualise the estimator  $\widetilde{S}_{\widehat{k}} := \max(\min(\widehat{S}_{\widehat{k}}, 1), 0)$  where obviously  $\|\widetilde{S}_{\widehat{k}} - S\|^2 \leq \|\widehat{S}_{\widehat{k}} - S\|^2$  holds.

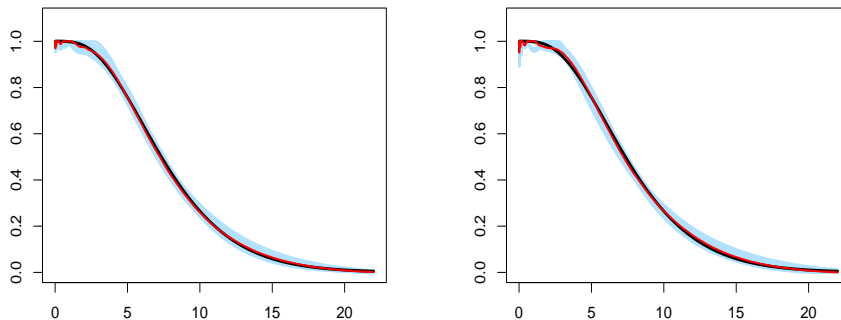


Figure 1: Considering the estimators  $\widetilde{S}_{\widehat{k}}$ , we depict 50 Monte-Carlo simulations with error densities  $g(x) = \mathbb{1}_{(0,1)}(x)$  (left) and  $g(x) = \mathbb{1}_{(0.5,1.5)}(x)$  (right) with  $n = 1000$ . The true survival function  $S$  of an  $\Gamma_{4,0.5}$  distribution is given by the black curve while the red curve is the point-wise empirical median of the 50 estimates.

## Bibliographie

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