

JUMP DETECTION IN GENERALIZED ERROR-IN-VARIABLES REGRESSION WITH AN APPLICATION TO AUSTRALIAN HEALTH TAX POLICIES¹

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Without measurement errors in predictors, discontinuity of a nonparametric regression function at unknown locations could be estimated using a number of existing approaches. However, it becomes a challenging problem when the predictors contain measurement errors. In this paper, an error-in-variables jump point estimator is suggested for a nonparametric generalized error-in-variables regression model. A major feature of our method is that it does not impose any parametric distribution on the measurement error. Its performance is evaluated by both numerical studies and theoretical justifications. The method is applied to studying the impact of Medicare Levy Surcharge on the private health insurance take-up rate in Australia.

1. Introduction. This paper is motivated by our attempt to study the impact of the Medical Levy Surcharge (MLS) tax policy on the take-up rate of the private health insurance (PHI) in Australia. People in Australia are liable of MLS (which is about 1 percent of their annual taxable incomes) if they do not buy PHI and their annual taxable incomes are above a certain level. For example, the income threshold for single individuals was \$50,000 per annum in the 2003–2004 financial year, where the dollar sign “\$” used here and throughout the paper represents the Australian Dollar (AUD). The major purposes of this tax policy were to give people more choices of health insurance and to take a certain pressure off the public medical system. It was expected that this policy would generate a jump in the PHI take-up rate around the taxable income threshold. The size of the jump could be used to evaluate the impact of the policy on the PHI take-up rate. However, the jump location may not be exactly at the threshold for the reasons given below. First, the \$500 MLS at the threshold could be lower than the net cost of PHI, and taxpayers usually consider buying PHI only when the MLS exceeds the cost of PHI at higher income levels. Second, the MLS is collected when taxpayers file their tax returns after the financial year is finished, while the decision to buy the PHI should be made before the financial year starts. Because it is difficult for people to predict their taxable incomes accurately, they may not be aware of the

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MLS issue until it occurs. So, the jump location is actually unknown and needs to be estimated properly before the jump size can be estimated. To estimate the jump location accurately is also helpful for understanding the demand for PHI in Australia. This is because from the costs of the PHI and the difference between the estimated jump point and the threshold, one can infer the true value of PHI to the taxpayers.

There is a large literature using tax changes as a source of variation in the after-tax price of health insurances. Most of these studies are for the US-employer-provided health insurances. See Gruber and Poterba (1994), Finkelstein (2002), Rodríguez and Stoyanova (2004), and Buchmueller, Didardo and Valletta (2011) for a few examples. Rarely is it the case that the tax changes could be argued as exogenous. Jumps caused by policy design, such as the MLS in Australia, have been argued to be exogenous locally for the individuals around it [Lee and Lemieux (2010)].

In the statistical literature, jump detection in regression functions has been discussed by several authors, including Joo and Qiu (2009), Müller (1992, 2002), Qiu (1991, 1994), Qiu and Yandell (1998), Wu and Chu (1993), and the references therein. See Qiu (2005) for an overview on this topic. All existing jump detection methods assume that the explanatory variable does not have any measurement error involved. Meanwhile, the existing literature on the error-in-variables regression modeling assumes that the measurement error distribution is known or it can be estimated reasonably well beforehand and that the related regression function is continuous. See, for example, Carroll, Maca and Ruppert (1999), Carroll et al. (2006), Comte and Taupin (2007), Cook and Stefanski (1994), Delaigle (2008), Delaigle and Meister (2007), Fan and Masry (1992), Fan and Truong (1993), Hall and Meister (2007), Liang and Wang (2005), Staudenmayer and Ruppert (2004), Stefanski (2000), Stefanski and Cook (1995), and Taupin (2001). Our case is much more complicated. The available data to us are drawn from the “1% Sample Unit Record File of Individual Income Tax Returns” for the 2003–2004 financial year, that was developed by the Australian Tax Office (ATO) for research purposes. Out of privacy consideration, the ATO intentionally perturbed the income data by multiplying random numbers to the income data, and the true distribution of the random numbers is unrevealed. Our major task here is to estimate a jump point and the jump magnitude in a regression model when the regressor contains measurement errors with an unknown distribution. This problem is much more challenging to handle, compared to the ones discussed in the papers mentioned above. For instance, the deconvolution kernel regression estimator proposed by Fan and Truong (1993) assumes that the characteristic function of the measurement error distribution is completely known. Such detailed knowledge of the measurement error distribution is unavailable in the current PHI problem. Also, when the error distribution was misspecified in the conventional deconvolution problems, Meister (2009) pointed out that the Mean Integrated Squared Error (MISE) of the deconvolution kernel estimator was not bounded from above. Therefore, this estimator can perform badly when the error distribution is not correctly specified.

In this paper, we propose a generalized error-in-variables regression model for describing the relationship between the PHI take-up rate and a person's annual taxable income. In the model, a jump point is included to accommodate the possible abrupt impact of MLS on the PHI take-up rate. A novel jump detector is proposed as well, which takes into account the measurement errors. One feature of our method is that it does not require the measurement error distribution to be specified beforehand, making it applicable to the current PHI problem and other real problems.

The remainder of the article is organized as follows. In Section 2, our proposed model and jump detector are described in detail. In Section 3, some statistical properties of the proposed jump detector are discussed. In Section 4, its numerical performance is evaluated. In Section 5, an in-depth analysis of the PHI data is presented. Several remarks conclude the article in Section 6. Some technical details are provided in a supplementary file.

2. Proposed methodology. Let $\{(W_i, Y_i) : i = 1, \dots, n\}$ be a sample of n independent and identically distributed (i.i.d.) observations from the models described below:

(i) The conditional distribution of $Y_i | X_i = x$ has probability density function (p.d.f.) or probability mass function (p.m.f.) from the exponential family

$$(2.1) \quad \exp \left\{ \frac{y\theta(x) - b(\theta(x))}{a(\phi)} + c(y, \phi) \right\},$$

where X_i is the i th observation of the unobservable explanatory variable X , Y_i is the i th observation of the response variable Y , $\theta(x)$ is the canonical parameter when $X_i = x$, ϕ is a scale parameter, $a(\phi)$, $b(\theta(x))$, and $c(y, \phi)$ are certain functions of ϕ , $\theta(x)$, and (y, ϕ) , respectively.

(ii) W_i is the observed value of X_i with a measurement error, and their relationship can be described by the model

$$(2.2) \quad W_i = X_i + \sigma_n U_i,$$

where $\sigma_n > 0$ denotes the standard deviation of the measurement error in X_i , and U_i is the standardized measurement error with mean 0 and variance 1. It is also assumed that U_i 's are i.i.d., U_i is independent of both X_i and Y_i , the distribution of U_i , denoted as f_U , and the distribution of X_i , denoted as f_X , are both unknown.

In model (2.1), $\theta(x)$ relates Y_i to X_i . And, model (2.1) includes many commonly used distributions (e.g., Normal, Poisson, Binomial) as special cases. In the current PHI problem, because ATO perturbed the income data by multiplying each original income observation by a random number, the income variables are used in log scale so that model (2.2) with additive measurement error is appropriate. More specifically, X_i and W_i denote the true and observed annual taxable incomes in log scale, respectively, Y_i denotes the status of PHI take-up (Y_i

equals 1 when a specific person buys the PHI and 0 otherwise), and $Y_i|X_i = x$ is assumed to follow the Bernoulli distribution with the probability of success being $p(x) = P(Y_i = 1|X_i = x)$. In such cases, the quantities $\theta(x)$, $a(\phi)$, $b(\theta(x))$, and $c(y, \phi)$ can be specified as follows:

$$\begin{aligned} \theta(x) &= \log\left(\frac{p(x)}{1 - p(x)}\right), & a(\phi) &= 1, \\ b(\theta) &= \log(1 + \exp(\theta)), & c(y, \phi) &= 0. \end{aligned}$$

As discussed in Section 1, the tax policy MLS is expected to generate a jump in $p(x)$. So, a jump in $\theta(x)$ is expected as well.

In the PHI problem, it is important to estimate the jump position in $\theta(x)$ in order to study the impact of the tax policy MLS on the PHI take-up. Our proposed jump detector is described below. The true jump position of $\theta(x)$ is assumed to be at s which is unknown. Without loss of generality, let us assume that the support of f_X is $[0, 1]$ and $s \in (0, 1)$. Let $m(x)$ denote the conditional mean of Y given $X = x$. Then, it can be checked from (2.1) that

$$m(x) = E(Y|X = x) = b'(\theta(x)).$$

For any given point $x \in (2h_n, 1 - 2h_n)$, let us consider its right-sided neighborhood $[x, x + h_n]$, where $h_n > 0$ is a bandwidth parameter. When there is no measurement error in X , $m(x+) = \lim_{\Delta x \rightarrow 0+} m(x + \Delta x)$ can be estimated reasonably well [cf. Qiu (2005), Chapter 2], by

$$(2.3) \quad \frac{\sum_{i=1}^n Y_i K_r\left(\frac{X_i - x}{h_n}\right)}{\sum_{i=1}^n K_r\left(\frac{X_i - x}{h_n}\right)},$$

where K_r is a decreasing kernel function with the right-sided support $(0, 1]$. In the case when X has measurement errors involved, the estimator in (2.3) is unavailable because X_i 's are no longer observable. It may be problematic if we simply replace X_i 's by W_i 's in (2.3) because we do not know whether a specific X_i is located on the right-hand side of x or not when its observed value W_i is on the right-hand side of x , due to the measurement error. However, as demonstrated in Figure 1, the following fact can be observed: if W_i is close to the true jump point s , then the corresponding unobservable X_i is likely to be on the other side of the jump location and, consequently, Y_i follows a distribution with the parameter $\theta(X_i)$ which could be very different from $\theta(W_i)$. A one-sided kernel estimator defined in (2.3) with the X_i 's replaced by the corresponding W_i 's actually averages observations on both sides of the jump location. Thus, the impact of the measurement error could be severe in such cases. On the other hand, in the case when W_i is farther away from s , such an impact becomes smaller. Based on this fact, let us consider a one-step-right neighborhood of x , defined to be $N_{n,r}(x; h_n) := (x + h_n, x + 2h_n)$, and define

$$(2.4) \quad \widehat{m}_{n,r}(x+) = \frac{\sum_{i=1}^n Y_i K_r\left(\frac{W_i - (x + h_n)}{h_n}\right)}{\sum_{i=1}^n K_r\left(\frac{W_i - (x + h_n)}{h_n}\right)},$$

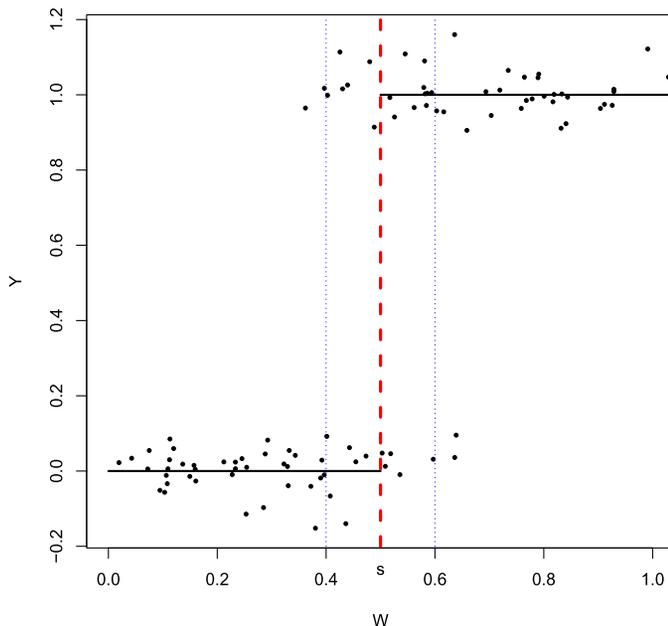


FIG. 1. The solid line denotes the conditional mean function $m(x) = E(Y|X = x)$ that has a jump at $s = 0.5$ (marked by the vertical dashed line). The dark points denote observations of (W, Y) where W is the observed value of X with measurement error involved. It can be seen that the values of X corresponding to those W values that are close to the true jump location (e.g., those fall between the two vertical dotted lines) are likely to be on both sides of the jump.

where the kernel function K_r is the same as the one in (2.3). If x is at the jump location, then $\hat{m}_{n,r}(x+)$ should be a weighted average of observations that are on the right-hand side of x since the observations in the neighborhood $N_{n,r}(x; h_n)$ are all quite far away from the jump point. This would limit the possibility of averaging observations on both sides of x , and thus diminish the impact of the measurement error. However, this estimator may still have some bias for estimating $m(x+)$ because the X values of most observations used in $\hat{m}_{n,r}(x+)$ are at least one bandwidth above x . To address this issue, let us define

$$(2.5) \quad \hat{m}_n(x+) = \frac{\sum_{i=1}^n Y_i K_r((W_i - x)/h_n) K^*(|\hat{m}_n^*(W_i+) - \hat{m}_{n,r}(x+)|/\rho_n)}{\sum_{i=1}^n K_r((W_i - x)/h_n) K^*(|\hat{m}_n^*(W_i+) - \hat{m}_{n,r}(x+)|/\rho_n)},$$

where the kernel function K_r is the same as the one in (2.3), K^* is another decreasing kernel function with support $[0, 1]$, $\rho_n = \max_{x \leq W_i \leq x+h_n} |\hat{m}_n^*(W_i+) - \hat{m}_{n,r}(x+)|$, and

$$\hat{m}_n^*(x+) = \frac{\sum_{j=1}^n Y_j K_r\left(\frac{W_j - x}{h_n}\right)}{\sum_{j=1}^n K_r\left(\frac{W_j - x}{h_n}\right)}$$

is the conventional one-sided kernel estimator of $m(x+)$. The intuitive explanation of (2.5) is as follows. From its definition, $\widehat{m}_n^*(W_i+)$ is mainly determined by observations close to W_i and these observations are mostly in the neighborhood $[x, x + h_n]$. For some of these observations, the corresponding X_i 's may be on the left-hand side of x and, thus, the impact of the measurement error on the one-sided estimator in (2.5) could be severe. On the other hand, the measurement error does not have much impact on $\widehat{m}_{n,r}(x+)$, as explained above. The bias in $\widehat{m}_{n,r}(x+)$ for estimating $m(x+)$, due to the fact that it uses many observations that are at least one bandwidth on the right-hand side of x , is considered significantly smaller than the bias in $\widehat{m}_n^*(W_i+)$ due to measurement error, because the former is due to the variation of $m(\cdot)$ in a small continuity region while the latter is caused by the jump. So, the difference $\widehat{m}_n^*(W_i+) - \widehat{m}_{n,r}(x+)$ can provide us a measure of the impact of the measurement error in W_i on estimation of $m(x+)$. If the difference is small, then the impact of the measurement error in W_i should be small. Otherwise, its impact should be large. The kernel function K^* in (2.5) aims to eliminate such an impact. Therefore, $\widehat{m}_n(x+)$ should provide a reasonable estimator for $m(x+)$. An estimator of $m(x-)$ can be constructed similarly to (2.5), which is denoted as $\widehat{m}_n(x-)$. Then, the true jump location s can be estimated by

$$(2.6) \quad \widehat{s}_n = \arg \max_{x \in (2h_n, 1-2h_n)} |\widehat{m}_n(x+) - \widehat{m}_n(x-)|,$$

and the corresponding jump magnitude d in $m(x)$ can be estimated by

$$(2.7) \quad \widehat{d}_n = \widehat{m}_n(\widehat{s}_n+) - \widehat{m}_n(\widehat{s}_n-).$$

It should be pointed out that, although the exponential family in (2.1) is parameterized using the canonical parameter $\theta(x)$, the mean parameter $m(x)$ is often easier to interpret in practice. For this reason, both the jump location and the jump magnitude are discussed above in terms of $m(x)$, instead of $\theta(x)$. In Section 3, we will show that under certain regularity conditions, \widehat{s}_n is a consistent estimator.

In the proposed jump detector (2.6), there is one parameter h_n to choose. According to Gijbels and Goderniaux (2004), jump detectors based on kernel smoothing in cases without measurement error depend heavily on the choice of bandwidth parameters. In simulation studies, the true jump location s could be known. Then, h_n can be chosen to be the one that minimizes $|\widehat{s}_n(h_n) - s|$, where \widehat{s}_n has been written as $\widehat{s}_n(h_n)$ for convenience of discussion. In practice, s is usually unknown. In such cases, we suggest the following bootstrap bandwidth selection procedure:

- For a given bandwidth value $h_n > 0$, apply the proposed jump detection procedure (2.4)–(2.6) to the original data set $\{(W_1, Y_1), (W_2, Y_2), \dots, (W_n, Y_n)\}$, and obtain an estimator of s , denoted as $\widehat{s}_n(h_n)$.
- Draw with replacement n times from the original data set to obtain the first bootstrap sample, denoted as $\{(\widetilde{W}_1^{(1)}, \widetilde{Y}_1^{(1)}), (\widetilde{W}_2^{(1)}, \widetilde{Y}_2^{(1)}), \dots, (\widetilde{W}_n^{(1)}, \widetilde{Y}_n^{(1)})\}$.
- Apply the proposed jump detection procedure (2.4)–(2.6) to the first bootstrap sample, and obtain the first bootstrap estimator of s , denoted as $\widetilde{s}_n^{(1)}(h_n)$.

- Repeat the previous two steps B times and obtain B bootstrap estimators of s : $\{\tilde{s}_n^{(1)}(h_n), \tilde{s}_n^{(2)}(h_n), \dots, \tilde{s}_n^{(B)}(h_n)\}$.
- Then, the bandwidth h_n is chosen to be the minimizer of

$$(2.8) \quad \min_{h_n} \frac{1}{B} \sum_{k=1}^B |\hat{s}_n(h_n) - \tilde{s}_n^{(k)}(h_n)|.$$

It should be pointed out that, as a byproduct of the above bootstrap bandwidth selection procedure, a confidence interval for s can be constructed from the empirical distribution of $\{\tilde{s}_n^{(1)}(\tilde{h}_n), \tilde{s}_n^{(2)}(\tilde{h}_n), \dots, \tilde{s}_n^{(B)}(\tilde{h}_n)\}$, where \tilde{h}_n denotes the bandwidth selected by the bootstrap. More specifically, for a given significance level $\alpha \in (0, 1)$, a $100(1 - \alpha)\%$ confidence interval for s is defined to be $(\tilde{s}_{n,\alpha/2}(\tilde{h}_n), \tilde{s}_{n,1-\alpha/2}(\tilde{h}_n))$, where $\tilde{s}_{n,\alpha/2}(\tilde{h}_n)$ and $\tilde{s}_{n,1-\alpha/2}(\tilde{h}_n)$ denote the $(\alpha/2)$ th and $(1 - \alpha/2)$ th quantiles of the empirical distribution of $\{\tilde{s}_n^{(1)}(\tilde{h}_n), \tilde{s}_n^{(2)}(\tilde{h}_n), \dots, \tilde{s}_n^{(B)}(\tilde{h}_n)\}$.

3. Statistical properties. In this section, we discuss some statistical properties of the proposed jump detector defined in (2.6). To this end, we have the theorem below.

THEOREM 1. *Assume that $\{(W_1, Y_1), (W_2, Y_2), \dots, (W_n, Y_n)\}$ are i.i.d. observations from models (2.1) and (2.2), and the following conditions are satisfied:*

- (1) $\theta(\cdot)$ is a bounded, piecewise continuous function with a single jump at s and its first-order derivative is also a bounded function,
- (2) $a(\cdot), b(\cdot), b'(\cdot),$ and $b''(\cdot)$ are all bounded and continuous functions,
- (3) $(b')^{-1}(\cdot)$ exists and it is strictly monotone and Lipschitz-1 continuous² in any compact subset of the range of $\theta(\cdot)$,
- (4) the support of f_X is $[0, 1]$ and $s \in (0, 1)$,
- (5) f_X is continuous, bounded, and positive on $(0, 1)$,
- (6) f_U is a continuous function and has a positive value at 0,
- (7) the kernel functions K^* and K_r are Lipschitz-1 continuous density functions with the same support $[0, 1]$,
- (8) the bandwidth h_n satisfies the conditions that $h_n = o(1)$, and $(\log n)^{1+\eta} / (n^{1/2}h_n) = o(1)$, for some $\eta > 1/2$.

Then, we have the following results:

- (i) If $\sigma_n/h_n = o(1)$, then

$$\hat{m}_n(x+) - \hat{m}_n(x-) = m(x+) - m(x-) + O\left(\frac{\sigma_n^{2/3}}{h_n^{2/3}}\right) + O(\beta_n) \quad a.s.,$$

²Given an interval $I \subset \mathcal{R}$, a function $g : I \rightarrow \mathcal{R}$ is called Lipschitz-1 continuous if there exists a real constant $C \geq 0$ such that, for all x_1 and $x_2 \in I$, $|g(x_1) - g(x_2)| \leq C|x_1 - x_2|$.

where $\beta_n = h_n + \frac{(\log n)^{1+\eta}}{n^{1/2}h_n}$.

(ii) If $\lim_{n \rightarrow \infty} \sigma_n/h_n = C$, for some $C > 0$, then

$$\begin{aligned} \widehat{m}_n(x+) - \widehat{m}_n(x-) &= m(x+) - m(x-) - (m(x+) - m(x-))C_{K,r} + O(\beta_n) \quad a.s., \end{aligned}$$

where

$$C_{K,r} = \frac{\int_0^1 K_r^{**}(w)P(|U| > w/C) dw}{\int_0^1 K_r^{**}(w) dw},$$

and

$$K_r^{**}(w) = K_r(w)K^* \left(\frac{\int_0^1 K_r(v)P(v+w < CU < v+1) dv}{\int_0^1 K_r(v)P(v < CU < v+1) dv} \right).$$

(iii) If the conditions in either (i) or (ii) hold, then we have

$$|\widehat{s}_n - s| = O(h_n), \quad a.s.$$

Theorem 1 shows that the jump detector defined in (2.6) provides a statistically consistent estimator of s under some regularity conditions. Its proof is given in a supplementary file [Kang et al. (2015)]. In result (ii) of the above theorem, if K^* and K_r are both decreasing on $[0, 1]$, then we have

$$\frac{\int_0^1 K_r^{**}(w)P(|U| > w/C) dw}{\int_0^1 K_r^{**}(w) dw} \leq \frac{\int_0^1 K_r(w)P(|U| > w/C) dw}{\int_0^1 K_r(w) dw}.$$

It can be checked that if the conventional kernel estimators are used when defining the jump detection criterion [i.e., $\widehat{m}_n^*(x+)$ is used], then the asymptotic bias for $\widehat{m}_n^*(x+) - \widehat{m}_n^*(x-)$ to estimate $m(x+) - m(x-)$ is $(m(x+) - m(x-)) \frac{\int_0^1 K_r(w)P(|U|>w/C)dw}{\int_0^1 K_r(w)dw}$, which is larger than the asymptotic bias $(m(x+) - m(x-)) \frac{\int_0^1 K_r^{**}(w)P(|U|>w/C)dw}{\int_0^1 K_r^{**}(w)dw}$ when we use $\widehat{m}_n(x+) - \widehat{m}_n(x-)$ to estimate $m(x+) - m(x-)$. Therefore, the second kernel function K^* used in (2.5) is helpful in reducing the asymptotic bias. In Theorem 1, it is required that the measurement error variance σ_n^2 tends to 0 when the sample size n increases. In the literature, it has been pointed out that this condition is needed for consistently estimating the regression function when its observations have measurement errors involved and when little prior information about the measurement error distribution is available [cf. Delaigle (2008)].

4. Numerical studies. In this section, we present some results regarding the numerical performance of the proposed jump detector described in Section 2, which are organized in two subsections. Section 4.1 includes some simulation examples related to the jump detector defined in (2.6). Section 4.2 compares the proposed jump detector to the difference-kernel-estimation (DKE) procedure that ignores the measurement error [cf. Qiu (2005), Section 3.2].

4.1. *Numerical performance of the proposed jump detector.* In this subsection, the performance of the proposed jump detector is evaluated using two simulated examples. In each example, we consider cases when the sample size n equals 100 or 200, $f_X \sim \text{Unif}[0, 1]$, and f_U is either a Normal, a Laplace, or a Uniform distribution, with $E(U) = 0$ and $\text{Var}(U)/\text{Var}(X)$ fixed at 15%. In each combination of n and f_U , the simulation is repeated 100 times. For each given bandwidth h_n , 100 values of the Absolute Error (AE), defined as $\text{AE}(h_n) = |\widehat{s}_n(h_n) - s|$, are computed. Their average is called the Mean Absolute Error (MAE) and is denoted as $\text{MAE}(h_n)$. The minimizer of $\text{MAE}(h_n)$ is called the *optimal bandwidth* and is denoted as h_{opt} . In each replicated simulation, we also compute a bandwidth value using the proposed bootstrap procedure. The average of such 100 bandwidth values is called the *bootstrap bandwidth*, denoted as h_{bt} . In each example, the values of h_{opt} , h_{bt} , $\text{MAE}(h_{\text{opt}})$, $\text{MAE}(h_{\text{bt}})$, and the empirical coverage probability (CP) of the 95% confidence interval (see its description at the end of Section 2) computed from the 100 replicated simulations are presented. In the case when $n = 100$ and f_U is Normal, the sample that gives the median value of $\text{AE}(h_{\text{opt}})$ is denoted as \mathcal{S}_{50} . For that sample, the estimated jump location by (2.6) with $h_n = h_{\text{bt}}$ and the corresponding 95% confidence interval for s will be presented. Throughout this section, if there is no further specification, the bootstrap sample size B is chosen to be 999, and K^* and K_r used in (2.4)–(2.6) are both chosen to be the Epanechnikov kernel function.

In the first example, the conditional distribution of $Y|X = x$ is assumed to follow the Normal distribution with density

$$\frac{1}{\sqrt{2\pi \times 0.01^2}} \exp\left\{-\frac{(y - \theta(x))^2}{2 \times 0.01^2}\right\},$$

where

$$\theta(x) = \begin{cases} -\sin(2\pi x), & \text{if } x \leq 0.5; \\ -\sin(2\pi x) + 1, & \text{otherwise.} \end{cases}$$

Figure 2 shows a realization of the sample \mathcal{S}_{50} , the true function $\theta(\cdot)$ (solid line), the estimated jump location \widehat{s}_n (vertical dashed line), and a 95% confidence interval for the true jump location s (vertical dotted lines). It can be seen that the proposed jump detector estimates the true jump location reasonably well. The numerical performance of the jump detector (2.6) based on 100 replicated simulations is summarized in Table 1. From the table, it can be seen that (i) the proposed jump detector estimates the true jump location reasonably well for various error distributions, (ii) the performance of \widehat{s}_n improves as the sample size n increases, (iii) the bootstrap bandwidths are slightly larger than the optimal bandwidths but they are quite close to each other, and (iv) the empirical coverage probabilities of the proposed confidence interval for s are all close to the nominal coverage probability 0.95.

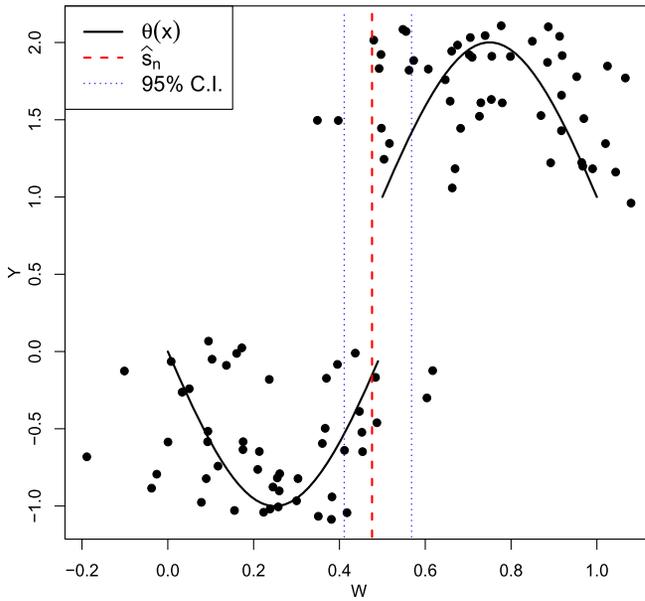


FIG. 2. A realization of \mathcal{S}_{50} in the first example with the true function of $\theta(\cdot)$ (solid line), the estimated jump location \hat{s}_n (vertical dashed line), and a 95% confidence interval for the true jump location s (vertical dotted lines).

Next, we discuss the second simulation example whose setting is made similar to that of the PHI data. Assume that the conditional distribution of $Y|X = x$ is Bernoulli with the probability of success being

$$p(x) = \begin{cases} 1 - x^2, & \text{if } x \in (0, 0.5]; \\ 0.5(1 - x)^2, & \text{if } x \in (0.5, 1). \end{cases}$$

Figure 3 shows a realization of \mathcal{S}_{50} with the true function of $p(\cdot)$ (solid line), the estimated jump location \hat{s}_n (vertical dashed line), and the 95% confidence interval for s (vertical dotted lines). It can be seen from the figure that the sample \mathcal{S}_{50}

TABLE 1
Numerical summary of the first simulation example based on 100 replicated simulations

n	f_U	h_{opt}	h_{bt}	MAE(h_{opt})	MAE(h_{bt})	CP
100	Normal	0.3000	0.3008	0.0290	0.0293	0.95
	Laplace	0.2931	0.2985	0.0292	0.0308	0.98
	Uniform	0.2767	0.2910	0.0335	0.0347	0.96
200	Normal	0.2991	0.3074	0.0232	0.0245	0.96
	Laplace	0.2902	0.3002	0.0191	0.0195	0.94
	Uniform	0.2721	0.2917	0.0232	0.0239	0.97

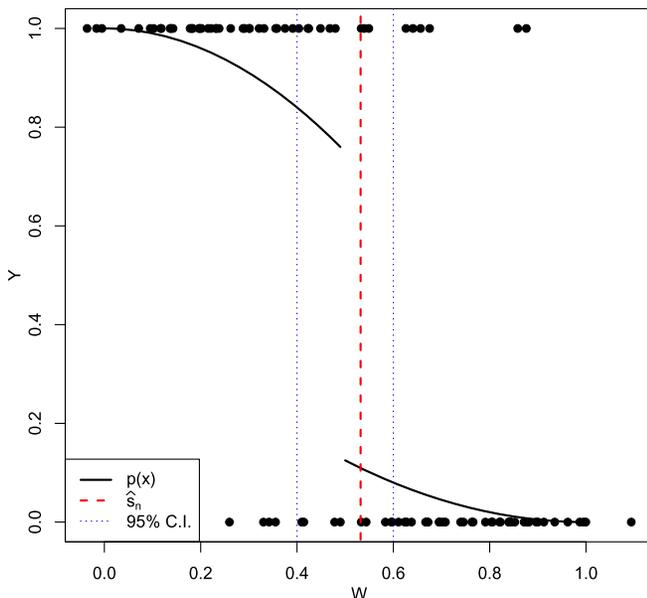


FIG. 3. A realization of \mathcal{S}_{50} in the second example with the true function of $p(\cdot)$ (solid line), the estimated jump location \hat{s}_n (vertical dashed line), and the 95% confidence interval for s (vertical dotted lines).

has quite severe measurement errors involved and that the proposed jump detector gives a reasonably good estimate of the true jump location. The numerical performance of the jump detector (2.6) based on 100 replicated simulations is summarized in Table 2. From the table, it can be seen that similar conclusions to those in the first example can be made here.

4.2. Comparison to the DKE estimator. The DKE procedure [see Section 3.2 in Qiu (2005) for a detailed discussion] provides a good estimator of the true jump position when there is no measurement error involved. In this subsection, we com-

TABLE 2
Numerical summary of the second simulation example based on 100 replicated simulations

n	f_U	h_{opt}	h_{bt}	$MAE(h_{opt})$	$MAE(h_{bt})$	CP
100	Normal	0.3329	0.3448	0.0407	0.0444	0.93
	Laplace	0.2758	0.2826	0.0397	0.0422	0.98
	Uniform	0.3203	0.3247	0.0479	0.0484	0.97
200	Normal	0.3122	0.3100	0.0363	0.0369	0.98
	Laplace	0.2820	0.2816	0.0326	0.0336	0.92
	Uniform	0.2878	0.2983	0.0352	0.0352	0.94

pare our proposed jump detector (2.6) with the DKE procedure in an artificial example with similar setup to that of the PHI data. The proposed jump detector is denoted as NEW and the DKE procedure is denoted as DKE. Assume that the conditional distribution of $Y|X = x$ is Bernoulli with the probability of success being

$$p(x) = \begin{cases} \frac{25}{36}x^2 + 0.15, & \text{if } x \in [0, 0.6); \\ 12(x - 0.8)^3 + 0.596, & \text{if } x \in [0.6, 1]. \end{cases}$$

It can be seen that $p(x)$ is a piecewise polynomial with a jump of size 0.1 at $x = 0.6$, as plotted in Figure 4. In the figure, the estimated function of $p(\cdot)$ by the local linear kernel (LLK) smoothing procedure is also shown. From the plot, it can be seen that the shape of $p(\cdot)$ is similar to that in the PHI study, which is shown in Figure 5. Note that the jump size in the PHI application is estimated to be 0.19 (cf. Section 5), while the jump size in this example is about half of that size. To make this numerical study in a similar setup to that of the PHI study, which has 9685 observations distributed in the interval $[9.8, 11.3]$, we choose the sample size n in the current example to be 6000 and the observations are in the design interval $[0, 1]$. In addition, f_U is chosen to be $N(0, 0.05^2)$, and f_X is either $\text{Unif}[0, 1]$, $\text{Beta}(2, 2)$, $\text{Beta}(3, 2)$, or $\text{Beta}(2, 3)$. In each case, the simulation is repeated 100 times, the optimal bandwidth is selected based on the 100 replicated simulations,

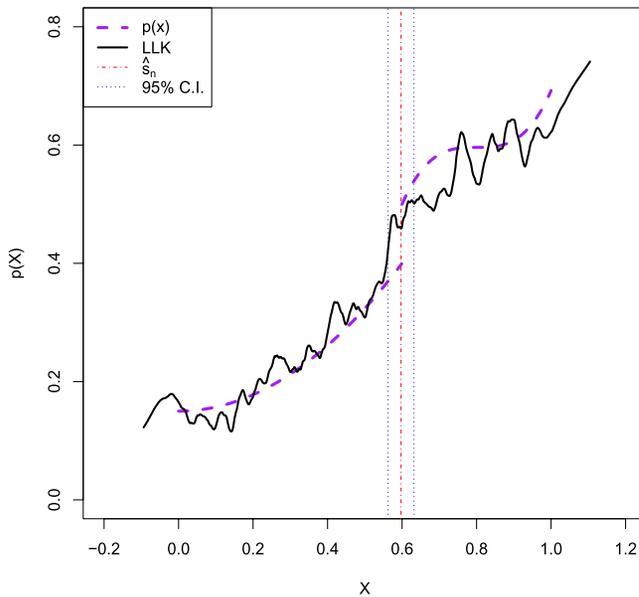


FIG. 4. The dashed line denotes the true curve of $p(\cdot)$, the solid line denotes the local linear kernel (LLK) estimate of $p(\cdot)$ using one realization of simulated data when $f_X(\cdot)$ is $\text{Unif}[0, 1]$, the vertical dot-dashed line denotes the estimated jump location \hat{s}_n , and the vertical dotted lines denote the 95% confidence interval for s .

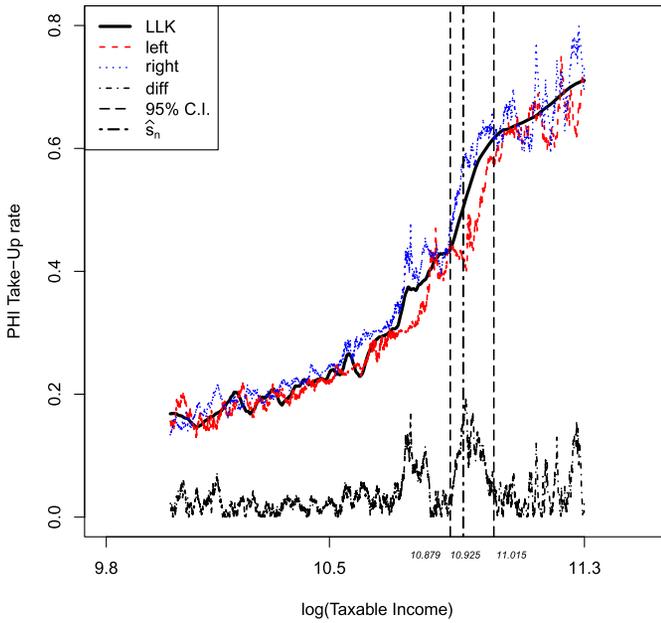


FIG. 5. The solid line denotes the local linear kernel (LLK) estimate of $p(\cdot)$ in the PHI example, the dashed line denotes the left-sided estimate of $p(\cdot)$, the dotted line denotes the right-sided estimate of $p(\cdot)$, the dot-dash line denotes the absolute difference between the two one-sided estimates of $p(\cdot)$, the long-dash vertical lines denote the 95% confidence interval for s , and the two-dash line denotes the estimated jump location \hat{s}_n .

and the mean and standard deviation of the 100 values of $|\hat{s}_n - s|$ and the 100 values of $|(\hat{m}_n(\hat{s}_n+) - \hat{m}_n(\hat{s}_n-) - 0.1|$ (i.e., the absolute bias of the estimated jump size) are computed, respectively. They are denoted as MAE, SDAE, MABJS, and SDABJS. The results are presented in Table 3.

TABLE 3

Numerical comparison of the proposed jump detector NEW with the DKE procedure based on 100 replicated simulations. MAE and SDAE denote the mean and standard deviation of the 100 values of $|\hat{s}_n - s|$. MABJS and SDABJS denote the mean and standard deviation of the 100 values of $|(\hat{m}_n(\hat{s}_n+) - \hat{m}_n(\hat{s}_n-) - 0.1|$

f_X	DKE				NEW			
	MAE	SDAE	MABJS	SDABJS	MAE	SDAE	MABJS	SDABJS
Unif[0, 1]	0.01718	0.00126	0.02598	0.00064	0.01532	0.00115	0.00511	0.00047
Beta(2, 2)	0.01547	0.00109	0.02475	0.00049	0.01329	0.00108	0.00961	0.00044
Beta(3, 2)	0.01607	0.00110	0.02593	0.00040	0.01305	0.00092	0.00494	0.00037
Beta(2, 3)	0.01810	0.00113	0.02707	0.00055	0.01690	0.00108	0.01031	0.00058

From Table 3, it can be seen that when there is measurement error involved, (i) the precision of the detected jump by the proposed jump detection procedure is better than that of the DKE procedure, across all different choices of f_X , (ii) the proposed procedure reduces the bias of the estimated jump size, which is consistent with our discussion in Section 3, and (iii) the proposed procedure has a slightly smaller variability for detecting the jump location and about the same variability for estimating the jump size, compared to the DKE procedure.

5. Analysis of the PHI data. In this section, we apply our proposed jump detector to the PHI data for evaluating the impact of MLS on the take-up rate of PHI.

The purposes of introducing PHI in Australia were to give consumers more choices and take some pressure off the public medical system. However, the PHI take-up rate by Australians was very low at the beginning when the PHI was first introduced in 1984, and the take-up rate had been declining toward the end of 1990s (the take-up rate was only about 31 percent at that time) until a series of policies (including MLS) were introduced. Impact of some of these policy measures (e.g., Lifetime Health Cover) have been studied in a few studies, including Butler (2002), Frech, Hopkins and MacDonald (2003), Palangkaraya and Yong (2005), and Palangkaraya et al. (2009). But the role of MLS has not been identified separately yet. The MLS was imposed in 1997 on high-income taxpayers who did not have private insurances. Between 1997–1998 and 2007–2008, the threshold of annual taxable income at which MLS was payable was \$50,000 for singles without children and \$100,000 for couples. For each dependent child in the household, the threshold increased by \$3000. So, people having children may lead to multiple jumps in the current PHI data. Unfortunately, we do not have information on the number of children in a family. Also, multiple jump locations within a relatively narrow range would be difficult to distinguish, given the measurement error involved in the PHI data. To mitigate the effect of multiple jumps due to people having children, this paper only focuses on singles in the current PHI data.

The data used here are from a confidentialized “1% Sample Unit Record File of Individual Income Tax Returns” for the 2003–2004 financial year, that was developed by ATO for research purposes. The file contains just over 109,000 records of individual tax returns and detailed information on income from various sources; different types of tax deductions; taxable income; and the take-up of PHI by the individuals. It also contains a limited number of demographic variables, including gender, age group, and marital status. In this paper, we focus on singles between 20 and 69 years old, who were all subject to the same income threshold of \$50,000 for the MLS. Therefore, the PHI take-up rate is expected to have a jump around that level of the annual taxable income. In the tax and transfer system or in the health insurance premium regime in Australia, there is no other differential treatment related to the PHI take-up. Other demographic covariates (such as gender and age) would not generate discontinuity in the take-up rate either as a function

of the annual taxable income. So, in the current PHI data, MLS seems to be the only factor responsible for the jump in the take-up rate.

As a method of confidentialization, ATO “perturbed” the income variables and the deductions, and provided the following information on the way the data was perturbed: several random numbers within a specified range for each individual were generated, which were converted into a rate (equal probability of being positive or negative) and which was then applied to the various components of the tax return. These rates were applied to the components in a way to try to maintain relationships with similar items. This was achieved by grouping the components into three broad categories: work or employment related income and deductions; investment income and deductions; and business and other income and deductions. Thus, there is some information about the measurement errors in the income data, but the actual distribution of the measurement errors is impossible to be identified based on the provided information. The sample was further restricted to minimize the number of income sources/deduction sources so that the distribution of the error term could be more homogeneous, according to the following criteria: (1) Only those who had positive earnings as the only sources of income were selected; (2) Individuals whose taxable income was not positive (which means their total tax deductions were not less than their earnings) were dropped; and (3) We further dropped individuals whose nonwork related deductions formed a significant part of their taxable income—specifically, we dropped those individuals whose work related deductions were less than 90 percent of earnings when the total deductions were more than 10 percent of earnings; whose total deductions were over 50 percent of earnings; or whose total deductions were all nonwork related and the total deductions were over 10 percent of their earnings.

The final sample for analysis contains 9685 records of individual tax returns. By a preliminary analysis, we found that about 26% of singles bought PHI in 2003–2004, and the PHI take-up rates for those whose annual taxable incomes were below \$50,000 and those whose annual taxable incomes were above that level were quite different. The PHI take-up rate for the former group was about 21% and it was about 57% for the latter group. Because ATO perturbed the income data by multiplying each original income observation by a random number, we used the income variable in log scale in our analysis, so that the additive measurement error assumption in (2) is valid here.

We then use our proposed jump detector (7) to estimate the jump position, in which the possible jump location is searched within [10, 11.25] (or, equivalently, [\$22,026, \$76,879] of annual taxable income). The results are shown in Figure 5, where the estimated function of $p(\cdot)$ by the local linear kernel (LLK) smoothing procedure is shown by the solid line, the left-sided and right-sided estimates of $p(\cdot)$ are shown by the dashed and dotted lines, respectively, their difference is shown by the dot-dashed line at the bottom of the plot, and the jump location estimate \hat{s}_n and the corresponding 95% confidence interval for s are shown by the vertical dot-dash and long-dash lines, respectively. In the plot, the related estimates look noisier near the right end because there were fewer people who had high incomes.

The estimated jump location is $\hat{s}_n = 10.9255$ ($\approx \$55,575$). The bandwidth chosen by the bootstrap procedure is 0.0792. The 95% confidence interval for s computed by the proposed bootstrap procedure is (10.8792, 11.0158) [$\approx (\$53,061, \$60,827)$], which implies that the true jump location s is significantly larger than 10.8198 ($\approx \$50,000$). This finding confirms our intuition that people usually act later than they are hit by the MLS. The estimated jump magnitude by (2.7) is 0.19. This number shows that the local effect at the MLS tax policy discontinuity is quite big. For individuals with only one income source, the policy can be considered locally exogenous because the observations to the left and right of (but close to) the jump position are more or less homogeneous except the policy treatment. It implies that, among the individuals whose annual taxable income is around \$55,575, MLS brings about an extra 19% of them onto the private health system. This also implies a negative price elasticity of PHI demand since the jump in the take-up rate can be seen as a response to a price discount in the premium.

6. Concluding remarks. We have proposed a generalized error-in-variables jump regression model for describing the relationship between people's annual taxable income and the PHI take-up rate in Australia. A novel jump detector is proposed as well, which can accommodate the possible measurement errors. A major feature of the proposed method is that it does not require much prior knowledge on the measurement error distribution, making it applicable in practice. Its performance is evaluated by both numerical studies and theoretical justifications. By the proposed method, we found that the actual jump in the PHI take-up rate, caused by the MLS tax policy, occurred at a larger taxable income value than the threshold value used in the policy.

There is much room for further improvement of the current method. First, the proposed jump detection method assumes that there is a single jump point at an unknown location. By the framework of jump regression analysis [cf. Qiu (2005)], it might be possible to extend it to cases when there are multiple jump points and the number of jump points could be either known or unknown. Second, this paper focuses on jump detection only. It requires much future research to develop an appropriate method to estimate a jump regression function from observed data with measurement errors. Third, it might be important to extend the current method to higher-dimensional cases.

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SUPPLEMENTARY MATERIAL

Supplement to “Jump detection in generalized error-in-variables regression with an application to Australian health tax policies” (DOI: [10.1214/15-AOAS814SUPP](https://doi.org/10.1214/15-AOAS814SUPP); .pdf). This supplemental file mainly gives the proof of Theorem 1.

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