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Nonparametric estimation of splicing points in skewed cost distributions: a kernel-based approach

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ABSTRACT

This paper proposes a nonparametric method for detecting splicing points within cost distributions. Leveraging techniques from change point detection, we adopt an asymmetric gamma kernel estimator to identify these transition points in often highly skewed distributions. Our method focuses on identifying locations within the distribution where a significant change occurs, allowing for the application of distinct models on either side of the splicing point. Theoretical properties of the estimator, including strong consistency and asymptotic normality, are established. In particular, it is demonstrated that the estimator is consistent with a faster convergence rate than the parametric one. Since this estimator tends to underestimate the splicing point in finite samples, we propose a bias correction method to enhance accuracy. Our proposed approach is validated through simulations and real data applications.

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1. Introduction

Estimating the precise location within a distribution where significant structural changes occur is a critical task in many areas, including economics, actuarial science, and finance. Such points, often referred to as splicing points, mark transitions where the distribution may exhibit different characteristics that are better captured by distinct models on either side. For example, in the context of income distributions, this might correspond to a threshold beyond which a different regime or model is necessary to accurately describe the 'elite income' tier. In the field of non-life insurance, a handful of severe losses within a collection of policies beyond some threshold can significantly contribute to the total claim amount. In geophysics and hydrology, studies examine catastrophic disasters from the perspective of significant event analysis. It is widely recognised that a single model cannot capture characteristics over the entire range of such distributions. Typically, in income distributions, the segment above a certain threshold is modelled separately from the bulk of the data,

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which lies below this threshold; see Cowell et al. (1998) and Jenkins (2017), for instance. Moreover, modelling the whole range of a loss distribution is of particular importance and interest in actuarial science, particularly in non-life insurance. Then, splicing (e.g., Klugman et al. 2019) or composite modelling (e.g., Cooray and Ananda 2005; Scollnik and Sun 2012) – the practice of modelling the parts below and above a threshold differently – is frequently employed.

The aim of this paper is to propose a nonparametric method of finding a threshold in these distributions. Accordingly, the words 'threshold' and 'splicing point' are used exchangeably hereinafter, whenever no confusion may arise. Our estimation strategy is grounded on prior knowledge (or stylised facts) about shapes of the underlying distributions of interest. Our particular focus is on the distributions of nonnegative economic and financial variables including incomes, wages, consumption expenditures, short-term interest rates, and actuarial losses. These variables are examples of cost variables. Distributions of cost variables have support on \mathbb{R}_+ with a natural boundary at the origin and are highly right-skewed, with a concentration of observations near the origin and fewer data points as values increase. As will be revealed shortly, we employ a specific kernel to capture these characteristics while avoiding possible misspecification by relying on a particular parametric model. In addition, we translate the problem of threshold estimation into that of change point detection, which has been actively studied in statistics. Our translation or interpretation of the problem is motivated by the original idea of splicing, which refers to the technique of combining two different probability density functions (pdfs) for the two regions below and above a threshold or a splicing point, and continuity of the distribution at this point is not required.¹

Following the literature on change point detection (e.g., Chu and Cheng 1996; Couallier 1999; Huh 2002), we take the absolute difference of two kernel density estimates as the diagnostic function and define its maximiser as the splicing point estimator.² Nonetheless, our procedure remarkably differs from previous ones. While they are designed to work in the central part of a distribution (i.e. near the peak), a threshold that divides the bulk and less dense regions is expected to be situated far from the peak of the distribution. To overcome such difficulty while exploiting prior information on shapes of cost distributions, we adopt the asymmetric gamma kernel by Chen (2000); our contribution is the first work in which an asymmetric kernel is applied for the problem of splicing point estimation in composite models of density or regression curves, to the best of our knowledge. For a data point $u \in \mathbb{R}_+$, a design point $x \in \mathbb{R}_+$ and a smoothing parameter b > 0, the gamma kernel is defined as

$$K_{G(x,b)}(u) = \frac{u^{x/b} \exp\left(-u/b\right)}{b^{x/b+1} \Gamma\left(x/b+1\right)} \mathbf{1}\left\{u \ge 0\right\},\tag{1}$$

where $\Gamma(a) = \int_0^\infty t^{a-1} \exp(-t) dt$ for a > 0 is the gamma function, and $\mathbf{1}\{\cdot\}$ denotes an indicator function. A nonnegative random variable *Z* is said to obey the gamma distribution having the shape parameter $\alpha > 0$ and the scale parameter $\beta > 0$, which is denoted as $G(\alpha, \beta)$ in shorthand notation hereinafter, if its pdf is given by $f(z) = z^{\alpha-1} \exp(-z/\beta)\mathbf{1}\{z \ge 0\}/\{\beta^{\alpha}\Gamma(\alpha)\}$. Observe that the gamma kernel can be interpreted as the pdf of G(x/b + 1, b).

The use of asymmetric kernels, particularly the gamma kernel described in (1), represents a significant methodological contribution in our threshold estimation approach.

This kernel offers several advantages over traditional symmetric kernels that are worth highlighting. The gamma kernel provides adaptive smoothing by automatically changing shapes across design points while requiring only a single smoothing parameter to generate a variety of shapes. This property enables effective calibration of right-skewed densities with support on \mathbb{R}_+ , such as those typically found in cost distributions.

Beyond these initial advantages, the gamma kernel offers additional benefits that strengthen our methodology. First, the gamma kernel naturally respects the boundary constraints of the data, eliminating the boundary bias problem that occurs with symmetric kernels near the origin. Since many financial and actuarial data sets have a natural lower bound at zero, this property ensures more accurate density estimation in the critical region near this boundary. Second, the gamma kernel maintains optimal convergence rates in mean integrated squared error (MISE) within the class of nonnegative kernel density estimators (see, e.g., Chen 2000, p. 477). Third, the gamma kernel's adaptability to different degrees of data concentration is especially beneficial when dealing with unequally distributed data. Its variable bandwidth property automatically provides more smoothing in regions with sparse data (often in the tails) and less smoothing in regions with dense data. Fourth, the asymmetric nature of the gamma kernel aligns naturally with the inherent asymmetry present in skewed distributions. This alignment improves the accuracy of threshold identification by better capturing the underlying structure of the data. Fifth, our method benefits from the gamma kernel's capability to maintain nonnegative estimates throughout the support range, which is mathematically consistent with the nonnegative nature of the variables we typically analyze in cost distribution applications.

Our splicing point estimator is super-consistent (i.e. its convergence rate exceeds \sqrt{n} , where *n* is the sample size) and asymptotically normal when suitably implemented. While the estimator shares its asymptotic properties with the existing literature (e.g., Chu and Cheng 1996; Couallier 1999; Huh 2002), our proof strategy is totally different. The gamma kernel admits neither the location-scale transformation $K_h(u - x) = K\{(u - x)/h\}/h$ with a bandwidth h(> 0) nor exchangeability between u and x, unlike standard symmetric kernels. Instead, approximations to the incomplete gamma, digamma and polygamma functions studied by Funke and Hirukawa (2019, 2024) are tailored for the technical proofs. This proof strategy is novel and of independent interest; see the Appendix and/or the Supplemental Material for more details. It is also demonstrated that a uniform approximation to our diagnostic function has a unique maximum. So far existence of the maximum in the diagnostic function has been simply suggested, not formally proven, in the literature, although it is a key ingredient for consistency of the splicing point estimator. A concern is that the proposed estimator tends to generate negative biases in finite samples. However, an elementary bias correction can instantly improve bias properties of the estimator without inflating its variance, and thus we advocate putting the bias-corrected version to practical use.

The integration of asymmetric gamma kernels into threshold selection methodology represents a significant methodological innovation in the statistical literature, as it combines the boundary-respecting properties of asymmetric kernels with rigorous mathematical foundations to deliver improved convergence properties and bias reduction – a contribution that addresses fundamental challenges in optimal threshold detection for data with natural boundaries.

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Acknowledging that threshold estimation is a notoriously difficult problem, many authors have proposed various threshold detection methods, mainly in the field of nonlife insurance. Below a summary of different types of approaches is provided with emphasis on specific difficulties that arise. The approaches are divided roughly into three categories, namely, (i) heuristic approaches, (ii) graphical diagnostics and (iii) automated procedures. For (i), a threshold is defined as a fixed quantile (DuMouchel 1983) or determined by some formula depending on the sample size (Loretan and Phillips 1994); see Scarrott and Mac-Donald (2012) for more details. Despite no theoretical justification, these methods are used by actuaries in practical applications.

Examples of (ii) include the Hill plot and its variants, which were extensively analysed by Kratz and Resnick (1996) for their theoretical properties and stability. The mean excess plot (Davison and Smith 1990) and various quantile-based visualisation techniques systematically reviewed by Drees et al. (2020) have become standard tools in extreme value analysis. Complementing these approaches, Reiss and Thomas (2007) develop a procedure to identify regions of stability among extreme value index estimates, whereas Neves and Fraga Alves (2004) provide further analysis on the tuning parameters required for optimal threshold selection. For practical applications where sample size considerations are paramount, Ferreira et al. (2003) propose the formula $k = \sqrt{n}$ as a systematic approach to determining appropriate thresholds. These approaches are easy to grasp and thus used regularly, whereas there is room for practitioners' discretion at the stage of identifying a threshold.

Recent research has shifted toward (iii) such as the minimum Kolmogorov-Smirnov (KS) distance procedure (Clauset et al. 2009; Drees et al. 2020), sequential goodness-of-fit testing (e.g., Bader et al. 2018), and the minimum quantile discrepancy and automated Eye-Balling methods (Danielsson et al. 2019). A significant category within these procedures involves selecting thresholds based on goodness-of-fit of the generalised Pareto distribution (GPD), where the threshold is chosen as the lowest level above which the GPD provides adequate fit to the exceedances (Dupuis 1999; Choulakian and Stephens 2001; Northrop and Coleman 2014). These approaches employ various techniques, including comparing the empirical distribution to the fitted GPD via goodness-of-fit tests (Wadsworth 2016) or by minimising the distance between them (Pickands 1975; Gonzalo and Olmo 2004), with the latter approach theoretically analysed again by Drees et al. (2020). The KS and Anderson-Darling tests are commonly applied in this context. Another related method is the root mean square error (RMSE) approach by Li et al. (2014), which measures the difference between analytical and observed cumulative distribution functions (cdfs) of exceedances at different thresholds, with the threshold having the lowest RMSE considered optimal. While these approaches are conceptually straightforward, error control remains challenging due to the ordered nature of the hypotheses, with standard multiple testing methods like false discovery rate (Benjamini 2010a, 2010b) not being directly applicable. Another promising approach treats the data as a mixture of distributions, with a GPD for the tail and another distribution for the bulk joined at the threshold (Mac-Donald et al. 2011; Wadsworth and Tawn 2012; Naveau et al. 2016). By considering the threshold as a parameter to estimate, these methods account for uncertainty from threshold selection in inferences, although care is needed to ensure the bulk and tail models remain robust to misspecification. These procedures circumvent arbitrariness but rely on a certain parametric model of the tail part including GPD. There is also a class of theoretically

motivated procedures that target the optimal sample fraction for specific estimation tasks, such as the estimation of high probabilities (Hall and Weissman 1997) or the Hill estimator.

Beyond these categories, Langousis et al. (2016) identify several additional specialised approaches for threshold selection. These include methods based on asymptotic results about estimators of tail distribution properties, such as the Jackson or the Lewis kernel statistics (Jackson 1967 resp. Lewis 1965), modified by Goegebeur et al. (2008) to enhance the performance of the Hill estimator. Another significant approach is the automated version of the mean residual life (MRL) plot. For more complex scenarios, resampling-based estimators have been developed, although these are computationally demanding and often require practitioners to select tuning parameters (Danielsson et al. 2001), which may render them unsuitable for smaller sample sizes (Ferreira et al. 2003). Our threshold detection procedure will be able to serve as a more objective and flexible alternative to these methods.

Furthermore, there will be potentially many applications of our proposal. The definition of a 'cost' is not strict, actually. Our threshold estimation procedure is expected to work equally for non-cost variables (e.g., quantities demanded, transaction volumes, etc.), as long as shapes of their distributions have similarities to those of costs. It can be also employed for threshold detection prerequisite for tracing out the evidence of illegal trading (James et al. 2023) and implementing extreme changes in changes (CIC) estimation (Sasaki and Wang 2024), for instance.

The remainder of this paper is organised as follows. Section 2 overviews the splicing point estimation using the gamma kernel and then recommends its practical implementation. In Section 3, convergence properties of the proposed estimator, namely, strong consistency and asymptotic normality, are explored. Section 4 conducts Monte Carlo simulations to compare finite-sample behaviours of our splicing point estimator with those of several existing competitive estimation methods. Our aim is to show numerically the advantage of our proposal over these alternatives. In Section 5, the proposed estimation approach is applied to a couple of real world datasets. Section 6 concludes. Proofs of theorems and propositions are provided in the Appendix. Proofs of lemmata are deferred to the online Supplementary Material.

This paper adopts the following notational conventions: $a_n \sim b_n$ means that a_n/b_n converges to 1; $a_n = o(b_n)$ signifies that a_n/b_n converges to 0; $a_n = O(b_n)$ means that a_n/b_n is bounded; and we say that $a_n \simeq b_n$ if there exist constants $0 < c_1 < c_2 < \infty$ so that $c_1a_n \leq b_n \leq c_2a_n$. For a function h(x) and a point $c, h(c^-) = \lim_{x \to c} h(x), h(c^+) = \lim_{x \to c} h(x)$ and $h^{(m)}(x) = d^m h(x)/dx^m$ denote the left and right limits, and the *m*th-order derivative, respectively. The abbreviation *a.s.* stands for 'almost surely'. Finally, the expression 'X $\stackrel{d}{=}$ Y' reads 'A random variable X obeys the distribution Y.'

2. Our proposal: an informal overview

2.1. Estimation of a splicing point

It is suspected that f(x), the pdf of a 'cost' variable $X \in \mathbb{R}_+$, is discontinuous at t_0 on a prespecified closed interval $I_0 := [\underline{t}, \overline{t}]$ with $0 < \underline{t} < \overline{t} < \infty$. It is assumed that the interval I_0 is situated in the upper region of the underlying cost distribution. Prior knowledge on the interval is not at all unrealistic, because quite often practitioners have a rough idea about

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the location of the threshold through, for example, preliminary threshold estimates, historical experiences and/or empirical quantiles. Against this background, our method can complement existing practices and objectify the graphical analysis procedures mentioned in Section 1.

The model below basically follows those of Chu and Cheng (1996) and Couallier (1999). A similar local structure can be also found in threshold detection problems for nonparametric regression (e.g., Wu and Chu 1993a, 1993b; Joo and Qiu 2009) and deconvolution (e.g., Delaigle and Gijbels 2006). It is assumed that the pdf f(x) for $x \in I_0$ can be modelled locally as

$$f(x) = g(x) + d_0 \mathbf{1} \{ x < t_0 \},$$
(2)

where g(x) is a sufficiently smooth function, and $t_0 \in I_0$ is the splicing point (or threshold). The local structure (2) also implies the jump size

$$d_0 := f(t_0^-) - f(t_0^+),$$

where $|d_0| \in (0, \infty)$ is assumed throughout.

Our problem is how to estimate the threshold t_0 nonparametrically. Suppose that there are *n i.i.d.* observations $\{X_i\}_{i=1}^n$ at hand. If t_0 were located in the bulk region, as in the existing literature, splitting the entire sample into two sub-samples near t_0 would cause no serious issue. Two sample sizes are roughly the same, and thus both left and right limits of a density can be estimated equally well. This is clearly not the case in our problem. Because t_0 is located in the upper region, sample-splitting near t_0 results in imbalance in sample sizes of two sub-samples and an imprecise density estimate from the right sub-sample. In view of this, we use the entire sample to estimate both limits of the pdf. To do so, we introduce 'shifted' gamma kernels $K_{G(x,b;\pm\Delta)}(\cdot)$, which are defined as pdfs of gamma distributions $G\{(x \pm \Delta)/b + 1, b\}$, i.e.

$$K_{G(x,b;\pm\Delta)}(u) := \frac{u^{(x\pm\Delta)/b} \exp\left(-u/b\right)}{b^{(x\pm\Delta)/b+1} \Gamma\left\{\left(x\pm\Delta\right)/b+1\right\}} \mathbf{1}\left\{u\geq0\right\},$$

where $b(=b_n > 0)$ is the smoothing parameter, $\Delta(=\Delta_n > 0)$ plays the role of a shift parameter, and each parameter shrinks towards zero at a certain rate. Obviously, $K_{G(x,b;\pm\Delta)}(\cdot)$ collapse to Chen's (2000) original gamma kernel (1) when $\Delta = 0$. The kernels can be interpreted as those designed to smooth the data off the target design point x by a margin of Δ . In addition, they put the maximum weight at sightly left or right of x because they have their modes at $x \pm \Delta$.

Our threshold estimator is derived from the difference between two density estimates, which is generated by introducing a shift parameter Δ . Let the shifted density estimators be

$$\hat{f}^{\pm}(x) := \hat{f}_{b,\Delta}^{\pm}(x) := \frac{1}{n} \sum_{i=1}^{n} K_{G(x,b;\pm\Delta)}(X_i).$$

Also define

$$\hat{J}(x) := \hat{f}^{-}(x) - \hat{f}^{+}(x),$$

where a single, common value is chosen for the smoothing parameter b in both density estimates, as in Chu and Cheng (1996), Couallier (1999) and Huh (2002). Following these articles, we also utilise $|\hat{J}(x)|$ as the diagnostic function for threshold detection. The estimator of the splicing point t_0 , denoted as \hat{t} , is defined as the maximiser of $|\hat{J}(x)|$ on $x \in I_0$, i.e.

$$\hat{t} := \arg \max_{x \in I_0} \left| \hat{J}(x) \right|.$$

2.2. Recommended estimation procedure

Monte Carlo results in Section 4 indicate that \hat{t} tends to underestimate t_0 . However, it turns out that the negative bias can be alleviated substantially, with no additional cost of spread, by an elementary bias correction. In practice, we recommend the bias-corrected version of the proposed estimator

$$\tilde{t} := \hat{t} + b$$

for a suitably chosen smoothing parameter b. A theoretical foundation of the bias correction can be found in Remark 3.2, and its necessity is also visualised in Figure 1. Superior finite-sample properties of \tilde{t} over \hat{t} are confirmed in Section 4.

We can compute \tilde{t} in the following steps:

- (1) Prespecify the interval I_0 that is likely to cover the splicing point t_0 .
- (2) Put the shift parameter Δ = b^α for α = 0.70 and select the smoothing parameter b via the modified likelihood cross-validation method given by (11)–(12). Both the exponent α in Δ and the choice method for b are based on our judgments from the Monte Carlo study and real data examples.
- (3) Find the maximiser of $|\hat{J}(x)|$ on $x \in I_0$ and take it as \hat{t} .
- (4) Obtain the bias-corrected estimate $\tilde{t} = \hat{t} + b$ using the value of *b* selected in Step (2).

3. Large-sample properties of the splicing point estimator

In this section convergence properties of the estimator \hat{t} are documented. Our particular focus is on its consistency and asymptotic normality. In the course of this, we demonstrate existence of a unique maximum in a certain uniform approximation to $|\hat{J}(x)|$ on $x \in I_0$, which constitutes a key condition for consistency of \hat{t} . The asymptotic distribution of \hat{t} also hints that a simple form of its leading bias enables us to derive the bias-corrected estimator \tilde{t} .

3.1. Regularity conditions

Convergence results below rely on the fact that $|\hat{J}(x)|$ can be approximated by the difference between two incomplete gamma functions. To deliver the results, we impose the following regularity conditions.

Assumption 3.1: $\{X_i\}_{i=1}^n \in \mathbb{R}_+$ are *i.i.d.* random variables.

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Assumption 3.2:

- (i) The pdf f(x) is uniformly bounded on $x \in \mathbb{R}_+$.
- (ii) The local structure (2) holds, $g^{(2)}(x)$ is uniformly bounded on $x \in \mathbb{R}_+$, and $g^{(3)}(x)$ is Lipschitz continuous and bounded on $x \in I_0$.

Assumption 3.3: Tuning parameters *b* and Δ satisfy *b*, $\Delta \rightarrow 0$,

$$\frac{b^{3/4}}{\Delta} + \frac{\Delta}{b^{1/2+\delta_1}} + \frac{b^{1/2-4\delta_1}}{n^{1-\delta_2}\Delta^2} \to 0$$
(3)

for some arbitrarily small $\delta_1, \delta_2 > 0$, and

$$\frac{\ln n}{nb^{3/2-\kappa}} = O(1) \tag{4}$$

for some $\kappa \in [0, 1)$, as $n \to \infty$.

All these assumptions are standard for uniform approximations to asymmetric kernel estimators. Similar conditions can be found, for example, in Funke and Hirukawa (2024). It follows from Assumption 3.2(ii) that $f^{(1)}(t_0^-) = f^{(1)}(t_0^+)$. Two boundedness conditions on derivatives of the smoothed component $g(\cdot)$ also serve as important ingredients for approximations to $E\{\hat{J}^{(p)}(x)\}$ on $x \in I_0$ for p = 0, 1, 2. This type of condition has been often imposed in simulation studies on change point detection (e.g., Wu and Chu 1993a, 1993b; Chu and Cheng 1996).

Assumption 3.3 controls the shrinkage rates of tuning parameters b and Δ . The condition (3) draws the following important conclusions: (i) $b = o(\Delta)$; (ii) $b^{1/2} = o(\Delta^2/b)$; and (iii) $\Delta = o(\Delta^3/b^{3/2})$. These are frequently used to controls remainder terms in the asymptotic expansions. It also follows from $b = o(\Delta)$ and $\Delta = o(b^{1/2})$ that although the shift parameter Δ should shrink to zero more slowly than the smoothing parameter b, the convergence rate of Δ must not be too slow (or must be faster than $b^{1/2}$, to be more precise). Couallier (1999), for instance, also imposes a similar rate requirement. The condition (3) also implies that

$$\frac{\ln n}{nb^{1/2}} = \left(\frac{b^{1/2-4\delta_1}}{n^{1-\delta_2}\Delta^2}\right) \left(\frac{\ln n}{n^{\delta_2}}\right) \left(\frac{\Delta}{b^{1/2+\delta_1}}\right)^2 b^{6\delta_1} \to 0.$$

This result serves as a prerequisite for Proposition 3.1, as will be revealed shortly. The other condition (4) is an additional technical requirement for strong uniform consistency of \hat{t} .

3.2. Consistency

Below asymptotic properties of the splicing point estimator \hat{t} are explored. Our analysis starts from a uniform approximation to $\hat{f}^{\pm}(x)$ on I_0 , which is documented in the next proposition. To save space, we adopt the following shorthand notation whenever no confusion may arise: $K_x^{\pm}(u) = K_{G(x,b;\pm\Delta)}(u)$; $K_x(u) = K_{G(x,b)}(u)$; $a^{\pm} = (x \pm \Delta)/b$; and $z_0 = t_0/b$.

Proposition 3.1: If Assumptions 3.1–3.3 hold, then

$$\sup_{x \in I_0} \left| E\left\{ \hat{f}^{\pm}(x) \right\} - \left\{ g\left(x\right) \pm g^{(1)}(x) \Delta + d_0 \int_0^{t_0} K_x^{\pm}(u) \, \mathrm{d}u \right\} \right| = O\left(b\right)$$
(5)

and

$$\sup_{x \in I_0} \left| \hat{f}^{\pm} (x) - E\left\{ \hat{f}^{\pm} (x) \right\} \right| = O\left(\sqrt{\frac{\ln n}{nb^{1/2}}}\right) \quad a.s.,$$
(6)

as $n \to \infty$.

A direct outcome from Proposition 3.1 is that

$$\sup_{x\in I_0} \left| \hat{J}(x) - E\left\{ \hat{J}(x) \right\} \right| = O\left(\sqrt{\frac{\ln n}{nb^{1/2}}}\right) \quad a.s.$$

It also follows from $||\hat{J}(x)| - |E\{\hat{J}(x)\}|| \le |\hat{J}(x) - E\{\hat{J}(x)\}|$ that

$$\left|\hat{J}(x)\right| = \left|E\left\{\hat{J}(x)\right\}\right| + O\left(\sqrt{\frac{\ln n}{nb^{1/2}}}\right)$$
 a.s.

uniformly on I_0 . In short, $|E\{\hat{J}(x)\}|$ constitutes the dominant term in $|\hat{J}(x)|$, or the effect of the location x on the magnitude of $|\hat{J}(x)|$ appears only in the value of $|E\{\hat{J}(x)\}|$ in a first-order asymptotic sense. This result also plays a key role in the proof of Theorem 3.1 below; see the Appendix for more details.

It follows from (5) and Assumption 3.2(ii) that $|E\{\hat{J}(x)\}|$ can be further approximated by

$$\left| E\left\{ \hat{J}\left(x\right) \right\} \right| := \left| d_0 \right| J\left(x\right) + O\left(\Delta\right)$$
(7)

uniformly on I_0 , where

$$J(x) = \left| \int_0^{t_0} K_x^-(u) \, \mathrm{d}u - \int_0^{t_0} K_x^+(u) \, \mathrm{d}u \right|$$

= $\int_0^{t_0} K_x^-(u) \, \mathrm{d}u - \int_0^{t_0} K_x^+(u) \, \mathrm{d}u$
= $P\left(a^- + 1, z_0\right) - P\left(a^+ + 1, z_0\right),$

and $P(a, z) := \gamma(a, z) / \Gamma(a)$ is a normalised version of the lower incomplete gamma function $\gamma(a, z) = \int_0^z t^{a-1} \exp(-t) dt$ for a, z > 0. The reasons why $P(a^- + 1, z_0) \ge P(a^+ + 1, z_0)$ holds are that $P(a^{\pm} + 1, z_0) = \Pr(Y^{\pm} \le z_0)$ for $Y^{\pm} \stackrel{d}{=} G(a^{\pm} + 1, 1)$ (i.e. $P(a^{\pm} + 1, z_0)$ are cdfs of Y^{\pm} evaluated at z_0) and that the larger the shape parameter is, the flatter the gamma distribution becomes.

We are about to demonstrate strong consistency of \hat{t} . Before proceeding, it is curious whether J(x) on $x \in I_0$ indeed has a unique maximum at t_0 (or within a shrinking neighbourhood of t_0 even if it is not maximised exactly at this point). In reality, however, it is

quite cumbersome to look into the local property of J(x) analytically. Fortunately, several approximations to the incomplete gamma function are available, and we rely on one of them. More specifically, we employ Equation (1) of Pagurova (1965) to approximate the normalised lower incomplete gamma functions $P(a^{\pm} + 1, z_0)$ around the standard normal cdf. The next proposition refers to properties of the approximation and the maximiser of the approximated function.

Proposition 3.2: If Assumption 3.3 holds, then the followings hold true.

(i) Define

$$Q(x) := \left(\frac{x+t_0}{x^{3/2}}\right) \phi\left(\frac{x-t_0}{\sqrt{bx}}\right),$$

where $\phi(\cdot)$ is the pdf of N(0, 1). Then,

$$\sup_{x\in I_0} \left| J(x) - Q(x)\left(\frac{\Delta}{b^{1/2}}\right) \right| = O\left(\frac{\Delta^3}{b^{3/2}}\right),$$

as $n \to \infty$.

(ii) Q(x) on I_0 has a unique maximum at $x = t^* \in (t_0 - b, t_0)$.

Before establishing strong consistency of \hat{t} , we show by some numerical illustration that maximising $|\hat{J}(x)|$ is a well-defined problem. A discontinuous density f(x) and the diagnostic function $|\hat{J}(x)|$ are drawn in Panel (a) of Figure 1. Model 1-A in Section 4 is chosen for this illustration. The density is discontinuous at $t_0 = 4$ with a magnitude of discontinuity $d_0 = 0.15$. The diagnostic function is computed from a Monte Carlo sample of sample size 500 under tuning parameters b = 0.05 and $\Delta = b^{0.70}$.

Propositions 3.1 and 3.2 jointly imply that $|\hat{J}(x)| \sim |d_0|J(x) \sim |d_0|Q(x)(\Delta/b^{1/2})$ holds in theory. This result tempts us to make a visual inspection of shapes of these three curves. The curves around the true threshold $t_0 = 4$ are plotted in Panel (b) of Figure 1. Notice that the panel magnifies the area surrounding t_0 to visualise preciseness of the approximations. It can be immediately found that all three curves are single-peaked around the true threshold, which confirms well-definedness of the optimisation problem. Approximating $|\hat{J}(x)|$ by $|d_0|J(x)$ looks decent, whereas the discrepancy between the two curves suggests that the approximation errors which are asymptotically negligible may not be ignored in finite samples. Furthermore, $|d_0|Q(x)(\Delta/b^{1/2})$ approximates $|d_0|J(x)$ quite well; rather, they are almost indistinguishable. For reference, Panel (b) also indicates maximisers of $|\hat{J}(x)|$ and Q(x) are $\hat{t} \approx 3.8693$ and $t^* \approx 3.9502$, respectively. It follows from b = 0.05 that the latter confirms Proposition 3.2(ii).

The function Q(x) has the following properties. Observe that

$$b^{-1/2}Q(x) = \sqrt{\frac{t_0^2/b}{2\pi x^3}} \exp\left\{-\frac{\left(t_0^2/b\right)(x-t_0)^2}{2t_0^2 x}\right\} \left(\frac{x+t_0}{t_0}\right),$$

where $\sqrt{(t_0^2/b)/(2\pi x^3)} \exp\{-(t_0^2/b)(x-t_0)^2/(2t_0^2x)\}$ is the pdf of the inverse Gaussian distribution $IG(t_0, t_0^2/b)$. Because the shape parameter of this distribution $t_0^2/b \to \infty$, the



Figure 1. The diagnostic function and its approximations.

pdf is close to a normal one for a sufficiently small b > 0. In addition, the distribution has mean t_0 and variance bt_0 . It follows that the pdf roughly behaves like $N(t_0, bt_0)$, and thus, heuristically, the shape of Q(x) also looks like a bell curve centred around t_0 .

Strong consistency of \hat{t} for t_0 is formally delivered in the theorem below.

Theorem 3.1: Let $c_n := b^{1/2+\delta_1}$ for δ_1 defined in Assumption 3.3. If Assumptions 3.1–3.3 hold, then $|\hat{t} - t_0| = O(c_n)$ a.s. as $n \to \infty$.

The proof of Theorem 3.1 closely follows that of Theorem 3.1 in Chu and Cheng (1996); see the Appendix for more details. Some readers may wonder why Theorem 2.1 of Newey and McFadden (1994) is not employed. There are two reasons for not relying on this theorem. First, we can immediately see the followings: Q(x) is uniquely maximised at $x = t^*$; I_0 is compact; and Q(x) is continuous. The problem is that while $(b^{1/2}/\Delta)|\hat{f}(x)|$ is uniformly approximated by $|d_0|Q(x)$ on I_0 , Q(x) still depends on *n* through *b*. Therefore, Theorem 2.1 of Newey and McFadden (1994) is not directly applicable. Second, while Theorem 2.1 of Newey and McFadden (1994) can lead to (weak) consistency $\hat{t} \stackrel{P}{\to} t_0$, it says nothing about the convergence rate. In contrast, Theorem 3.1 above and $c_n = o(b^{1/2})$ jointly establish that $|\hat{t} - t_0| = o_p(b^{1/2})$. As will be seen in Lemma A.8 in the Appendix, the weak consistency of \hat{t} with this rate plays a key role in establishing the asymptotic normality of \hat{t} .

3.3. Asymptotic normality

The theorem below documents asymptotic normality of \hat{t} . The asymptotic distribution is derived indirectly, as in Chu and Cheng (1996, Theorem 1), Couallier (1999, Théorème 2) and Delaigle and Gijbels (2006, Theorem 3.1). The indirect derivation comes from the fact that \hat{t} solves the first-order condition $\hat{f}^{(1)}(\hat{t}) = 0$. Then, a mean-value expansion of the left-hand side around $\hat{t} = t_0$ is made, and suitable approximations to the incomplete gamma, digamma and polygamma functions are utilised in the expansion; see the Supplemental Material for more details. This is possible because unlike f(x), its estimates $\hat{f}^{\pm}(x)$ are smooth functions even at t_0 due to differentiability of shifted gamma kernels $K_x^{\pm}(\cdot)$ with respect to x.

Theorem 3.2: If Assumptions 3.1–3.3 hold, then

$$\sqrt{\frac{n}{b^{1/2}}}\left\{\hat{t} - t_0 - (-b)\right\} \stackrel{d}{\to} N\left(0, V_0\right) := N\left(0, \frac{3\sqrt{\pi}t_0^{1/2}}{4d_0^2}\left\{\frac{f(t_0^-) + f(t_0^+)}{2}\right\}\right)$$

as $n \to \infty$.

Remark 3.1: While it is difficult to obtain asymptotic bias and variance of \hat{t} in light of the indirect nature, the asymptotic distribution in Theorem 3.2 implies the first two moments of \hat{t} . The dominant bias term of \hat{t} is -b regardless of the position of t_0 . The expression of the term is much simpler than what is obtained by Couallier (1999, Théorème 2). The difference arises from different assumptions on the local structure of the pdf f on I_0 ; invoke that our Assumption 3.2(ii) follows the setup by Chu and Cheng (1996). As discussed in Remark 3.2 below, simplicity of the dominant bias term enables us to make the bias correction of \hat{t} straightforward. Moreover, V_0 , the coefficient of the dominant variance term, suggests that the larger the magnitude of discontinuity $|d_0|$, the easier the estimation of t_0 . It can be also recognised that the farther t_0 moves away from the origin, the less precise its estimator becomes. Finally, both bias and variance terms are free of the shift parameter Δ . A similar result is obtained in Théorème 2 of Couallier (1999); to put it another way, Δ does not affect convergence properties of \hat{t} in a first-order asymptotic sense.

Remark 3.2: As will be seen in the next section, \hat{t} tends to yield negative biases in finite samples, i.e. it is likely to underestimate the location of the splicing point, which coincides with what Theorem 3.2 predicts. However, the theorem also suggests that the bias can be corrected straightforwardly by adding b to \hat{t} . This is the foundation of the bias-corrected estimator $\tilde{t} = \hat{t} + b$ described in Section 2.2. Indeed, the proofs of Lemma A.6 and Theorem 3.2 jointly imply that the leading bias of \tilde{t} is $O(\Delta^2)$, whereas its variance is still $O(b^{1/2}/n)$. It will be confirmed in the Monte Carlo study shortly that \tilde{t} is a remedy for better finite-sample properties.

Remark 3.3: An approximation to the mean squared error (AMSE) of \hat{t} is

AMSE
$$(\hat{t}) = b^2 + \frac{b^{1/2}}{n}V_0 = O\left(b^2 + \frac{b^{1/2}}{n}\right),$$
 (8)

where $O(b^2)$ and $O(b^{1/2}/n)$ terms are leading squared bias and variance of \hat{t} , respectively. The AMSE for the threshold parameter implied by Théorème 2 of Couallier (1999) is in the form of $O(h^4 + h/n)$, where h is the the bandwidth for standard symmetric kernels. It can be found that this AMSE and (8) are of the same order of magnitude by recognising that $b \simeq h^2$. Furthermore, it follows from (8) that no bias-variance trade-off occurs, because a smaller b makes both squared bias and variance terms smaller.

Remark 3.4: Some readers may wonder how to pick *b* and Δ for super-consistency of \hat{t} . Then, for arbitrarily small $\delta_1, \delta_2 > 0$ as given in Assumption 3.3, put $\Delta \simeq b^{\alpha}$ for some $\alpha \in (1/2 + \delta_1, 3/4)$ and $b \simeq n^{-\beta}$ for some $\beta \in (0, (1 - \delta_2)/(2\alpha - 1/2 + 4\delta_1))$. It

is straightforward to see that such Δ and b jointly satisfy (3). In addition, when $\alpha \in (1/2, 3/4)$, we have

$$\frac{1-\delta_2}{1+4\delta_1} < \frac{1-\delta_2}{2\alpha - 1/2 + 4\delta_1} < \frac{2(1-\delta_2)}{1+12\delta_1},$$

where the two bounds $(1 - \delta_2)/(1 + 4\delta_1)$ and $2(1 - \delta_2)/(1 + 12\delta_1)$ are slightly below 1 and 2, respectively. Using this, we may draw the following three conclusions on the convergence rate of \hat{t} :

- (1) We are always allowed to pick $\beta > 1/2$. Then, $AMSE(\hat{t}) = o(n^{-1})$, or \hat{t} becomes super-consistent.
- (2) It is even possible to set $\beta = 2/3$, in particular. This value balances orders of magnitude in the squared bias and variance so that $O(b^2) = O(b^{1/2}/n) = O(n^{-4/3})$. As a consequence, $AMSE(\hat{t}) = O(n^{-4/3})$. It is also clear that the AMSE convergence rate of \hat{t} is determined by the exponent β . $AMSE(\hat{t}) = O(b^2)$ (i.e. the squared bias dominates) for $\beta \le 2/3$, and $AMSE(\hat{t}) = O(b^{1/2}/n)$ (i.e. the squared bias becomes asymptotically negligible) otherwise. The latter case corresponds to an 'undersmoothing' scenario so that $nb^{3/2} \rightarrow 0$ holds. As a consequence, the asymptotic normality statement in Theorem 3.2 reduces to $\sqrt{n/b^{1/2}}(\hat{t} t_0) \stackrel{d}{\rightarrow} N(0, V_0)$.
- (3) The best possible rate is $AMSE(\hat{t}) = O(n^{-2+\varepsilon})$ for an arbitrarily small $\varepsilon > 0$. The
- rate can be attained by setting α and β slightly above 1/2 and slightly below 2, respectively. Chu and Cheng (1996) and Couallier (1999) also report that their threshold estimators can attain the same convergence rate under the best case scenario.

Furthermore, it is not hard to see that for Δ and b defined above, we can always find some $\kappa \in [0, 1)$ satisfying (4). To see this, observe that (4) holds if $nb^{3/2-\kappa} \to \infty$ at a polynomial rate. The rate requirement is attained for conclusion (1) by setting β slightly above 1/2 and $\kappa = 0$. For conclusion (2), $\beta = 2/3$ and any $\kappa \in (0, 1)$ can jointly establish a polynomial divergence of $nb^{3/2-\kappa}$. Finally, for conclusion (3), β slightly below 2 and κ slightly below 1 lead to $nb^{3/2-\kappa} \to \infty$ at a polynomial rate.

Remark 3.5: As long as f(x) can be locally modelled as or well-approximated by (2), both \hat{t} and \tilde{t} become super-consistent when implemented as in Remark 3.4. It follows that regardless of whether a parametric (e.g., GPD) or nonparametric model (e.g., Markovitch and Krieger 2000) is fitted to the upper part, our splicing point estimator can be safely used as a threshold estimate without deteriorating the convergence rate for the model of the upper part of the distribution. In addition, Table 1 of Clauset et al. (2009) lists examples of non-power law distributions that behave like the GPD. Clauset et al. (2009) even argue that fitting a power law distribution in their procedure has nothing to do with a plausible match of the distribution with the data, and they recommend a goodness-of-fit test as a post-estimation analysis. Super-consistency of our estimators does no harm to convergence rates of the test statistics, either.

4. Finite-sample performance

4.1. Monte Carlo design

We consider two alternative models in the simulation study. For each model, 1000 Monte Carlo replications of $\{X_i\}_{i=1}^n$ with sample size $n \in \{250, 500\}$ are simulated.

In the first case, the univariate random variable $X \in \mathbb{R}_+$ is drawn from a *log-normal*-like distribution. What differs from a usual log-normal distribution is that a quadratic term is added to the pdf on the interval $[0, t_0) = [0, 4)$. Specifically, the pdf f(x) is

$$f(x) = \left\{\frac{1}{1 + (2/3)Dt_0}\right\} \left[\frac{1}{x\sigma\sqrt{2\pi}}\exp\left\{-\frac{(\ln x - \mu)^2}{2\sigma^2}\right\} + S(x)\right], (\mu, \sigma) = \left(\frac{1}{5}, \frac{3}{4}\right),$$

where

$$S(x) := D\left\{1 - \left(\frac{x - t_0}{t_0}\right)^2\right\} \mathbf{1} \{x < t_0\}$$

and

$$d_0 = f(t_0^-) - f(t_0^+) = \frac{D}{1 + (2/3) Dt_0}.$$

The shift parameter *D* takes two values, and D = 1/4, 3/22 yield $d_0 = 0.15$, 0.10, respectively. The former and latter cases are labelled as 'Model 1-A' and 'Model 1-B'. It also follows from $f^{(1)}(t_0^-) = f^{(1)}(t_0^+)$ that (2) is satisfied in the neighbourhood of the splicing point $t_0 = 4$.

This design is in some sense similar to that of Chu and Cheng (1996), who consider the density f(x) constructed by splicing left and right sides of two normal distributions with zero mean but different variances. In their design, $f(0^-) \neq f(0^+)$ but $f^{(1)}(0^-) = f^{(1)}(0^+)$, and thus (2) holds in the neighbourhood of the origin (aside from the fact that the threshold lies in the middle part).

In the second case, the nonnegative random variable *X* is generated by some distribution spliced at $t_0 = 4$. The pdf f(x) in this case takes the general form

$$f(x) = f_L(x) \mathbf{1} \{x < t_0\} + (1 - c_L) f_R(x) \mathbf{1} \{x \ge t_0\},\$$

where $f_L(x)$ is some density function truncated at t_0 , $f_R(x)$ is another density function with support on $[t_0, \infty)$, and $c_L := \int_0^{t_0} f(x) \, dx = \int_0^{t_0} f_L(x) \, dx$ ensures unity of the integral of f(x) over its entire support \mathbb{R}_+ ; in other words, $f_L(x)$ and $f_R(x)$ represent bulk and upper part models, respectively. This scenario is labelled as 'Model 2'.

Throughout the Weibull distribution with density

$$f_L(x) = \frac{\kappa}{\lambda} \left(\frac{x}{\lambda}\right)^{\kappa-1} \exp\left\{-\left(\frac{x}{\lambda}\right)^{\kappa}\right\}, \quad (\kappa, \lambda) = \left(3, \frac{11}{4}\right),$$

is considered as the bulk part of Model 2. Densities of the following distributions are examined for the tail part, and three cases are denoted as 'Model 2-A', 'Model 2-B' and 'Model 2-C', depending on the corresponding tail model:

$$f_{R}(x) = \begin{cases} \frac{1}{s} \left\{ 1 + \frac{\zeta(x-t_{0})}{s} \right\}^{-(1+1/\zeta)} \mathbf{1} \left\{ x \ge t_{0} \right\}, (\zeta, s) = \left(\frac{1}{4}, 4\right) & [A: GPD] \\ \frac{k}{\ell} \exp\left\{ \left(\frac{1}{\ell}\right)^{k} \right\} \left(\frac{x-t_{0}+1}{\ell}\right)^{k-1} \exp\left\{ - \left(\frac{x-t_{0}+1}{\ell}\right)^{k} \right\} \mathbf{1} \left\{ x \ge t_{0} - 1 \right\}, \\ (k,\ell) = \left(\frac{1}{4}, 1\right) & [B: Translated Weibull] \\ \frac{1}{\varsigma} \sqrt{\frac{2}{\pi}} \exp\left\{ - \frac{(x-t_{0})^{2}}{2\varsigma^{2}} \right\} \mathbf{1} \left\{ x \ge t_{0} \right\}, \varsigma = 4\sqrt{\frac{2}{\pi}} \quad [C: Half-Normal] \end{cases}$$

where $f(t_0^-) = f_L(4)$ and $f(t_0^+) = (1 - c_L)f_R(4)$.

Drees et al. (2020, p.83) argue that discontinuity of the density at the threshold is an easy scenario for the threshold detection method by Clauset et al. (2009). Model 2-A is most favourable to existing threshold detection methods because of the chosen level of discontinuity. The tail part in Model 2-B, also known as a stretched exponential distribution, reflects that again it seemingly behaves like a power law distribution (Clauset et al. 2009). Because this pdf is unbounded at the boundary of the support $t_0 - 1$, it is truncated at t_0 . Model 2-C adopts a (shifted) half-normal distribution for the tail part. This distribution has a normal-type thin tail. Because our splicing point estimator is grounded on no particular parametric model for the tail part, we are curious to see how tail thickness influences finite-sample properties of our estimator.

Each of three cases violates (2) by construction but may be more realistic, because it is hard to judge whether the local structure (2) indeed holds in real data. Moreover, the bulk model and the jump size $d_0 = f(t_0^-) - f(t_0^+)$ are common across three cases of Model 2. The only difference is the tail modelling. In short, Model 2 is designed to investigate how robust our estimator is against violation of an important regularity condition and how our estimator behaves toward different tail models.

Table 1 presents the mode of the distribution of *X*, the constant c_L , left and right limits of the density at the splicing point $f(t_0^{\pm})$, and the jump size $d_0 = f(t_0^{-}) - f(t_0^{+})$. More than 95% of observations concentrate on the interval $[0, t_0)$ (i.e. in the bulk region) in each model, and all models but Model 2-C have polynomially decaying tails. These features reasonably mimic properties of cost distributions.

Our estimation procedure for t_0 is implemented as follows. There are two optimizations required, namely, (i) the one for tuning parameters (b, Δ) and (ii) the other for the search of the splicing point. For (i), Remark 3.4 suggests $\alpha \in (1/2, 3/4)$, and thus we restrict our attention to four values, namely, $\alpha \in \{0.55, 0.60, 0.65, 0.70\}$. A few crossvalidation (CV) methods for *b* are investigated, and their details are deferred to the next section. For each CV method, candidates of *b* are taken from 100 equally-spaced grids over the interval [0.005, 0.500]. For (ii), after (b, Δ) are determined, the threshold location

М	odel	Distribution	Mode	CL	$f(t_0^-)$	$f(t_{0}^{+})$	d ₀
1	А	Log-Normal + Quadratic ($D = 1/4$)	0.7499	0.9659	0.1728	0.0228	0.1500
	В	Log-Normal + Quadratic ($D = 3/22$)	0.7240	0.9583	0.1279	0.0279	0.1000
2	Α	Splicing with Weibull & GPD	2.4023	0.9539	0.1063	0.0115	0.0948
	В	Splicing with Weibull & Translated Weibull	2.4023	0.9539	0.1063	0.0115	0.0948
	С	Splicing with Weibull & Half-Normal	2.4023	0.9539	0.1063	0.0115	0.0948

Table 1. Characteristic numbers of underlying distributions.

is searched via a numerical optimisation routine for the diagnostic function $|\hat{J}(x)|$ on the interval $I_0 = [3, 5]$.³ Once the splicing point estimator \hat{t} is obtained as the maximiser of $|\hat{J}(x)|$, the bias-corrected estimator is computed as $\tilde{t} = \hat{t} + b$.

Finite-sample performances of \hat{t} and \tilde{t} are compared with those of existing (i) kernelsmoothed competitive threshold estimation procedures and (ii) automated threshold detection methods. For (i), we focus on the procedure by Chu and Cheng (1996) [CC]. The CC diagnostic function is $|\hat{J}_{CC}(x)| = |\hat{f}_1(x) - \hat{f}_2(x)|$, where $\hat{f}_j(x) = (nh)^{-1} \sum_{i=1}^n K_j\{(X_i - x)/h\}$, j = 1, 2, for the kernels K_1 and K_2 to be specified shortly and a common bandwidth h(> 0). As in our method, the maximiser of $|\hat{J}_{CC}(x)|$ on I_0 is defined as the threshold estimator. The kernels K_1 and K_2 are fourth-order polynomial ones. These are

$$K_1(u) = (0.4857 - 3.8560u + 2.8262u^2 + 19.1631u^3 + 11.9952u^4) \times \mathbf{1} \{ u \in [-1, 0.2012] \},\$$

and $K_2(u) = K_1(-u)$ for all u. These kernels are also employed for threshold detection in nonparametric regression curves by Wu and Chu (1993a, 1993b). The CC procedure is implemented as in ours. After the bandwidth value is found via grid search for a CV criterion in the next section, the threshold location is searched via numerical optimisation for the diagnostic function $|\hat{J}_{CC}(x)|$ on I_0 .

For (ii), we investigate the followings: (a) the minimum KS distance procedure between the empirical and GPD-based distribution functions by Clauset et al. (2009) [KS]; (b) the minimum quantile discrepancy criterion for the mean absolute deviation between empirical and GPD-based quantiles by Danielsson et al. (2019) [Q-MAD]; (c) the minimum quantile discrepancy criterion for the sup-norm between empirical and GPD-based quantiles by Danielsson et al. (2019) [Q-SUP]; (d) the automated Eye-Balling method based on tail index estimates by Danielsson et al. (2019) [AEB]; and (e) the Anderson-Darling sequential testing procedure by Bader et al. (2018) [ADST]. For (e), candidates of thresholds are 20 empirical percentiles from 50.0% until 97.5% with an increment of 2.5%, i.e. $\{50.0\%, 52.5\%, \ldots, 95.0\%, 97.5\%\}$. The 5% level of significance is used for testing, and *p*-values for multiple tests are adjusted by the ForwardStop procedure.

All simulations are conducted on R. In particular, R-packages poweRlaw, tea and eva are employed to implement automated threshold detection methods (a), (b)–(d) and (e), respectively.

4.2. Smoothing parameter selection

Selecting the smoothing parameter *b* is the most important practical issue. In our context, values of (b, Δ) must be determined before threshold location search so that the diagnostic function can be fixed on I_0 . However, Remark 3.3 does not help resolve this issue. There is no optimal choice for *b* on the basis of the bias-variance trade-off. Theorem 3.2 provides no guidance for Δ , either, because it does not automatically guarantee that any Δ satisfying (3) works equally well in finite samples. Furthermore, to the best of our knowledge, there is no decisive conclusion on selecting the tuning parameter in the context of threshold estimation; in fact, Chu and Cheng (1996) adopt fixed bandwidths in their Monte Carlo study.

Taking the dependence of Δ on *b* into account, we tailor Huh's (2012) approach to construct a few CV criterion functions. Before proceeding, put $\Delta = b^{\alpha}$ for a given α . Accordingly, $\hat{f}^{\pm}(x)$ are rewritten as $\hat{f}_{b}^{\pm}(x;\alpha)$, which signify the dependence of density estimates on (b, α) . Also let

$$\hat{f}_{b,-i}^{\pm}\left(x;\alpha\right) := \frac{1}{n-1} \sum_{j=1, j \neq i}^{n} K_{x}^{\pm}\left(X_{j}\right)$$

be density estimates using the sample with the *i*th observation eliminated. Finally, denote the number of observations falling into I_0 as $n_0 := \sum_{i=1}^n \mathbf{1}\{X_i \in I_0\}$.

The CV criterion function by Huh (2012) is defined as the sum of CV criteria for two density estimates that construct the diagnostic function for threshold location estimation. We incorporate this idea into three CV criterion functions. The minimiser of each criterion function is taken as the corresponding CV smoothing parameter. The first one is the least-squares cross-validation (LSCV) criterion. It is defined as

$$CV_{LS}(b;\alpha) = CV_{LS}^{-}(b;\alpha) + CV_{LS}^{+}(b;\alpha), \qquad (9)$$

where

$$CV_{LS}^{\pm}(b;\alpha) := \int_{I_0} \left\{ \hat{f}_b^{\pm}(x;\alpha) \right\}^2 \, dx - \frac{2}{n_0} \sum_{i:X_i \in I_0} \hat{f}_{b,-i}^{\pm}(X_i;\alpha) \,. \tag{10}$$

The remaining two criteria are likelihood-based ones. One is the simple likelihood cross-validation (LCV) criterion, which is analogous to $\hat{L}_2(h)$ of Marron (1985) and equation (2.1) of Van Es (1991). It is given by

$$CV_L(b;\alpha) = CV_L^-(b;\alpha) + CV_L^+(b;\alpha),$$

where

$$CV_{L}^{\pm}(b;\alpha) := -\sum_{i:X_{i} \in I_{0}} \ln \left\{ \hat{f}_{b,-i}^{\pm}(X_{i};\alpha) \right\}$$

is the negative log-likelihood. The other is the modified LCV (MLCV) criterion, which corresponds to $\hat{L}_5(h)$ of Marron (1985) and equation (2.2) of Van Es (1991). It takes the form of

$$CV_{ML}(b;\alpha) = CV_{ML}^{-}(b;\alpha) + CV_{ML}^{+}(b;\alpha), \qquad (11)$$

where

$$CV_{ML}^{\pm}(b;\alpha) := -\left[\sum_{i:X_i \in I_0} \ln\left\{\hat{f}_{b,-i}^{\pm}(X_i;\alpha)\right\} - \sum_{i=1}^n \int_{I_0} K_{X_i}^{\pm}(u) \, du\right]$$
$$= -\left[\sum_{i:X_i \in I_0} \ln\left\{\hat{f}_{b,-i}^{\pm}(X_i;\alpha)\right\} - \sum_{i=1}^n \left\{P\left(\frac{X_i \pm \Delta}{b} + 1, \frac{\bar{t}}{b}\right) - P\left(\frac{X_i \pm \Delta}{b} + 1, \frac{\bar{t}}{b}\right)\right\}\right], \quad (12)$$

and the second term is intended to eliminate the endpoint effect of the interval $I_0 = [\underline{t}, \overline{t}]$. Corresponding threshold location estimates are labelled 'SG-LS', 'SG-L' and 'SG-ML', where 'SG' abbreviates 'shifted gamma'. For their bias-corrected versions, we put '-BC' at the end.

Finally, choosing the bandwidth *h* is also required to implement the CC procedure. The LSCV analogous to (9)–(10) (i.e. finding a minimiser of the sum of two LSCV criteria for $\hat{f}_1(x)$ and $\hat{f}_2(x)$) is adopted.

4.3. Results

Table 2 presents several performance measures of threshold estimators. These include the bias, standard deviation and RMSE of each threshold estimator over 1000 Monte Carlo samples. In addition, for CC and SG estimators, Monte Carlo averages and standard deviations (in parentheses) of CV tuning parameters are reported for reference. Furthermore, there is no guarantee that automated threshold detection methods necessarily yield threshold estimates falling into the interval I_0 . For these methods, percentages of threshold estimates inside I_0 out of 1000 Monte Carlo samples are provided.

Properties of the tail parts in Model 2-C and other four models differ; while the tail for the former decays exponentially fast, the latter has a polynomially decaying tail. Therefore, it is reasonable to evaluate the Monte Carlo results from the former and the latter separately.

4.3.1. Models 1-A, 1-B, 2-A, and 2-B

We start from examining the results from automated threshold detection methods. Q-MAD generates the smallest RMSE for Models 1-A, 1-B and 2-A, whereas ADST yields the smallest RMSE for Model 2-B. In addition, more than 90% of estimates from Q-MAD are inside I_0 for each model and sample size, despite no restriction on the parameter space. While Models 2-A and 2-B are thought to be more favourable than Models 1-A and 1-B for these automated methods, Q-MAD is comparable to SG methods in terms of RMSE for the latter case. There are also general tendencies of underestimation by KS and Q-MAD and overestimation by AEB. No clear tendencies can be observed in ADST. In particular, the degree of overestimation by AEB is considerable.

CC also looks comparable to SG methods. It consistently overestimates the location of the splicing point. However, the bias reduces with the sample size, and jointly by the decrease in dispersion, its RMSE becomes smaller as the sample size is larger.

There is also a general tendency in the results from SG methods. It can be immediately found that the initial estimate \hat{t} tends to be negatively biased, as Theorem 3.2 predicts. Moreover, a short b makes both the bias and variance small, as suggested in Remark 3.3. Accordingly, it can be reasonably conjectured that a CV algorithm that can generate a small smoothing parameter value \hat{b} will contribute to the initial estimate \hat{t} with good quality. Clearly, MLCV alone fulfills this requirement. In contrast, LSCV and LCV consistently give a large value of \hat{b} , which leads to a considerably biased \hat{t} . This is the source of their poor performance, and the sizable bias cannot be corrected completely even after adding a large \hat{b} to \hat{t} .

In what follows, we look into SG-ML and SG-ML-BC more carefully. Although the shift parameter Δ does not enter the asymptotic normality result in Theorem 3.2, the choice of

s.
s.

				<i>n</i> =	250				<i>n</i> =	500			
Estimator	α	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }
					Model 1-	A: Log-Normal	+ Quadratic (L	D = 1/4					
CC	_	0.2260	0.6340	0.6731	0.0116	(0.0092)	_ `	0.0487	0.5775	0.5796	0.0073	(0.0037)	_
KS	_	-0.8770	0.5392	1.0295	_	(-)	64.4%	-0.5234	0.7046	0.8778	_	(-)	70.7%
Q-MAD	_	-0.2990	0.1569	0.3376	_	(–)	100.0%	-0.3229	0.1962	0.3778	_	(_)	99.7%
Q-SUP	_	-0.1039	0.4014	0.4146	_	(–)	97.6%	0.0800	0.6708	0.6755	_	(_)	91.3%
AEB	_	0.6615	0.8305	1.0618	_	(–)	72.9%	0.7181	0.5745	0.9197	_	(–)	70.8%
ADST	_	-0.2662	0.7930	0.8365	_	(-)	78.1%	0.2605	0.5359	0.5958	_	(-)	90.2%
SG-LS	0.55	-0.6680	0.4375	0.7985	0.3302	(0.212)	_	-0.7461	0.4079	0.8503	0.3716	(0.2027)	_
	0.60	-0.6578	0.4423	0.7927	0.3263	(0.2206)	_	-0.7351	0.4149	0.8441	0.3666	(0.2066)	_
	0.65	-0.6501	0.4444	0.7874	0.3221	(0.2220)	_	-0.7185	0.4243	0.8345	0.3585	(0.2113)	_
	0.70	-0.6333	0.4497	0.7767	0.3141	(0.2243)	_	-0.7053	0.4310	0.8266	0.3518	(0.2147)	_
SG-LS-BC	0.55	-0.3378	0.2250	0.4058	_	(-)	_	-0.3745	0.2078	0.4283	_	(-)	_
	0.60	-0.3315	0.2273	0.4020	_	(-)	_	-0.3685	0.2107	0.4245	_	(-)	_
	0.65	-0.3280	0.2282	0.3996	_	(-)	_	-0.3600	0.2155	0.4196	_	(-)	_
	0.70	-0.3192	0.2313	0.3942	_	(-)	_	-0.3535	0.2192	0.4160	_	(-)	_
SG-L	0.55	-0.5628	0.4462	0.7182	0.2725	(0.2076)	_	-0.6063	0.4269	0.7415	0.2914	(0.1987)	_
	0.60	-0.5118	0.4494	0.6811	0.2449	(0.2052)	_	-0.5363	0.4320	0.6886	0.2548	(0.1989)	_
	0.65	-0.4758	0.4468	0.6527	0.2262	(0.2011)	_	-0.4898	0.4288	0.6510	0.2311	(0.1939)	_
	0.70	-0.4590	0.4433	0.6381	0.2160	(0.1966)	_	-0.4608	0.4219	0.6247	0.2148	(0.1877)	_
SG-L-BC	0.55	-0.2904	0.2474	0.3814	_	(-)	_	-0.3149	0.2357	0.3933	_	(-)	_
	0.60	-0.2668	0.2546	0.3688	_	(-)	_	-0.2814	0.2417	0.3710	_	(-)	_
	0.65	-0.2497	0.2575	0.3586	_	(-)	_	-0.2587	0.2437	0.3554	_	(-)	_
	0.70	-0.2430	0.2598	0.3557	_	(-)	_	-0.2460	0.2437	0.3463	_	(-)	_
SG-ML	0.55	0.1375	0.4524	0.4728	0.0315	(0.0210)	_	0.3265	0.5146	0.6094	0.0249	(0.0183)	_
	0.60	0.0465	0.3836	0.3864	0.0335	(0.0196)	_	0.1547	0.4346	0.4613	0.0297	(0.0173)	_
	0.65	-0.0035	0.3204	0.3204	0.0351	(0.0188)	_	0.0608	0.3497	0.3550	0.0326	(0.0166)	_
	0.70	-0.0352	0.2737	0.2759	0.0364	(0.0187)	_	0.0101	0.2831	0.2832	0.0342	(0.0157)	_
SG-ML-BC	0.55	0.1690	0.4378	0.4693	_	(-)	_	0.3514	0.4993	0.6105	_	(-)	_
	0.60	0.0800	0.3714	0.3799	_	(-)	_	0.1845	0.4216	0.4602	_	(-)	_
	0.65	0.0316	0.3098	0.3114	_	(-)	_	0.0933	0.3390	0.3516	_	(-)	_
	0.70	0.0011	0.2639	0.2639	-	(-)	-	0.0443	0.2741	0.2777	-	(-)	-
												(cor	ıtinued).

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Table 2. Continued.	
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				<i>n</i> =	250					<i>n</i> =	500		
Estimator	α	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }
					Model 1-	-B: Log-Normal	+ Quadratic (D = 3/22)					
CC	_	0.3200	0.6223	0.6998	0.0139	(0.0126)	_	0.1409	0.5942	0.6107	0.0086	(0.0056)	_
KS	_	-0.9832	0.6012	1.1525	_	(-)	59.4%	-0.6664	0.6955	0.9632	_	(-)	66.0%
Q-MAD	_	-0.3227	0.2688	0.4200	_	(-)	99.3%	-0.3638	0.2716	0.4540	_	(-)	99.5%
Q-SUP	_	-0.0720	0.6380	0.6421	_	(-)	92.5%	0.1549	0.9658	0.9782	_	(-)	83.7%
AEB	_	0.9418	0.8936	1.2983	_	(-)	60.2%	1.0100	0.6164	1.1832	_	(-)	53.5%
ADST	_	-0.6039	1.0622	1.2219	_	(-)	56.5%	0.0711	0.9103	0.9131	_	(-)	72.3%
SG-LS	0.55	-0.9517	0.2014	0.9727	0.4740	(0.1070)	_	-0.9940	0.0711	0.9965	0.4967	(0.0392)	_
	0.60	-0.9472	0.2109	0.9703	0.4717	(0.1115)	_	-0.9923	0.0808	0.9956	0.4958	(0.0444)	_
	0.65	-0.9363	0.2304	0.9642	0.4658	(0.1220)	_	-0.9914	0.0856	0.9951	0.4953	(0.0468)	_
	0.70	-0.9303	0.2398	0.9607	0.4625	(0.1273)	_	-0.9856	0.1130	0.9921	0.4925	(0.0590)	_
SG-LS-BC	0.55	-0.4777	0.0973	0.4875	_	(-)	_	-0.4973	0.0321	0.4983	_	(-)	_
	0.60	-0.4755	0.1029	0.4865	_	(-)	_	-0.4965	0.0367	0.4979	_	(-)	_
	0.65	-0.4705	0.1122	0.4837	_	(-)	_	-0.4961	0.0390	0.4976	_	(-)	_
	0.70	-0.4679	0.1165	0.4822	_	(-)	_	-0.4931	0.0545	0.4961	_	(-)	_
SG-L	0.55	-0.9171	0.2572	0.9525	0.4554	(0.1321)	_	-0.9809	0.1295	0.9894	0.4897	(0.0660)	_
	0.60	-0.8958	0.2857	0.9403	0.4437	(0.1469)	_	-0.9652	0.1737	0.9807	0.4819	(0.0875)	_
	0.65	-0.8789	0.3064	0.9308	0.4347	(0.1556)	_	-0.9605	0.1840	0.9780	0.4789	(0.0939)	_
	0.70	-0.8711	0.3138	0.9259	0.4304	(0.1592)	_	-0.9510	0.2031	0.9724	0.4740	(0.1029)	_
SG-L-BC	0.55	-0.4617	0.1294	0.4794	_	(-)	_	-0.4912	0.0649	0.4955	_	(-)	_
	0.60	-0.4521	0.1439	0.4745	_	(-)	_	-0.4833	0.0878	0.4912	_	(-)	_
	0.65	-0.4443	0.1573	0.4713	_	(-)	_	-0.4816	0.0925	0.4904	_	(-)	_
	0.70	-0.4407	0.1616	0.4694	_	(-)	_	-0.4770	0.1029	0.4880	_	(-)	_
SG-ML	0.55	0.0586	0.5252	0.5285	0.0379	(0.0278)	_	0.2704	0.5556	0.6179	0.0308	(0.0246)	_
	0.60	-0.0557	0.4497	0.4532	0.0423	(0.0279)	_	0.0642	0.4729	0.4772	0.0398	(0.0253)	_
	0.65	-0.1146	0.3854	0.4021	0.0450	(0.0275)	_	-0.0434	0.3983	0.4007	0.0454	(0.0259)	_
	0.70	-0.1498	0.3557	0.3859	0.0468	(0.0275)	_	-0.0943	0.3433	0.3560	0.0489	(0.0260)	_
SG-ML-BC	0.55	0.0965	0.5064	0.5155	_	(-)	_	0.3012	0.5360	0.6149	_	(-)	_
	0.60	-0.0133	0.4332	0.4334	_	(-)	_	0.1040	0.4549	0.4667	_	(-)	_
	0.65	-0.0696	0.3706	0.3771	_	(-)	_	0.0020	0.3822	0.3822	_	(-)	_
	0.70	-0.1030	0.3408	0.3560	_	(–)	-	-0.0455	0.3280	0.3311	-	(–)	_

(continued).

Tabl	e 2.	Continued.	
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				n =	250				<i>n</i> =	500			
Estimator	α	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }
					Mod	lel 2-A: Splicing	with Weibull	& GPD					
CC	_	0.3261	0.6679	0.7432	0.0104	(0.0228)	_	0.1999	0.6443	0.6746	0.0060	(0.0023)	_
KS	_	-1.3273	0.9856	1.6533	_	(-)	9.1%	-0.6556	1.8320	1.9458	_	(-)	16.2%
Q-MAD	_	-0.2051	0.7303	0.7585	_	(-)	97.5%	0.1488	1.4884	1.4958	_	(-)	90.5%
Q-SUP	_	0.3085	1.6298	1.6588	_	(-)	86.4%	1.0775	2.9302	3.1221	_	(-)	78.4%
AEB	_	4.3704	2.9114	5.2514	_	(–)	8.1%	4.2648	2.0070	4.7135	_	(–)	2.9%
ADST	_	1.0151	1.9948	2.2382	_	(-)	49.1%	2.1329	1.5330	2.6266	_	(-)	23.3%
SG-LS	0.55	-0.6238	0.1250	0.6362	0.1503	(0.0644)	_	-0.6202	0.0914	0.6269	0.1528	(0.0450)	_
	0.60	-0.6268	0.1260	0.6393	0.1550	(0.0675)	_	-0.6237	0.0914	0.6304	0.1578	(0.0473)	_
	0.65	-0.6290	0.1272	0.6417	0.1583	(0.0705)	_	-0.6265	0.0919	0.6333	0.1613	(0.0495)	_
	0.70	-0.6308	0.1282	0.6437	0.1608	(0.0728)	_	-0.6285	0.0935	0.6354	0.1635	(0.0514)	_
SG-LS-BC	0.55	-0.4736	0.0848	0.4811	_	(-)	_	-0.4674	0.0615	0.4714	_	(-)	_
	0.60	-0.4717	0.0841	0.4792	_	(-)	_	-0.4659	0.0601	0.4698	_	(-)	_
	0.65	-0.4708	0.0840	0.4782	_	(-)	_	-0.4652	0.0592	0.4690	_	(-)	_
	0.70	-0.4700	0.0839	0.4775	_	(-)	_	-0.4649	0.0592	0.4687	_	(-)	_
SG-L	0.55	-0.6196	0.1186	0.6309	0.1848	(0.0668)	_	-0.6224	0.0831	0.6279	0.1862	(0.0470)	_
	0.60	-0.6216	0.1233	0.6337	0.1767	(0.0660)	_	-0.6242	0.0871	0.6303	0.1779	(0.0466)	_
	0.65	-0.6231	0.1282	0.6362	0.1718	(0.0648)	_	-0.6259	0.0898	0.6323	0.1728	(0.0459)	_
	0.70	-0.6239	0.1346	0.6383	0.1692	(0.0638)	_	-0.6275	0.0915	0.6342	0.1701	(0.0452)	_
SG-L-BC	0.55	-0.4348	0.0715	0.4407	_	(-)	_	-0.4362	0.0491	0.4389	_	(-)	_
	0.60	-0.4449	0.0759	0.4513	_	(-)	_	-0.4464	0.0522	0.4494	_	(-)	_
	0.65	-0.4514	0.0811	0.4586	_	(-)	_	-0.4531	0.0547	0.4564	_	(-)	_
	0.70	-0.4547	0.0881	0.4632	_	(-)	_	-0.4574	0.0566	0.4609	_	(-)	_
SG-ML	0.55	-0.4484	0.3354	0.5600	0.0333	(0.0082)	_	-0.4259	0.3665	0.5619	0.0331	(0.0079)	_
	0.60	-0.4518	0.3036	0.5443	0.0332	(0.0074)	_	-0.4568	0.2998	0.5464	0.0332	(0.0064)	_
	0.65	-0.4506	0.2968	0.5395	0.0335	(0.0073)	_	-0.4697	0.2632	0.5384	0.0337	(0.0057)	_
	0.70	-0.4334	0.3124	0.5342	0.0340	(0.0075)	_	-0.4733	0.2508	0.5356	0.0343	(0.0057)	_
SG-ML-BC	0.55	-0.4152	0.3313	0.5311	_	(-)	_	-0.3928	0.3612	0.5336	_	(-)	_
	0.60	-0.4186	0.3005	0.5153	_	(-)	_	-0.4237	0.2964	0.5171	_	(-)	_
	0.65	-0.4171	0.2940	0.5102	_	(-)	_	-0.4360	0.2609	0.5081	_	(-)	_
	0.70	-0.3994	0.3092	0.5051	-	(-)	-	-0.4390	0.2487	0.5046	-	(-)	-
												(con	tinued).

Table 2. Continued.	
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				<i>n</i> =	= 250			n = 500					
Estimator	α	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }
	Model 2-B: Splicing with Weibull & Translated Weibull												
CC	_	0.3302	0.6788	0.7548	0.0116	(0.0351)	_	0.1609	0.6716	0.6906	0.0058	(0.0020)	_
KS	_	-1.4714	0.9325	1.7420	_	(-)	12.4%	-1.0546	3.0694	3.2455	_	(-)	17.3%
Q-MAD	_	-0.2656	1.3378	1.3639	_	(-)	97.2%	0.4715	4.2932	4.3190	_	(-)	93.6%
Q-SUP	_	0.1266	1.8973	1.9015	_	(-)	92.1%	2.7196	16.7906	17.0094	_	(-)	80.1%
AEB	_	6.4959	13.3971	14.8889	_	(-)	39.8%	3.7522	5.4395	6.6081	_	(-)	37.6%
ADST	_	-0.6069	0.7234	0.9442	_	(-)	85.3%	-0.6242	0.4701	0.7814	_	(-)	95.2%
SG-LS	0.55	-0.6123	0.1278	0.6255	0.1384	(0.0591)	_	-0.5991	0.0968	0.6068	0.1364	(0.0427)	_
	0.60	-0.6154	0.1283	0.6287	0.1425	(0.0619)	_	-0.6024	0.0974	0.6103	0.1405	(0.0448)	_
	0.65	-0.6175	0.1296	0.6309	0.1453	(0.0645)	_	-0.6050	0.0982	0.6130	0.1434	(0.0467)	_
	0.70	-0.6193	0.1307	0.6329	0.1473	(0.0665)	_	-0.6069	0.0993	0.6150	0.1453	(0.0484)	_
SG-LS-BC	0.55	-0.4739	0.0899	0.4824	_	(-)	_	-0.4626	0.0670	0.4675	_	(-)	_
	0.60	-0.4730	0.0888	0.4812	_	(-)	_	-0.4620	0.0658	0.4667	_	(-)	_
	0.65	-0.4721	0.0883	0.4803	_	(-)	_	-0.4616	0.0652	0.4662	_	(-)	-
	0.70	-0.4720	0.0881	0.4801	_	(-)	_	-0.4616	0.0649	0.4662	_	(-)	-
SG-L	0.55	-0.6036	0.1328	0.6181	0.1502	(0.0621)	_	-0.5977	0.0968	0.6055	0.1477	(0.0443)	-
	0.60	-0.6028	0.1397	0.6188	0.1428	(0.0614)	_	-0.5969	0.1013	0.6054	0.1404	(0.0438)	-
	0.65	-0.6022	0.1461	0.6196	0.1386	(0.0605)	_	-0.5969	0.1041	0.6059	0.1364	(0.0430)	-
	0.70	-0.6023	0.1518	0.6211	0.1368	(0.0596)	_	-0.5976	0.1056	0.6068	0.1346	(0.0423)	-
SG-L-BC	0.55	-0.4534	0.0857	0.4615	_	(-)	_	-0.4500	0.0611	0.4541	_	(-)	-
	0.60	-0.4600	0.0925	0.4692	_	(-)	_	-0.4565	0.0656	0.4612	_	(-)	-
	0.65	-0.4635	0.0993	0.4741	_	(-)	_	-0.4605	0.0687	0.4656	_	(-)	-
	0.70	-0.4654	0.1061	0.4774	_	(-)	_	-0.4630	0.0706	0.4683	_	(-)	-
SG-ML	0.55	-0.4056	0.3411	0.5300	0.0325	(0.0087)	_	-0.3769	0.3688	0.5273	0.0350	(0.0101)	-
	0.60	-0.4099	0.3227	0.5217	0.0326	(0.0082)	_	-0.4048	0.3125	0.5114	0.0355	(0.0090)	-
	0.65	-0.3956	0.3177	0.5074	0.0329	(0.0078)	_	-0.4224	0.2595	0.4958	0.0359	(0.0080)	-
	0.70	-0.3742	0.3280	0.4976	0.0330	(0.0077)	_	-0.4118	0.2559	0.4848	0.0359	(0.0078)	-
SG-ML-BC	0.55	-0.3732	0.3370	0.5028	_	(-)	_	-0.3418	0.3625	0.4982	_	(-)	-
	0.60	-0.3774	0.3194	0.4944	_	(-)	_	-0.3693	0.3079	0.4808	_	(-)	-
	0.65	-0.3628	0.3147	0.4802	_	(-)	_	-0.3865	0.2567	0.4640	_	(-)	-
	0.70	-0.3412	0.3249	0.4711	_	(-)	_	-0.3758	0.2532	0.4532	_	(-)	_

(continued).

Table 2. Continued.

		n = 250						n = 500					
Estimator	α	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }	Bias	SD	RMSE	ĥ	or \hat{b}	%{ <i>I</i> ₀ }
Model 2-C: Splicing with Weibull & Half-Normal													
CC	_	0.3235	0.6634	0.7381	0.0104	(0.0228)	_	0.1936	0.6369	0.6657	0.0060	(0.0023)	_
KS	_	-1.3201	0.3891	1.3762	_	(-)	6.4%	-1.2673	0.7162	1.4557	_	(-)	2.2%
Q-MAD	_	-0.3307	0.3061	0.4506	_	(_)	99.3%	-0.2735	0.3994	0.4841	_	(_)	96.5%
Q-SUP	_	0.0444	0.9760	0.9770	_	(_)	87.8%	0.1620	1.3087	1.3187	_	(-)	84.8%
AEB	_	1.6486	1.0892	1.9760	_	(_)	30.2%	1.7211	0.7767	1.8882	_	(–)	19.0%
ADST	_	-0.0373	1.3417	1.3422	_	(-)	51.0%	1.1259	1.1772	1.6289	_	(-)	39.7%
SG-LS	0.55	-0.6274	0.1236	0.6394	0.1549	(0.0661)	_	-0.6243	0.0900	0.6308	0.1576	(0.0463)	_
	0.60	-0.6304	0.1244	0.6425	0.1598	(0.0692)	_	-0.6279	0.0901	0.6343	0.1628	(0.0486)	_
	0.65	-0.6328	0.1256	0.6451	0.1634	(0.0721)	_	-0.6308	0.0905	0.6373	0.1666	(0.0506)	_
	0.70	-0.6344	0.1271	0.6470	0.1659	(0.0748)	_	-0.6330	0.0913	0.6395	0.1691	(0.0527)	_
SG-LS-BC	0.55	-0.4725	0.0825	0.4796	_	(-)	_	-0.4667	0.0596	0.4705	_	(-)	_
	0.60	-0.4706	0.0818	0.4776	_	(-)	_	-0.4651	0.0582	0.4687	_	(-)	_
	0.65	-0.4693	0.0818	0.4764	_	(-)	_	-0.4642	0.0573	0.4677	_	(-)	_
	0.70	-0.4685	0.0819	0.4756	_	(-)	_	-0.4639	0.0568	0.4674	_	(-)	_
SG-L	0.55	-0.6234	0.1163	0.6342	0.1948	(0.0690)	_	-0.6266	0.0805	0.6318	0.1970	(0.0487)	_
	0.60	-0.6257	0.1207	0.6373	0.1861	(0.0684)	_	-0.6288	0.0845	0.6344	0.1881	(0.0485)	_
	0.65	-0.6275	0.1255	0.6399	0.1808	(0.0674)	_	-0.6307	0.0872	0.6367	0.1827	(0.0479)	_
	0.70	-0.6284	0.1321	0.6422	0.1779	(0.0663)	_	-0.6325	0.0890	0.6388	0.1795	(0.0471)	_
SG-L-BC	0.55	-0.4287	0.0683	0.4341	_	(-)	_	-0.4296	0.0463	0.4321	_	(-)	_
	0.60	-0.4396	0.0720	0.4455	_	(-)	_	-0.4406	0.0492	0.4434	_	(-)	_
	0.65	-0.4467	0.0771	0.4533	_	(-)	_	-0.4480	0.0515	0.4510	_	(-)	_
	0.70	-0.4505	0.0843	0.4584	_	(-)	_	-0.4530	0.0533	0.4562	_	(-)	_
SG-ML	0.55	-0.4538	0.3291	0.5606	0.0337	(0.0086)	_	-0.4331	0.3675	0.5680	0.0340	(0.0085)	—
	0.60	-0.4587	0.3067	0.5517	0.0336	(0.0078)	_	-0.4606	0.2958	0.5473	0.0338	(0.0068)	—
	0.65	-0.4638	0.2877	0.5458	0.0339	(0.0075)	_	-0.4755	0.2557	0.5399	0.0344	(0.0062)	—
	0.70	-0.4526	0.2988	0.5423	0.0345	(0.0078)	_	-0.4852	0.2422	0.5423	0.0352	(0.0063)	—
SG-ML-BC	0.55	-0.4202	0.3252	0.5313	_	(-)	_	-0.3991	0.3618	0.5387	_	(-)	_
	0.60	-0.4251	0.3035	0.5223	_	(-)	_	-0.4267	0.2922	0.5172	_	(-)	_
	0.65	-0.4299	0.2852	0.5159	_	(-)	_	-0.4411	0.2532	0.5086	_	(-)	_
	0.70	-0.4181	0.2959	0.5122	—	(-)	—	-0.4500	0.2400	0.5100	_	(-)	_

Note: 'Bias', 'SD' and 'RMSE' are biases, standard deviations and RMSEs of splicing point or threshold estimates over 1000 Monte Carlo samples, respectively. For kernel-smoothed threshold location estimation procedures, Monte Carlo averages and standard deviations (in parentheses) of CV tuning parameters are presented under the heading ' \hat{h} or \hat{b} '. For automated threshold detection methods, percentages of threshold estimates falling into I_0 are reported under the heading ' \Re_{I_0} '.

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 Δ (or the exponent α , to be more precise) matters in finite samples. For each model and sample size, smoothing parameter values via MLCV do not vary much across four values of α . On the other hand, $\alpha = 0.70$ (i.e. the smallest shift) always results in the smallest RMSE among four initial SG-ML estimates. Because bias correction is made by simply adding \hat{b} to the initial estimate \hat{t} , additional variability through bias correction (i.e. estimation error of the correction term) is not introduced. As a consequence, superiority of the smallest shift is maintained after the bias correction. Indeed SG-ML-BC with $\alpha = 0.70$ outperforms others in terms of RMSE for Models 1-A and 1-B. Its RMSE is not the smallest for Models 2-A and 2-B; an exception is Model 2-B with n = 500, to be more precise. For these cases, however, the RMSE is simply larger than those from other bias corrected SG methods, and it is still smaller than those from CC and five automated methods.

4.3.2. Model 2-C

It is of particular interest to see how threshold estimators behave in case of a thin tail. Q-MAD continues to perform best among five automated threshold detection methods for each sample size. It produces the smallest RMSE, and again more than 90% of estimates fall within I_0 . The performance of CC is also qualitatively similar to above. The RMSE of SG-ML with $\alpha = 0.70$ becomes worse than that of Q-MAD, and as a result, Q-MAD outperforms SG-ML-BC in terms of RMSE even after the bias correction. It is worth emphasising that the RMSE of SG-ML with $\alpha = 0.70$ is smaller than those of SG-LS and SG-L, and bias correction using a sizable \hat{b} merely improves their RMSEs.

However, it is dubious whether this can be thought of as 'satisfactory' for the automated threshold detection method. Q-MAD is designed to detect the point at which deviation from a GPD occurs, and the exponentially decaying tail contradicts the premise of a thick tail. In this standpoint, results from other automated methods could be more reasonable. Another lesson learnt here is that based on no particular parametric model in the tail part, SG methods identify a jump point of the distribution as the threshold, regardless of its tail thickness.

4.3.3. Summary

Invoke that the local structure (2) holds in the neighbourhood of the true threshold location for two cases of Model 1, whereas it is violated for three cases of Model 2. It can be confirmed from Monte Carlo results that SG-ML-BC with $\alpha = 0.70$ is of most practical relevance and importance, because its RMSE is by far the smallest in favourable scenarios and in general less than those from other competing methods even in unfavourable scenarios with deviation from (2).

5. Real data examples

5.1. Data description

In this section our threshold estimation procedure is applied to two datasets on cost variables that are made publicly available. The one is of actuarial losses and the other of wages. Below each dataset is discussed in detail.

The first dataset is of Danish fire insurance losses. Since the seminal analysis by McNeal (1997), the dataset has been arguably most popularly chosen for empirical studies

on non-life insurance. We extract the one named danish in R-package SMPracticals. The dataset contains 2,492 fire insurance losses denominated in millions of Danish kroner from 1980 to 1990.

The second dataset is taken from Merged Outgoing Rotation Group Earnings Data of the US Current Population Survey (CPS), also known as CPS Labour Extract, which is available on the webpage of the National Bureau of Economic Research. We extract hourly wages (the variable earnhre) earned by males from the dataset in 1979. Before proceeding, all observations denominated originally in US cents are converted into US dollars. The original sample size is 54,769. Considering the computation burden of kernel smoothing, we downsize the dataset to a sub-sample of sample size 2, 709 by random sampling, which roughly accounts for 5% of the original.

Table 3 reports summary statistics of the datasets. Each distribution is right-skewed and thus reasonably represents stylised facts of a 'cost' distribution. It can be also found that the distribution of the downsized dataset of US male wages well represents that of the original.

5.2. Estimation details

Based on simulation results in Section 4, we compare SG-ML(-BC) with $\alpha = 0.70$ with KS, Q-MAD, Q-SUP, AEB, ADST, and CC. The tuning parameter for each of SG-ML(-BC) and CC is taken from 100 equally-spaced grids over the interval [0.005, 0.500], and then the threshold estimate \hat{t} is searched via a numerical optimisation routine of the corresponding diagnostic function.

The prespecified interval I_0 for each dataset roughly covers the upper part of the distribution. In addition, Cooray and Ananda (2005) and Scollnik and Sun (2012) estimate several different parametric composite models from the Danish fire insurance dataset and obtain threshold estimates ranging roughly from 1 through 3. Using the same dataset, Reynkens et al. (2017) report the threshold estimate of 17 via a graphical method. The interval I_0 for this dataset also incorporates these empirical findings.

In addition, to implement ADST, we adopt the 5% level of significance and the ForwardStop procedure for *p*-value adjustments in multiple tests. Candidates of thresholds are 54 empirical percentiles from 20.0% until 99.5% with an increment of 1.5%, i.e. {20.0%, 21.5%, ..., 98.0%, 99.5%}, for Danish fire insurance losses and 34 empirical percentiles from 50.0% until 99.5% with an increment of 1.5%, i.e. {50.0%, 51.5%, ..., 98.0%, 99.5%}, for US male hourly wages. The range of percentiles for each dataset roughly coincides with the corresponding interval I_0 .

5.3. Results

Table 4 presents the estimation results. For the Danish fire insurance data, there is no consensus in threshold estimates among five GPD-based detection methods, except that those from KS and ADST are fairly close each other. The estimate from SG-ML is close to those from KS and ADST, so is the one from its bias-corrected version SG-ML-BC. In contrast, the result from CC is problematic and recognised as an example of estimation failure in that the threshold estimate is a corner solution.

Table 3. Descriptive statistics of datasets.

Data	п	Mean	SD	SK	Min.	Q1	Q2	Q3	90%	95%	99%	Max.
				(A) Danisl	h Fire Insurance	Losses (in mil	lions of Danish	n kroner)				
Original	2,492	3.063	7.975	19.884	0.313	1.157	1.634	2.646	5.080	8.406	24.614	263.250
					(B) US Male H	ourly Wages (i	n US dollars)					
Original	54,769	6.136	2.972	3.013	0.500	3.850	5.630	8.000	9.980	11.000	14.793	99.990
Downsized	2, 709	6.080	2.858	1.525	1.000	3.750	5.550	8.000	9.954	11.000	14.640	40.000

Note: n = sample size; Mean = average; SD = standard deviation; SK = skewness; Min. = minimum value; Q1 = first quartile; Q2 = median (i.e. second quartile); Q3 = third quartile; 90% = 90% quantile; 95% = 95% quantile; 99% = 99% quantile; and Max. = maximum value.

Data	Estimator	Estimate of t ₀	Estimator	α	I ₀	Estimate of t ₀	\hat{h} or \hat{b}
		(A) Danish Fire Insur	ance Losses (in n	nillions of D	anish kroner)	
Original	KS	1.375	CC	_	[1, 30]	30.000	0.005
5	Q-MAD	29.037					
	Q-SUP	11.123	SG-ML	0.70	[1, 30]	1.861	0.235
	AEB	25.288	SG-ML-BC	_	_	2.096	_
	ADST	1.406					
		(B) US Ma	le Hourly Wages	(in US dolla	ars)		
Downsized	KS	10.400	ć	_	[5, 15]	14.998	0.005
	Q-MAD	10.000					
	Q-SUP	10.000	SG-ML	0.70	[5, 15]	9.524	0.140
	AEB	13.500	SG-ML-BC	_	_	9.664	_
	ADST	16.000					

Tab	le 4	 Resu 	ts for	real	data	examp	les.
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On the other hand, all five GPD-based methods detect the threshold in the US male wage distribution at 10 or larger, and in particular, KS, Q-MAD and Q-SUP produce very similar threshold estimates. SG-ML and SG-ML-BC also yield threshold estimates near 10, which is considerably close to the results from Q-MAD and Q-SUP. Once again, CC may have failed to estimate the threshold, because its estimate lies almost on the boundary. Moreover, the threshold estimate by ADST corresponds to the 99.5% empirical percentile; in other words, this method suggests deviations from a GPD at all percentiles considered.

As the location of the threshold is unknown in each distribution, it is hard to judge among all threshold estimators considered. Nonetheless, it is safe to say that SG-ML(-BC) can serve as a good alternative to existing threshold detection methods.

6. Conclusion

It is widely recognised that a single model cannot describe the while range of a cost distribution well. Accordingly, it is of growing importance and interest to find the location of a threshold at which two different models are spliced, whereas this estimation problem is known to be notoriously difficult. This paper has explored a method of estimating the threshold of a cost distribution nonparametrically. Development of our method stems from tailoring the existing techniques for change point detection in statistics to stylised facts of cost distributions. The diagnostic function is the absolute difference of two kernel density estimates, and the maximiser of the function over a prespecified interval is defined as the threshold estimator. Because it is located in the right tail region, we propose to compute two density estimates using the entire sample and shifted gamma kernels. Our proposed estimator is shown to be super-consistent and asymptotically normal when suitably implemented. The proof strategy is also new in that approximations to the incomplete gamma, digamma and polygamma functions are utilised. Since the estimator tends to underestimate the threshold location and its dominant bias term takes a simple form, we advocate correcting its bias in a straightforward manner. It is confirmed in the Monte Carlo study that as a result of the bias correction, the negative bias is substantially eliminated with no additional price of spread. Several data-based methods of choosing the smoothing parameter are also assessed via simulations. Our proposal is finally applied to two kinds of real world datasets. Judging from simulation results and real data examples, we recommend SG-ML-BC for practical use.

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Notes

- 1. Many authors (e.g., Cooray and Ananda 2005; Scollnik and Sun 2012) additionally impose differentiability of the pdf at the splicing point to make the entire density smooth and reduce the number of parameters. This practice has computational advantage now that the splicing point can be expressed as a function of other model parameters. However, such a restriction on parameters results in less flexibility. From this viewpoint, Reynkens et al. (2017) first estimate the splicing point and then compute estimates of remaining model parameters.
- 2. An alternative approach is proposed by Desmet et al. (2010), who transform kernel density estimation into kernel regression estimation via prebinning and applying an existing method of discontinuity detection for nonparametric regression curves. This approach has the disadvantage that prebinning substantially reduces the sample size which can be used for regression estimation.
- 3. The reason why different algorithms are utilised for (i) and (ii) is as follows. While a numerical optimisation routine substantially reduces computation time for (i), it often finds local extrema and corner solutions because of a high degree of nonlinearity in CV criteria. A grid search can circumvent these issues. In contrast, $|\hat{J}(x)|$ is concave on I_0 , and a numerical optimisation routine helps expedite computation for (ii).

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Data availability statement

The datasets used in Section 5 are openly available; see Section 5.1 for more details.

Disclosure statement

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Appendix

The Appendix provides technical proofs of theorems and propositions. To save space, we defer proofs of lemmata to the Supplemental Material. Before proceeding, additional shorthand notation is introduced, and a few useful formulae related to the gamma function are presented.

A.1 Additional notation

 $\Psi(x) = d \ln \Gamma(x)/dx = \Gamma^{(1)}(x)/\Gamma(x) \text{ and } \Psi^{(m)}(x) = d^m \Psi(x)/dx^m \text{ signify the digamma and} polygamma functions, respectively. In addition, the following notation is adopted in the proofs:$ $<math display="block">R(a,z) = z^a \exp(-z)/\Gamma(a+1) \text{ for } a, z > 0; \dot{K}_c^{\pm}(u) = \partial K_x^{\pm}(u)/\partial x|_{x=c}; \ddot{K}_c^{\pm}(u) = \partial^2 K_x^{\pm}(u)/\partial x^2|_{x=c}; \\ \ddot{K}_c^{\pm}(u) = \partial^3 K_x^{\pm}(u)/\partial x^3|_{x=c}; H_i = \dot{K}_{t_0}^{-}(X_i) - \dot{K}_{t_0}^{+}(X_i); \text{ and } z^{\pm} = (t_0 \pm \Delta)/b.$

A.2 Useful formulae on the gamma function

Stirling's formula:

$$\Gamma(a+1) = \sqrt{2\pi} a^{a+1/2} \exp(-a) \left\{ 1 + \frac{1}{12a} + O(a^{-2}) \right\} \text{ as } a \to \infty.$$
(A1)

Recursive formula for the lower incomplete gamma function:

$$\gamma (a+1,z) = a\gamma (a,z) - z^a \exp(-z) \quad \text{for } a, z > 0.$$
 (A2)

Recursive formula for the polygamma function:

$$\Psi^{(m)}(a+1) = \Psi^{(m)}(a) + \frac{(-1)^m m!}{a^{m+1}} \quad \text{for } a > 0 \text{ and } m \in \{0, 1, 2, \ldots\}.$$
 (A3)

A.3 Proof of Proposition 3.1

The proof requires the following lemmata.

Lemma A.1: As $b \rightarrow 0$,

$$\sup_{x \in I_0} \left| \Psi\left(\frac{x}{b} + 1\right) - \left\{ \ln\left(\frac{x}{b}\right) + \frac{b}{2x} \right\} \right| = O\left(b^2\right),$$
$$\sup_{x \in I_0} \left| \Psi^{(1)}\left(\frac{x}{b} + 1\right) - \left(\frac{b}{x} - \frac{b^2}{2x^2}\right) \right| = O\left(b^3\right), \text{ and}$$
$$\sup_{x \in I_0} \left| \Psi^{(m)}\left(\frac{x}{b} + 1\right) \right| = O\left(b^m\right) \text{ for } m \ge 2$$

Lemma A.2: Let $C_0 := 2 \max\{\underline{t}^{-3/2}, \overline{t}^{1/2}\} + \underline{t}^{-1/2}$. Then, as $n \to \infty$,

$$\sup_{(x,u)\in I_0\times\mathbb{R}_+}\left|\dot{K}_x^{\pm}(u)\right|\leq b^{-3/2}\sqrt{\frac{2}{\pi}}C_0.$$

Lemma A.3 (Van der Vaart and Wellner 1996, Lemma 2.2.9): Let X_1, \ldots, X_n be independent random variables with bounded ranges [-M, M] and zero means. Then,

$$\Pr\left(\left|\sum_{i=1}^{n} X_{i}\right| > x\right) \le 2 \exp\left\{-\frac{x^{2}}{2\left(\nu + Mx/3\right)}\right\}$$

for all x and $v \geq Var(\sum_{i=1}^{n} X_i)$.

A.3.1 Proof of Proposition 3.1

Proof of (5): Because

$$E\left\{\hat{f}^{\pm}(x)\right\} = \int_{0}^{\infty} K_{x}^{\pm}(u) g(u) \, \mathrm{d}u + d_{0} \int_{0}^{t_{0}} K_{x}^{\pm}(u) \, \mathrm{d}u,$$

it suffices to show that

$$\sup_{x \in I_0} \left| \int_0^\infty K_x^{\pm}(u) g(u) \, \mathrm{d}u - \left\{ g(x) \pm g^{(1)}(x) \, \Delta \right\} \right| = O(b) \,. \tag{A4}$$

Observe that $\int_0^\infty K_x^{\pm}(u)g(u) \, du = E\{g(X^{\pm})\}$ for $X^{\pm} \stackrel{d}{=} G(a^{\pm} + 1, b)$. A second-order Taylor expansion of $g(X^{\pm})$ around $X^{\pm} = x$ yields

$$E\left\{g\left(X^{\pm}\right)\right\} = g\left(x\right) + g^{(1)}\left(x\right)E\left(X^{\pm} - x\right) + \frac{1}{2}E\left\{g^{(2)}\left(\bar{x}^{\pm}\right)\left(X^{\pm} - x\right)^{2}\right\}$$

for some \bar{x}^{\pm} on the line segment joining X^{\pm} and x. Notice that $|E\{g^{(2)}(\bar{x}^{\pm})(X^{\pm}-x)^2\}| \leq \sup_{x \in \mathbb{R}_+} |g^{(2)}(x)| |E(X^{\pm}-x)^2$, where $\sup_{x \in \mathbb{R}_+} |g^{(2)}(x)| < \infty$ by Assumption 3.2(ii). Furthermore, by the property of gamma random variables and Assumption 3.3, $E(X^{\pm}-x) = \pm \Delta + b$ and $E(X^{\pm}-x)^2 = xb + \Delta^2 \pm 3\Delta b + 2b^2 = \{x + o(1)\}b$. Then, (A4) immediately follows.

Proof of (6): This part can be established by a minor modification of the proof of Theorem 2 in Funke and Hirukawa (2025). However, we present the proof for this part in full in order for this paper to be self-contained. For ease of exposition, let $a_n := \sqrt{\ln n/(nb^{1/2})}$, $\zeta_{in}^{\pm}(x) := (1/n)[K_x^{\pm}(X_i) - E\{K_x^{\pm}(X_i)\}]$, $N_n := n^{1+\epsilon} b^{-3/2}$ for a sufficiently small $\epsilon > 0$, and $|I_0| := \overline{t} - \underline{t}$. Then, the proof takes the following two steps.

- (1) Split the interval I_0 into N_n equally-spaced grids to create N_n sub-intervals with length $N_n^{-1}|I_0|$, and replace the supremum with a maximisation over finite N_n sub-intervals.
- (2) Employ Lemma A.3 (Bernstein's inequality) to bound the remainder term.

Step 1

Let I_j , $j = 1, ..., N_n$, be the *j*th sub-interval. Also let x_j be the right-most point in I_j with $x_0 \equiv \underline{t}$ and $x_{N_n} \equiv \overline{t}$. Suppose that the design point *x* falls into I_j . By the mean-value theorem, Lemma A.2 and Assumption 3.3,

$$\begin{aligned} \left| K_x^{\pm} \left(u \right) - K_{x_j}^{\pm} \left(u \right) \right| &\leq \sup_{\left(x, u \right) \in I_0 \times \mathbb{R}_+} \left| \dot{K}_x^{\pm} \left(u \right) \right| \sup_{x \in I_j} \left| x - x_j \right| \\ &\leq O\left(b^{-3/2} \right) O\left(N_n^{-1} \right) = O\left\{ n^{-(1+\epsilon)} \right\} \leq O\left(a_n \right). \end{aligned}$$

It follows from C_r -inequality that

$$\max_{1 \le j \le N_n} \sup_{x \in I_j} \left| \sum_{i=1}^n \zeta_{in}^{\pm}(x) - \sum_{i=1}^n \zeta_{in}^{\pm}(x_j) \right| = O(a_n).$$
(A5)

Step 2

Before employing Bernstein's inequality in Lemma A.3, we must determine two bounds M and v. First, it holds that for a given $x > \Delta(> 0)$,

$$K_x^{\pm}(u) \le \frac{b^{-1/2} \left(x \pm \Delta\right)^{-1/2} \{1 + o(1)\}}{\sqrt{2\pi}}.$$
 (A6)

This can be confirmed by recognising that $K_x^{\pm}(u)$ is maximised at $u = x \pm \Delta$ and employing (A1). Then, by (A6),

$$\sup_{(x,u)\in I_0\times\mathbb{R}_+}\left|K_x^{\pm}\left(u\right)\right|\leq b^{-1/2}\sqrt{\frac{2}{\pi\,\underline{t}}},$$

and thus, by C_r -inequality,

$$\left|\zeta_{in}^{\pm}(x)\right| \leq 2\sqrt{\frac{2}{\pi t}}n^{-1}b^{-1/2} = 2\sqrt{\frac{2}{\pi t}}\frac{a_n^2}{\ln n} =: M.$$

Second, Assumption 3.2(i) implies that there is some constant $\overline{C} \in (0, \infty)$ so that $\sup_{x \in \mathbb{R}_+} f(x) \leq 1$ \overline{C} . Then,

$$Var\left\{\sum_{i=1}^{n}\zeta_{in}^{\pm}(x)\right\} = \sum_{i=1}^{n} Var\left\{\zeta_{in}^{\pm}(x)\right\}$$
$$= \sum_{i=1}^{n} E\left\{\zeta_{in}^{\pm}(x)\right\}^{2}$$
$$\leq \sum_{i=1}^{n} E\left\{\frac{1}{n}K_{x}^{\pm}(X_{i})\right\}^{2}$$
$$\leq \frac{\overline{C}}{n}\int_{0}^{\infty}\left\{K_{x}^{\pm}(u)\right\}^{2} du.$$

Also let

$$A^{\pm}(x) := \frac{b^{-1}\Gamma\left(2a^{\pm}+1\right)}{2^{2a^{\pm}+1}\Gamma^{2}\left(a^{\pm}+1\right)}.$$
(A7)

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Observe that

$$\int_0^\infty \left\{ K_x^{\pm}(u) \right\}^2 \, \mathrm{d}u = A^{\pm}(x) \int_0^\infty \frac{u^{2a^{\pm}} \exp\left\{ -u/(b/2) \right\}}{(b/2)^{2a^{\pm}+1} \Gamma\left(2a^{\pm}+1\right)} \, \mathrm{d}u = A^{\pm}(x) \,,$$

because the integrand in the middle term is the pdf of $G(2a^{\pm} + 1, b/2)$. Using (A1), $(x \pm \Delta)^{-1/2} = x^{-1/2}\{1 + o(1)\}$ and $o(1) \le 1$ for a sufficiently large *n* gives

$$A^{\pm}(x) = \frac{b^{-1/2} \left(x \pm \Delta\right)^{-1/2} \left\{1 + o\left(1\right)\right\}}{2\sqrt{\pi}} \le \frac{b^{-1/2}}{\sqrt{\pi t}}$$

uniformly on I_0 . In sum,

$$\operatorname{Var}\left\{\sum_{i=1}^{n}\zeta_{in}^{\pm}(x)\right\} \leq \frac{\overline{C}}{\sqrt{\pi t}}n^{-1}b^{-1/2} = \frac{\overline{C}}{\sqrt{\pi t}}\frac{a_{n}^{2}}{\ln n} =: v.$$

Lemma A.3 establishes that for such *M* and *v* and an arbitrarily chosen K > 0,

$$\Pr\left\{\left|\sum_{i=1}^{n}\zeta_{in}^{\pm}(x)\right| > K\sqrt{\frac{\overline{C}}{\sqrt{\pi \underline{t}}}}a_{n}\right\} \le 2\exp\left\{-\frac{K^{2}\ln n}{2\left(1+\frac{2}{3}\sqrt{\frac{2}{\pi \underline{t}}}Ka_{n}/\sqrt{\frac{\overline{C}}{\sqrt{\pi \underline{t}}}}\right)}\right\}$$

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It follows from $a_n = o(1)$ that $(2/3)\sqrt{2/(\pi \underline{t})}Ka_n/\sqrt{\overline{C}/\sqrt{\pi \underline{t}}} \le 1$ holds for a sufficiently large *n*. Accordingly,

$$\Pr\left\{\left|\sum_{i=1}^{n} \zeta_{in}^{\pm}(x)\right| > K\sqrt{\frac{\overline{C}}{\sqrt{\pi \underline{t}}}}a_n\right\} \le 2\exp\left\{-\frac{K^2\ln n}{2(1+1)}\right\} = 2n^{-\frac{K^2}{4}}.$$

In the end,

$$\Pr\left\{\max_{1\leq j\leq N_n}\left|\sum_{i=1}^n \zeta_{in}^{\pm}(x_j)\right| > K\sqrt{\frac{\overline{C}}{\sqrt{\pi \underline{t}}}}a_n\right\}$$

$$\leq \sum_{i=1}^{N_n}\max_{1\leq j\leq N_n}\Pr\left\{\left|\sum_{i=1}^n \zeta_{in}^{\pm}(x_j)\right| > K\sqrt{\frac{\overline{C}}{\sqrt{\pi \underline{t}}}}a_n\right\}$$

$$\leq N_n \cdot \max_{1\leq j\leq N_n}\Pr\left\{\left|\sum_{i=1}^n \zeta_{in}^{\pm}(x_j)\right| > K\sqrt{\frac{\overline{C}}{\sqrt{\pi \underline{t}}}}a_n\right\}$$

$$= O\left(N_n n^{-\frac{K^2}{4}}\right).$$

Moreover, (4) implies that

$$b^{-3/2} = O\left\{n^{\frac{1}{1-\kappa}} \left(\frac{b^{\frac{\kappa}{2}}}{\ln n}\right)^{\frac{1}{1-\kappa}}\right\} \le O\left(n^{\frac{1}{1-\kappa}}\right),$$

where the last inequality holds because $b^{\kappa/2}/\ln n$ is convergent. Then, picking $K = 2\sqrt{2(1+\epsilon) + 1/(1-\kappa)}$ yields $N_n n^{-K^2/4} = O\{n^{-(1+\epsilon)}\}$ so that

$$\sum_{n=1}^{\infty} \Pr\left\{ \max_{1 \le j \le N_n} \left| \sum_{i=1}^n \zeta_{in}^{\pm} \left(x_j \right) \right| > K \sqrt{\frac{\overline{C}}{\sqrt{\pi \underline{t}}}} a_n \right\} \le \sum_{n=1}^{\infty} O\left(\frac{1}{n^{1+\epsilon}} \right) < \infty$$

Therefore, by the Borel-Cantelli lemma,

$$\max_{1 \le j \le N_n} \left| \sum_{i=1}^n \zeta_{in}^{\pm} \left(x_j \right) \right| = O\left(a_n \right) \quad a.s..$$
(A8)

It follows from (A5) and (A8) that

$$\sup_{x \in I_0} \left| \hat{f}^{\pm}(x) - E\left\{ \hat{f}^{\pm}(x) \right\} \right|$$

$$\leq \max_{1 \leq j \leq N_n} \left| \sum_{i=1}^n \zeta_{in}^{\pm}(x_j) \right| + \max_{1 \leq j \leq N_n} \sup_{x \in I_j} \left| \sum_{i=1}^n \zeta_{in}^{\pm}(x) - \sum_{i=1}^n \zeta_{in}^{\pm}(x_j) \right|$$

$$= O(a_n) \quad a.s..$$

This completes the proof.

A.4 Proof of Proposition 3.2

Proof of (i): Dividing both sides of (A2) by $\Gamma(a^{\pm} + 1)$ and using $\Gamma(a^{\pm} + 1) = a^{\pm}\Gamma(a^{\pm})$, we have

$$P(a^{\pm}+1,z_0) = P(a^{\pm},z_0) - R(a^{\pm},z_0)$$

Then, J(x) can be rewritten as

$$J(x) = \{ P(a^{-}, z_{0}) - P(a^{+}, z_{0}) \} - \{ R(a^{-}, z_{0}) - R(a^{+}, z_{0}) \}.$$
(A9)

Let y^{\pm} solve the equation $z_0 = a^{\pm} + \sqrt{a^{\pm}}y^{\pm}$. Then,

$$y^{\pm} = \frac{z_0 - a^{\pm}}{\sqrt{a^{\pm}}} = \frac{t_0 - (x \pm \Delta)}{\sqrt{b (x \pm \Delta)}}.$$

Now Equation (1) of Pagurova (1965) is applied to $P(a^{\pm}, z_0)$, where this equation can be viewed as the Edgeworth expansion for a gamma cdf. In fact,

$$P(a^{\pm}, z_0) = P\left(a^{\pm}, a^{\pm} + \sqrt{a^{\pm}}y^{\pm}\right) = \Phi\left(y^{\pm}\right) - \frac{1}{3\sqrt{a^{\pm}}}\phi^{(2)}\left(y^{\pm}\right) + R_{a^{\pm}},$$
(A10)

where $\Phi(\cdot)$ is the cdf of N(0, 1). Also observe that $\phi^{(j)}(x) =: (-1)^j H_j(x)\phi(x)$, where the $H_j(x)$ are Hermite polynomials (i.e. $H_0(x) = 1$, $H_1(x) = x$, $H_2(x) = x^2 - 1$, $H_3(x) = x^3 - 3x$,...). The remainder term $R_{a^{\pm}}$ in (A10) is $(1/a^{\pm})$ times higher-order derivatives of $\phi(x)$ evaluated at y^{\pm} . It follows from (A9) and (A10) that

$$J(x) = \left\{ \Phi(y^{-}) - \Phi(y^{+}) \right\} - \frac{1}{3} \left\{ \frac{\phi^{(2)}(y^{-})}{\sqrt{a^{-}}} - \frac{\phi^{(2)}(y^{+})}{\sqrt{a^{+}}} \right\} + (R_{a^{-}} - R_{a^{+}}) - \left\{ R(a^{-}, z_{0}) - R(a^{+}, z_{0}) \right\}.$$

Part (i) is established if all the followings are demonstrated:

$$\sup_{x \in I_0} \left| \left\{ \Phi\left(y^-\right) - \Phi\left(y^+\right) \right\} - Q\left(x\right) \left(\frac{\Delta}{b^{1/2}}\right) \right| = O\left(\frac{\Delta^3}{b^{3/2}}\right); \tag{A11}$$

$$\sup_{x \in I_0} \left| \frac{\phi^{(2)}(y^-)}{\sqrt{a^-}} - \frac{\phi^{(2)}(y^+)}{\sqrt{a^+}} \right| = O\left(\frac{\Delta^3}{b^{3/2}}\right);$$
(A12)

$$\sup_{x \in I_0} |R_{a^-} - R_{a^+}| = O\left(\frac{\Delta^3}{b^{3/2}}\right); \quad \text{and}$$
(A13)

$$\sup_{x \in I_0} \left| R\left(a^-, z_0 \right) - R\left(a^+, z_0 \right) \right| = O\left(\frac{\Delta^3}{b^{3/2}} \right).$$
(A14)

Now, a third-order Taylor expansion of $\Phi(y^{\pm})$ around $\Delta = 0$ yields

$$\Phi\left(y^{\pm}\right) = \Phi\left(\frac{t_0 - x}{\sqrt{bx}}\right) \mp \left(\frac{t_0 + x}{2x^{3/2}}\right) \phi\left(\frac{t_0 - x}{\sqrt{bx}}\right) \left(\frac{\Delta}{b^{1/2}}\right) + \frac{1}{2} \left\{ \left(\frac{t_0 + x}{4x^3}\right) \phi^{(1)} \left(\frac{t_0 - x}{\sqrt{bx}}\right) \left(\frac{\Delta^2}{b}\right) + \left(\frac{3t_0 + x}{4x^{5/2}}\right) \phi\left(\frac{t_0 - x}{\sqrt{bx}}\right) \left(\frac{\Delta^2}{b^{1/2}}\right) \right\} + O\left(\frac{\Delta^3}{b^{3/2}}\right),$$
(A15)

where the $O(\Delta^3/b^{3/2})$ rate is uniform on I_0 . Observing that the terms involving odd orders of Δ survive after taking the difference between $\Phi(y^-)$ and $\Phi(y^+)$ and recognising that the $O(\Delta/b^{1/2})$ term can be rewritten as $\mp (1/2)Q(x)(\Delta/b^{1/2})$ establish (A11).

For (A12), notice that

$$\frac{1}{\sqrt{a^{\pm}}} = \frac{b^{1/2}}{\sqrt{x}} \left\{ 1 \mp \frac{\Delta}{2x} + O\left(\Delta^2\right) \right\},\tag{A16}$$

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where the $O(\Delta^2)$ rate is uniform on I_0 . Moreover, a second-order Taylor expansion of $\phi^{(2)}(y^{\pm})$ around $\Delta = 0$ yields

$$\phi^{(2)}\left(y^{\pm}\right) = \phi^{(2)}\left(\frac{t_0 - x}{\sqrt{bx}}\right) \mp \left(\frac{t_0 + x}{2x^{3/2}}\right)\phi^{(3)}\left(\frac{t_0 - x}{\sqrt{bx}}\right)\left(\frac{\Delta}{b^{1/2}}\right) + O\left(\frac{\Delta^2}{b}\right),$$

where the $O(\Delta^2/b)$ rate is again uniform on I_0 . By (3),

$$\sup_{x\in I_0} \left| \frac{\phi^{(2)}\left(y^{-}\right)}{\sqrt{a^{-}}} - \frac{\phi^{(2)}\left(y^{+}\right)}{\sqrt{a^{+}}} \right| \le O\left(b^{1/2}\right) O\left(\frac{\Delta}{b^{1/2}}\right) = O\left(\Delta\right) = o\left(\frac{\Delta^3}{b^{3/2}}\right) \le O\left(\frac{\Delta^3}{b^{3/2}}\right),$$

and thus (A12) is also shown. By a similar argument, jointly with $1/a^{\pm} = O(b)$ uniformly on I_0 ,

$$\sup_{x \in I_0} |R_{a^-} - R_{a^+}| \le O(b) O\left(\frac{\Delta}{b^{1/2}}\right) = O\left(\Delta b^{1/2}\right) = o\left(\frac{\Delta^3}{b^{3/2}}\right) \le O\left(\frac{\Delta^3}{b^{3/2}}\right),$$

which yields (A13).

Finally, using (A1) yields

$$R(a^{\pm}, z_0) = \frac{z_0^{a^{\pm}} \exp(-z_0)}{\Gamma(a^{\pm} + 1)} = \left\{ \frac{1 + O(1/a^{\pm})}{\sqrt{2\pi}} \right\} \frac{1}{\sqrt{a^{\pm}}} \left\{ \left(\frac{z_0}{a^{\pm}}\right)^{a^{\pm}} \exp(a^{\pm} - z_0) \right\},$$
 (A17)

where $\{1 + O(1/a^{\pm})\}/\sqrt{2\pi} = O(1)$ uniformly on I_0 . In addition,

$$\left(\frac{z_0}{a^{\pm}}\right)^{a^{\pm}} \exp\left(a^{\pm} - z_0\right) =: \exp\left\{a^{\pm} \left(\ln \rho^{\pm} + 1 - \rho^{\pm}\right)\right\}$$

for $\rho^{\pm} := z_0/a^{\pm} > 0$. It follows from $\ln \rho^{\pm} + 1 - \rho^{\pm} \le 0$ that $\exp\{a^{\pm}(\ln \rho^{\pm} + 1 - \rho^{\pm})\} \le 1 \le O(1)$. Then, by (A16),

$$\sup_{x \in I_0} \left| R\left(a^-, z_0\right) - R\left(a^+, z_0\right) \right| \le O\left(\Delta b^{1/2}\right) = o\left(\frac{\Delta^3}{b^{3/2}}\right) \le O\left(\frac{\Delta^3}{b^{3/2}}\right)$$

and thus (A14) is also proven.

Proof of (ii): Notice that

$$Q^{(1)}(x) = \frac{q(x)}{2bx^{7/2}}\phi\left(\frac{x-t_0}{\sqrt{bx}}\right),$$
(A18)

where $q(x) := -bx(x + 3t_0) - (x - t_0)(x + t_0)^2$. Heuristically, $q(x) \approx -(x - t_0)(x + t_0)^2$ for $b \approx 0$, and we can see that Q(x) is maximised at $x \approx t_0$. To be more precise, the maximiser of Q(x) is given by the solution of q(x) = 0 or $-bx(x + 3t_0) = (x - t_0)(x + t_0)^2$. It is not hard to see that for a sufficiently small b > 0, the latter has a unique solution on \mathbb{R}_+ .

Let t^* be the solution. Our argument so far suggests that $t^* \approx t_0$ for a sufficiently small b > 0. Now we even conjecture that $t^* = t_0 + cb$ for some $|c| < \infty$. To verify this conjecture, consider that $0 = q(t^*) = -b\{(t_0 + cb)(4t_0 + cb) + c(2t_0 + cb)^2\}$. Because b > 0, c solves $(t_0 + cb)(4t_0 + cb) + c(2t_0 + cb)^2 = 0$. This equation can be further rewritten as $b(c + 1)(c + 2t_0/b)^2 = -t_0c$. Observe that the left- and right-hand sides are cubic and downward sloping linear functions of c, respectively. For a sufficiently small b > 0, $-2t_0/b < -1$ holds, and in this case we can recognise that the equation has a unique solution $c \in (-1, 0)$. This completes the proof.

A.5 Proof of Theorem 3.1

Because $|\hat{t} - t_0| \le |\hat{t} - t^*| + |t^* - t_0|$ and we have already known in Proposition 3.2(ii) that $|t^* - t_0| \le b$, it suffices to demonstrate that

$$\left|\hat{t} - t^*\right| = O\left(c_n\right) \quad a.s.. \tag{A19}$$

The proof for (A19) closely follows the one for Theorem 1 of Chu and Cheng (1996). We keep using the same notation as in the proof of (6) in Proposition 3.1. In addition, define $\Upsilon := \{x : x \in I_0, |x - t^*| > c_n |I_0|\}$. Also let *w* be an element of $E_n := \{x_0, x_1, \dots, x_{N_n}\}$ that is closest to t^* , i.e.

$$|w-t^*| = \min_{0 \le j \le N_n} |x_j - t^*| \Leftrightarrow w = \arg\min_{s \in E_n} |s-t^*|$$

To show (A19), we need to prove that $\Pr\{\hat{t} \in \Upsilon\} = 0$. However, the set $\{\sup_{x \in \Upsilon} |\hat{J}(x)| \ge |\hat{J}(w)|\}$ is larger than the set $\{\hat{t} \in \Upsilon\}$, and thus $\Pr\{\hat{t} \in \Upsilon\} \le \Pr\{\sup_{x \in \Upsilon} |\hat{J}(x)| \ge |\hat{J}(w)|\}$ is the case. In what follows, we establish that

$$\Pr\left\{\sup_{x\in\Upsilon}\left|\hat{f}(x)\right| \ge \left|\hat{f}(w)\right|\right\} = 0.$$
(A20)

Let $x^* := \arg \sup_{x \in \Upsilon} |E\{\hat{J}(x)\}|$. Using (7) and Proposition 3.2 and invoking that $\Delta = o(\Delta^3/b^{3/2})$, we have

$$\left| E\left\{ \hat{J}\left(w\right) \right\} \right| - \sup_{x \in \Upsilon} \left| E\left\{ \hat{J}\left(x\right) \right\} \right| = \left| d_0 \right| \left\{ Q\left(w\right) - Q\left(x^*\right) \right\} \left(\frac{\Delta}{b^{1/2}}\right) + O\left(\frac{\Delta^3}{b^{3/2}}\right).$$
(A21)

For a sufficiently large *n*, *w* is closer to t^* than x^* . Hence, $Q(w) - Q(x^*) > 0$ is the case. By the mean value theorem, $Q(w) - Q(x^*) = Q^{(1)}(\varpi)(w - x^*)$ also holds, where $\varpi := \lambda x^* + (1 - \lambda)w$ for $\lambda (= \lambda_n) \in (0, 1)$. Combining these, we may write $Q(w) - Q(x^*) = |Q^{(1)}(\varpi)| |w - x^*|$.

Below the lower bound of $|Q^{(1)}(\varpi)||w - x^*|$ is obtained. Before proceeding, the following bounds are found to be useful:

$$|w - t^*| \le N_n^{-1} |I_0| = n^{-(1+\epsilon)} b^{3/2} |I_0|;$$
 (A22)

$$|x^* - t^*| \ge c_n |I_0|;$$
 (A23)

$$t_0 - t^* \in (0, b);$$
 and (A24)

$$\left|x^* - t_0\right| \le \eta b^{1/2} \left|I_0\right| \text{ for some constant } \eta > 0.$$
(A25)

While (A22) and (A23) follow from the definitions of w and x^* , respectively, (A24) is the direct outcome of Proposition 3.2(ii). To obtain (A25), suppose that some point $x' \in I_0$ and t_0 are far apart in the sense that $|x' - t_0| = O(b^{\varrho})$ for some $\varrho \in [0, 1/2)$. For such x', $|x' - t_0|/\sqrt{bx'} = O(b^{\varrho-1/2})$ diverges, and thus $Q(x') \to 0$ exponentially fast. Accordingly, $|E\{\hat{J}(x')\}| = O(\Delta^3/b^{3/2}) = o(\Delta/b^{1/2})$; in other words, such x' cannot be x^* . Therefore, a sensible range for x^* must be the one given in (A25).

It follows from (A18) that $|Q^{(1)}(\varpi)| = \{|q(\varpi)|/(2b\varpi^{7/2})\}\phi\{(\varpi - t_0)/\sqrt{b\varpi}\}\)$. The lower bound of this quantity can be found as follows. By (A22), (A24) and (A25),

$$\begin{split} |\varpi - t_0| &\leq \lambda \left| x^* - t_0 \right| + (1 - \lambda) \left| w - t_0 \right| \\ &\leq \lambda \left| x^* - t_0 \right| + (1 - \lambda) \left(\left| w - t^* \right| + \left| t^* - t_0 \right| \right) \\ &\leq \lambda \eta b^{1/2} \left| I_0 \right| + (1 - \lambda) \left\{ n^{-(1+\epsilon)} b^{3/2} \left| I_0 \right| + b \right\} \\ &= b^{1/2} \left| I_0 \right| \left[\lambda \eta + (1 - \lambda) \left\{ n^{-(1+\epsilon)} b + \frac{b^{1/2}}{\left| I_0 \right|} \right\} \right]. \end{split}$$

Because $n^{-(1+\epsilon)}b + b^{1/2}/|I_0|$ is convergent, we may put $n^{-(1+\epsilon)}b + b^{1/2}/|I_0| \le \eta$ for a sufficiently large *n* so that $|\varpi - t_0| \le \eta |I_0| b^{1/2}$. By $\varpi \in I_0 = [\underline{t}, \overline{t}]$, we also have $|\varpi - t_0|/\sqrt{b\varpi} \le \eta |I_0|/\sqrt{\underline{t}}$ so

that $\phi\{(\overline{\omega} - t_0)/\sqrt{b\overline{\omega}}\} = \phi\{|\overline{\omega} - t_0|/\sqrt{b\overline{\omega}}\} \ge \phi(\eta|I_0|/\sqrt{\underline{t}})$. Moreover, $|q(\overline{\omega})| \ge |(\overline{\omega} - t_0)(\overline{\omega} + t_0)^2| - |b\overline{\omega}(\overline{\omega} + 3t_0)| = |\overline{\omega} - t_0|(\overline{\omega} + t_0)^2 - b\overline{\omega}(\overline{\omega} + 3t_0)$.

$$\begin{aligned} |\varpi - t_0| &\geq |\varpi - t^*| - |t_0 - t^*| \\ &\geq \lambda |x^* - t^*| - \{(1 - \lambda) |w - t^*| + |t_0 - t^*|\} \\ &\geq \lambda c_n |I_0| - \{(1 - \lambda) n^{-(1 + \epsilon)} b^{3/2} |I_0| + b\}. \end{aligned}$$

Invoke that $|Q^{(1)}(\varpi)| = \{Q(w) - Q(x^*)\}/|w - x^*|$. For a sufficiently large $n, w \approx t^*$ is the case, and Q(x) in the vicinity of w is nearly flat. If ϖ were located near w, we would have $|Q^{(1)}(\varpi)| \approx |Q^{(1)}(t^*)| = 0$. In reality, $|Q^{(1)}(\varpi)|$ is still positive (although it is close to zero); in short, $\lambda \approx 0$ would contradict positivity of $|Q^{(1)}(\varpi)|$. On the other hand, concavity of Q(x) in the neighbourhood of t^* implies that to maintain positivity of $|Q^{(1)}(\varpi)|$, we must take ϖ near x^* or set λ far away from zero. Therefore, we are allowed to take $\lambda > 1/2$ (and thus $1 - \lambda < 1/2$) for a sufficiently large n so that

$$|\varpi - t_0| \ge \frac{1}{2} c_n |I_0| \left[1 - \frac{b}{c_n} \left\{ n^{-(1+\epsilon)} b^{1/2} |I_0| + \frac{2}{|I_0|} \right\} \right]$$

It follows from $(b/c_n)\{n^{-(1+\epsilon)}b^{1/2}|I_0| + 2/|I_0|\} = o(1)$ that $(b/c_n)\{n^{-(1+\epsilon)}b^{1/2}|I_0| + 2/|I_0|\} \le 1/2$ holds for a sufficiently large *n*, and thus $|\varpi - t_0| \ge (|I_0|/4)c_n$. Using $\varpi \in I_0 = [\underline{t}, \overline{t}]$ yields

$$\left|q\left(\varpi\right)\right| \geq \frac{|I_{0}|}{4}c_{n}\left(\underline{t}+t_{0}\right)^{2}-b\bar{t}\left(\bar{t}+3t_{0}\right)=\frac{c_{n}}{4}\left\{\left|I_{0}\right|\left(\underline{t}+t_{0}\right)^{2}-4\frac{b}{c_{n}}\bar{t}\left(\bar{t}+3t_{0}\right)\right\}$$

Again by $4(b/c_n)\bar{t}(\bar{t}+3t_0) = o(1)$, we may put $4(b/c_n)\bar{t}(\bar{t}+3t_0) \le (1/2)|I_0|(\underline{t}+t_0)^2$ for a sufficiently large *n*. Therefore, $|q(\varpi)| \ge (|I_0|/8)(\underline{t}+t_0)^2c_n$, and thus

$$\left|Q^{(1)}\left(\varpi\right)\right| \geq \frac{\left|I_{0}\right|\left(\underline{t}+t_{0}\right)^{2}}{16\overline{t}^{7/2}}\phi\left(\frac{\eta\left|I_{0}\right|}{\sqrt{\underline{t}}}\right)\left(\frac{c_{n}}{b}\right).$$
(A26)

For the lower bound of $|w - x^*|$, it follows from (A22) and (A23) that

$$|w-x^*| \ge |x^*-t^*| - |w-t^*| \ge c_n |I_0| - N_n^{-1} |I_0| = c_n \left\{ 1 - n^{-(1+\epsilon)} \frac{b^{3/2}}{c_n} \right\} |I_0|.$$

The fact that $n^{-(1+\epsilon)}b^{3/2}/c_n = o(1)$ implies that $n^{-(1+\epsilon)}b^{3/2}/c_n \le 1/2$ for a sufficiently large *n*. Then, we have

$$|w - x^*| \ge \frac{|I_0|}{2}c_n.$$
 (A27)

Substituting (A26) and (A27) into (A21) gives

$$\left|E\left\{\hat{J}(w)\right\}\right| - \sup_{x\in\Upsilon} \left|E\left\{\hat{J}(x)\right\}\right| \ge 3L + O\left(\frac{\Delta^3}{b^{3/2}}\right),$$

where

$$L = L_n := \frac{|d_0| |I_0|^2 \left(\underline{t} + t_0\right)^2}{96\overline{t}^{7/2}} \phi\left(\frac{\eta |I_0|}{\sqrt{\underline{t}}}\right) \left(\frac{c_n^2 \Delta}{b^{3/2}}\right).$$

By this result and a straightforward calculation,

$$\sup_{x\in\Upsilon} \left| \hat{J}(x) \right| - \left| \hat{J}(w) \right| \le 2 \sup_{x\in I_0} \left| \hat{J}(x) - E\left\{ \hat{J}(x) \right\} \right| - 3L + O\left(\frac{\Delta^3}{b^{3/2}}\right).$$

However, by the definition of *L* and (3), each of $\sqrt{\ln n/(nb^{1/2})}$ and $\Delta^3/b^{3/2}$ is of smaller order of magnitude than $c_n^2 \Delta/b^{3/2} = b^{-1/2+2\delta_1} \Delta$, and thus

$$2\left[\sup_{x\in I_0}\left|\hat{J}(x)-E\left\{\hat{J}(x)\right\}\right|-L\right]+\left\{O\left(\frac{\Delta^3}{b^{3/2}}\right)-L\right\}<0$$

with probability one. Therefore, $\Pr\{\sup_{x\in\Upsilon} |\hat{J}(x)| < |\hat{J}(w)|\} = 1$, and (A20) is established. This completes the proof.

A.6 Proof of Theorem 3.2

The proof requires the following lemmata.

Lemma A.4: For $a, z, \lambda > 0$ and $m \in \{0, 1, 2, ...\}$,

$$\lambda^{m} \frac{\gamma (a+m+1,z)}{\Gamma (a+1)} =: p_{m} P(a,z) - r_{m} R(a,z),$$

where $p_{m+1} = \lambda(a + m + 1)p_m$, $r_{m+1} = \lambda(a + m + 1)r_m + (\lambda z)^{m+1}$, and $p_0 = r_0 = 1$.

Lemma A.5: As $n \to \infty$,

$$P(z^{\pm}, z_0) = \frac{1}{2} \mp \frac{1}{\sqrt{2\pi}\sqrt{t_0}} \left(\frac{\Delta}{b^{1/2}}\right) + O\left(\frac{\Delta^2}{b^{1/2}}\right), \text{ and}$$
$$R(z^{\pm}, z_0) = \frac{b^{1/2}}{\sqrt{2\pi}\sqrt{t_0}} \left[1 \mp \frac{\Delta}{2t_0} - \frac{1}{2t_0} \left(\frac{\Delta^2}{b}\right) + O\left\{\max\left(b, \frac{\Delta^3}{b}\right)\right\}\right].$$

Lemma A.6: As $n \to \infty$,

$$E\left\{\left(\frac{b^{1/2}}{\Delta}\right)\hat{J}^{(1)}\left(t_{0}\right)\right\} \rightarrow -\sqrt{\frac{2}{\pi}}\frac{d_{0}}{t_{0}^{3/2}}$$

Lemma A.7: As $n \to \infty$,

$$Var\left\{\sqrt{\frac{nb^{5/2}}{\Delta^2}}\hat{j}^{(1)}(t_0)\right\} \to \frac{3}{2\sqrt{\pi}t_0^{5/2}}\left\{\frac{f(t_0^-) + f(t_0^+)}{2}\right\}.$$

Lemma A.8: If $|\hat{t} - t_0| = o_p(b^{1/2})$, then, as $n \to \infty$,

$$\left(\frac{b^{3/2}}{\Delta}\right)\hat{J}^{(2)}\left(\zeta\right) \xrightarrow{p} -\sqrt{\frac{2}{\pi}}\frac{d_0}{t_0^{3/2}}$$

for any ξ on the line segment joining \hat{t} and t_0 .

Lemma A.9: As $n \to \infty$, $E|H_i|^3 = O(\Delta^3/b^4)$.

A.6.1 Proof of Theorem 3.2

By a mean-value expansion of the first-order condition $\hat{J}^{(1)}(\hat{t}) = 0$, we have

$$0 = \hat{f}^{(1)}(t_0) + \hat{f}^{(2)}\begin{pmatrix} \circ \\ t \end{pmatrix} (\hat{t} - t_0) = E\left\{\hat{f}^{(1)}(t_0)\right\} + \left[\hat{f}^{(1)}(t_0) - E\left\{\hat{f}^{(1)}(t_0)\right\}\right] + \hat{f}^{(2)}\begin{pmatrix} \circ \\ t \end{pmatrix} (\hat{t} - t_0)$$
(A28)

for some $\stackrel{\circ}{t}$ on the line segment joining \hat{t} and t_0 . Theorem 3.1 indicates that the condition for Lemma A.8 is satisfied.

Rearranging (A28), we obtain

$$\sqrt{\frac{n}{b^{1/2}}} \left[\hat{t} - t_0 - \left\{ -\frac{E\left(\hat{j}^{(1)}\left(t_0\right)\right)}{\hat{j}^{(2)}\left(\stackrel{\circ}{t}\right)} \right\} \right] \\
= -\sqrt{\frac{n}{b^{1/2}}} \left[\frac{\hat{j}^{(1)}\left(t_0\right) - E\left\{\hat{j}^{(1)}\left(t_0\right)\right\}}{\hat{j}^{(2)}\left(\stackrel{\circ}{t}\right)} \right].$$
(A29)

Then, by Lemmata A.6 and A.8, the leading bias term for \hat{t} becomes

$$-\frac{E\left\{\hat{j}^{(1)}(t_{0})\right\}}{\hat{j}^{(2)}\left(\overset{\circ}{t}\right)} = -b + o_{p}(b).$$

To demonstrate asymptotic normality of $\sqrt{n/b^{1/2}}(b^{3/2}/\Delta)[\hat{J}^{(1)}(t_0) - E\{\hat{J}^{(1)}(t_0)\}]$, we also check Lyapunov's condition. This quantity can be expressed as

$$\sum_{i=1}^{n} Y_{i} := \sum_{i=1}^{n} \sqrt{\frac{b^{5/2}}{n\Delta^{2}}} \{H_{i} - E(H_{i})\}.$$

Then, by C_r -inequality, Jensen's inequality (due to the convexity of y^3 for $y \ge 0$) and Lemma A.9,

$$E|Y_i|^3 \le 8\left(\frac{b^{5/2}}{n\Delta^2}\right)^{3/2} E|H_i|^3 = O\left(n^{-3/2}b^{-1/4}\right)$$

It also follows from Lemma A.7 that $Var(Y_i) = O(n^{-1})$. Therefore,

$$\frac{\sum_{i=1}^{n} E|Y_i|^3}{\left\{\sum_{i=1}^{n} Var(Y_i)\right\}^{3/2}} = O\left(\frac{1}{\sqrt{nb^{1/2}}}\right) \to 0,$$

and Lyapunov's condition is established.

Now we are allowed to employ a central limit theorem, in conjunction with Lemma A.7, to obtain

$$\sqrt{\frac{n}{b^{1/2}}} \left(\frac{b^{3/2}}{\Delta}\right) \left[\hat{j}^{(1)}\left(t_{0}\right) - E\left\{\hat{j}^{(1)}\left(t_{0}\right)\right\}\right] \stackrel{d}{\to} N\left(0, \frac{3}{2\sqrt{\pi}t_{0}^{5/2}}\left\{\frac{f(t_{0}^{-}) + f(t_{0}^{+})}{2}\right\}\right).$$

Then, it holds that for the right-hand side of (A29),

$$-\sqrt{\frac{n}{b^{1/2}}} \left[\frac{\left(b^{3/2}/\Delta \right) \left\{ \hat{j}^{(1)}\left(t_{0}\right) - E\left(\hat{j}^{(1)}\left(t_{0}\right) \right) \right\}}{\left(b^{3/2}/\Delta \right) \hat{j}^{(2)}\left(\overset{\circ}{t} \right)} \right] \stackrel{d}{\to} N\left(0, \frac{3\sqrt{\pi}t_{0}^{1/2}}{4d_{0}^{2}} \left\{ \frac{f(t_{0}^{-}) + f(t_{0}^{+})}{2} \right\} \right)$$

by Lemma A.8. This completes the proof.