

**Statistica Sinica Preprint No: SS-12-304R2**

<b>Title</b>	Estimation of ordinary differential equation parameters using constrained local polynomial regression
<b>Manuscript ID</b>	SS-12-304R2
<b>URL</b>	<a href="http://www.stat.sinica.edu.tw/statistica/">http://www.stat.sinica.edu.tw/statistica/</a>
<b>DOI</b>	10.5705/ss.2012.304
<b>Complete List of Authors</b>	A. Adam Ding and Hulin Wu
<b>Corresponding Author</b>	Hulin Wu
<b>E-mail</b>	hwu@bst.rochester.edu
Notice: Accepted version subject to English editing.	

## Estimation of Ordinary Differential Equation Parameters Using Constrained Local Polynomial Regression

A. Adam Ding<sup>1</sup> and Hulin Wu<sup>2</sup>

<sup>1</sup>*Department of Mathematics, Northeastern University,  
360 Huntington Ave., Boston, MA 02115, U.S.A.*

*Email: a.ding@neu.edu*

<sup>2</sup>*Department of Biostatistics and Computational Biology,  
University of Rochester School of Medicine and Dentistry,  
601 Elmwood Avenue, Box 630, Rochester, New York 14642, U.S.A.*

*Email: Hulin.Wu@urmc.rochester.edu*

*Abstract:* We propose a new method to use a constrained local polynomial regression to estimate the unknown parameters in ordinary differential equation models with a goal of improving the smoothing-based two-stage pseudo-least squares estimate. The equation constraints are derived from the differential equation model and are incorporated into the local polynomial regression in order to estimate the unknown parameters in the differential equation model. We also derive the asymptotic bias and variance of the proposed estimator. Our simulation studies show that our new estimator is clearly better than the pseudo-least squares estimator in estimation accuracy with a small price of computational cost. An application example on immune cell kinetics and trafficking for influenza infection further illustrates the benefits of the proposed new method.

*Key words and phrases:* Constrained optimization; Local polynomial smoothing; Ordinary differential equation.

**Corresponding author:** Hulin Wu, Ph.D., Professor  
Department of Biostatistics and Computational Biology  
University of Rochester School of Medicine and Dentistry  
601 Elmwood Avenue, Box 630, Rochester, New York 14642, USA

Tel. 585-275-6767, Fax 585-273-1031

Email: Hulin.Wu@urmc.rochester.edu

## 1 Introduction

Differential equations are widely used to describe and quantify dynamic systems in many scientific fields. The so-called inverse problem of differential equation models, i.e., the estimation of unknown parameters based on experimental data of state variables, is quite challenging because the standard nonlinear least squares method may fail due to convergence problems, local minima and high computational cost. Recently alternative methods based on nonparametric smoothing have been proposed and investigated in statistical literature by Poyton, Varziri, McAuley, McLellan & Ramsay (2006), Ramsay, Hooker, Campbell & Cao (2007), Chen & Wu (2008), Liang & Wu (2008), Brunel (2008). These alternative methods intend to improve the computational efficiency and stability of the nonlinear least squares method with a price of reduced estimation accuracy.

A general nonlinear ordinary differential equation model can be written as

$$\frac{dX(t)}{dt} = F\{X(t); \theta\} \quad (1.1)$$

where  $X(t) = \{X_1(t), \dots, X_d(t)\}^T$  is a  $d$ -dimensional state vector,  $\theta = (\theta_1, \dots, \theta_q)^T$  is a  $q$ -dimensional vector of unknown parameters, and  $F(\cdot) = \{F_1(\cdot), \dots, F_d(\cdot)\}^T$  is a known nonlinear function vector. Note that the proposed methodology with minor modifications is also applicable to more general differential equations with input variables. The process  $X(t)$  is usually measured with noise and we observe

$$Y(t) = X(t) + e(t) \quad (1.2)$$

where the measurement error  $e(t)$  is independent of  $X(t)$  with mean zero and a covariance matrix  $\Sigma_e$ .

Denote the solution to the differential equation (1.1) as  $X(t; \theta)$ . Generally  $X(t; \theta)$  does not have an analytic solution and needs to be obtained by solving the differential equations numerically. This results in computationally intensive and often numerically unstable estimation for parameter  $\theta$ . To avoid numerically solving the differential equations, the nonparametric smoothing techniques were applied to the observed process to estimate the parameters  $\theta$  via multiple-stage procedures in Poyton, Varziri, McAuley, McLellan & Ramsay (2006), Ramsay, Hooker, Campbell & Cao (2007), Chen & Wu (2008), Liang & Wu (2008) and Brunel (2008). Particularly, Liang & Wu (2008) proposed using the local polynomial estimation as the smoothing technique in the first stage,

and obtained the pseudo-least square estimator for  $\theta$  in the second estimation stage. The differential equation (1.1) was only used in the second estimation stage while the first stage of the local polynomial smoothing did not use the information of differential equations, which results in a significant reduction of estimation efficiency of the pseudo-least squares estimator compared to the nonlinear least squares estimator. In this paper, we intend to propose a new approach to improve the Liang and Wu's pseudo-least squares estimator by creatively combining the local polynomial smoothing and differential equation information. We expect that the new method may gain more in estimation accuracy with a small price in computational cost.

## 2 Differential equation-Constrained Local Polynomial Regression

### 2.1 Notation and method

We assume that the process  $Y(t)$  is observed at time points  $t_1, t_2, \dots, t_n$ . So the measurement model (1.2) can be rewritten as

$$Y_i = Y(t_i) = X(t_i) + e(t_i), \quad i = 1, \dots, n. \quad (2.1)$$

For notational simplicity, we present our model and method for the univariate case, i.e.,  $d = 1$ . However, the proposed methodologies and theoretical results can be easily extended to the general case of  $d > 1$ . In particular, we will illustrate this point in our simulation studies and real data analysis by applying the proposed method to the multivariate cases in Section 3.

We can estimate  $X(t)$  and its derivative at any time point  $t$  by the nonparametric local polynomial smoothing of observed  $Y_i$ s,  $i = 1, \dots, n$ . Particularly, we can obtain the smoothing estimates for the process  $\hat{X}(t_k^*)$  and its derivative  $\hat{X}'(t_k^*)$  over a grid of time points  $t = t_1^*, t_2^*, \dots, t_m^*$ . Liang & Wu (2008) proposed a two-stage estimation procedure for differential equation parameter estimation: (1) use the local polynomial smoothing over the grid of observed time points  $t_i$  to yield estimates for  $\hat{X}(t_i)$  and  $\hat{X}'(t_i)$ ,  $i = 1, 2, \dots, n$  in the first stage; and (2) estimate differential equation parameters  $\theta$  using the pseudo-least squares estimator,

$$\hat{\theta}_{PLS} = \underset{\theta}{\operatorname{argmin}} \sum_{i=1}^n [\hat{X}'(t_i) - F\{\hat{X}(t_i); \theta\}]^2 \omega(t_i),$$

where  $\omega(t_i)$  is an appropriate weight function. In general, we can extend the Liang and Wu's procedure over a time grid of size  $m$  which can be larger than the number of original measurements  $n$ . That is,

$$\hat{\theta}_{PLS} = \underset{\theta}{\operatorname{argmin}} \sum_{k=1}^m [\hat{X}'(t_k^*) - F\{\hat{X}(t_k^*); \theta\}]^2 \omega(t_k^*).$$

This modified pseudo-least squares estimator is shown to converge at  $n^{-1/2}$  rate (Liang & Wu, 2008, 2010; Fang, Wu & Zhu, 2011).

Notice that Liang & Wu (2008)'s first stage smoothing was done without using the differential equation information. The differential equation was only used in the second stage to estimate  $\theta$  based on the first stage smoothing results. The separation of these two stages results in a significant reduction in estimation accuracy of the differential equation parameters. Poyton, Varziri, McAuley, McLellan & Ramsay (2006) and Ramsay, Hooker, Campbell & Cao (2007) used a spline approach to combine the smoothing stage with the differential equation information together, which produced a more accurate and stable estimate. They tried to minimize a criterion combining the residuals in smoothing fits to observations  $Y_i$ 's and deviation of the smoothing fits from the differential equation model. Motivated by these ideas, in this paper we propose to incorporate the differential equation information into the local polynomial smoothing to estimate the differential equation parameters  $\theta$  jointly with the state variable  $\hat{X}(t_k^*)$  and  $\hat{X}'(t_k^*)$ ,  $k = 1, 2, \dots, m$ . We expect this new method to improve upon the Liang-Wu's pseudo-least squares estimator in estimation accuracy with a small price of computational cost.

The standard local  $p$ th-order polynomial regression estimates  $X(t)$  and its derivative up to order  $p$  at time  $t$  can be obtained by minimizing the objective function

$$\sum_{i=1}^n \left\{ Y_i - \left( \alpha + \sum_{j=1}^p \beta_j (t_i - t)^j \right) \right\}^2 K_h(t_i - t) \quad (2.2)$$

where  $K(\cdot)$  is a symmetric kernel function,  $K_h(\cdot) = K(\cdot/h)/h$ , and  $h$  is the bandwidth. Then  $X(t)$  can be estimated by that of  $\alpha$  and the derivatives  $X^{(j)}(t)/j!$  can be estimated by those of  $\beta_j$ ,  $j = 1, 2, \dots, p$  (Fan & Gijbels, 1996). Considering the differential equation (1.1), we notice that the local polynomial coefficients  $(\alpha, \beta_1)$  in (2.2) should satisfy  $\beta_1 = F(\alpha; \theta)$ . The higher derivatives  $X^{(j)}(t)$  can similarly be expressed as functions of  $X(t)$  through equation (1.1). For example, denote  $D_X(X; \theta) = \partial F(X; \theta)/\partial X$ , then

$$\frac{d^2 X(t)}{dt^2} = \frac{\partial}{\partial X} F\{X(t); \theta\} \frac{dX(t)}{dt} = D_X\{X(t); \theta\} F\{X(t); \theta\}.$$

Denote  $F^{(1)}(X; \theta) = D_X(X; \theta)F(X; \theta)$ , differential equation (1.1) implies that we should restrict  $\beta_2 = F^{(1)}(\alpha; \theta)/2$ . Similarly, let  $D_X^{(j)}(X; \theta) = \partial F^{(j)}(X; \theta)/\partial X$ , and  $F^{(j)}(X; \theta) = D_X^{(j-1)}(X; \theta)F(X; \theta)$  with  $F^{(0)}(X; \theta) = F(X; \theta)$  and  $D_X^{(0)}(X; \theta) = D_X(X; \theta)$ , then we have

$$X^{(j)}(t) = \frac{d^j X(t)}{dt^j} = F^{(j-1)}\{X(t); \theta\}.$$

Thus, we have the general differential equation constraints

$$\beta_j = \frac{X^{(j)}(t)}{j!} = \frac{F^{(j-1)}\{X(t); \theta\}}{j!} = \frac{F^{(j-1)}\{\alpha; \theta\}}{j!}, \quad j = 1, \dots, p. \quad (2.3)$$

Hence, after plugging in the above differential equation constraints, the objective function of the local polynomial regression (2.2) can be reformulated as follows,

$$\sum_{i=1}^n [Y_i - \{\alpha + \sum_{j=1}^p \frac{F^{(j-1)}(\alpha; \theta)}{j!} (t_i - t)^j\}]^2 K_h(t_i - t). \quad (2.4)$$

The optimization of (2.4) jointly over  $\alpha$  and  $\theta$  provides estimates  $\hat{\alpha} = \hat{X}(t)$  and  $\hat{\theta}$  simultaneously.

However, the optimization of (2.4) is unlikely to provide a good estimate for  $\hat{\theta}$  since it only uses the differential equation constraint of  $X(t)$  at one time point  $t$ . Following the ideas in Liang & Wu (2008) and Brunel (2008), we could estimate the differential equation parameter  $\theta$  by integrating the above objective function over the grid of time points  $t = t_1^*, t_2^*, \dots, t_m^*$ . That is, we can minimize the objective function

$$\sum_{k=1}^m \sum_{i=1}^n [Y_i - \{\alpha_k + \sum_{j=1}^p \frac{F^{(j-1)}(\alpha_k; \theta)}{j!} (t_i - t_k^*)^j\}]^2 K_h(t_i - t_k^*) \omega(t_k^*), \quad (2.5)$$

with respect to  $\xi = (\alpha_1, \dots, \alpha_m, \theta)^T$ , where  $\omega(t_k^*)$  are nonnegative weights over the time grid as suggested by Brunel (2008) and the bandwidth  $h$  can be determined by the cross-validation approach or the plug-in method as suggested by Liang & Wu (2008). The  $\hat{\xi}$  that minimizes (2.5) is called the differential equation constrained local polynomial estimator.

For a general nonlinear function  $F$  of the differential equation model, the optimization of (2.5) becomes a nonlinear minimization problem, thus we may lose the computational efficiency of the original local polynomial fitting. To solve this problem, we consider a linear estimator that results from one iteration of the Gauss-Newton optimization

of (2.5) at a previous estimate  $\xi^* = (\alpha_1^*, \dots, \alpha_m^*, \theta^*)^T$ . In matrix notation, the objective function (2.5) is  $[Y - G(\xi)]^T W [Y - G(\xi)]$ , where  $Y = (Y_1, \dots, Y_n, \dots, Y_1, \dots, Y_n)^T$  is a  $(nm)$ -dimensional vector with the observations  $Y_i$ 's repeated  $m$  times,  $G = (G_{1,1}, \dots, G_{n,1}, \dots, G_{1,m}, \dots, G_{n,m})^T$  with

$$G_{i,k}(\xi) = G_{i,k}(\alpha_k, \theta) = \left\{ \alpha_k + \sum_{j=1}^p \frac{F^{(j-1)}(\alpha_k; \theta)}{j!} (t_i - t_k^*)^j \right\}$$

and  $W$  is a  $nm \times nm$  diagonal weight matrix, that is,

$$\text{Diag}\{\omega(t_1^*)K_h(t_1 - t_1^*), \dots, \omega(t_1^*)K_h(t_n - t_1^*), \dots, \omega(t_m^*)K_h(t_1 - t_m^*), \dots, \omega(t_m^*)K_h(t_n - t_m^*)\}.$$

Let  $J = (\partial G / \partial \alpha_1, \dots, \partial G / \partial \alpha_m, \partial G / \partial \theta_1, \dots, \partial G / \partial \theta_q)_{\xi = \xi^*}$  denote the  $nm \times (m + q)$  Jacobian matrix evaluated at  $\xi = \xi^*$ . Then a Gauss-Newton iteration minimizes (2.5) with  $G(\xi)$  replaced by its linear approximation  $G(\xi^*) + J(\xi - \xi^*)$ . This results in the weighted linear least squares estimator

$$\hat{\xi} = (J^T W J)^{-1} J^T W \tilde{Y}, \quad (2.6)$$

where  $\tilde{Y} = Y - G(\xi^*) + J\xi^*$  is a  $(nm)$ -dimensional vector.

Note that the selection of bandwidth  $h$  and  $m$  is an important issue for practice. Here we suggest to select the bandwidth  $h$  using the plug-in method according to the recommendations by Liang and Wu (2008). It works very well in our numerical simulations and real data analysis in Section 3. Selection of  $m$  for data-augmentation is less critical based on our simulation results. In theory, the larger  $m$  is better if the computational cost does not increase too much. Thus, we can select  $m$  as large as we can afford the computational cost in practice. In addition, our proposed method can be adapted to handle the case with partially observed state variables or observed functions of state variables in principle. But in this case, it may be difficult to find the initial values for the unobserved state variables to implement the proposed estimation algorithm.

## 2.2 Asymptotic property for $\hat{\theta}$

**Theorem 1** *We assume the following technical conditions*

(1) *The differential equation (1.1) holds over a time interval  $[a_0, b_0]$  and have a bounded solution  $X(t)$ . We observe  $Y_i(t)$  from model (2.1) at  $t = t_i \in [a_0, b_0]$ ,  $i = 1, \dots, n$ . The differential equation parameters  $\theta$  are jointly estimated with  $\alpha_i = X(t_i^*)$  over a time grid  $t_i^* \in [a_0, b_0]$ ,  $i = 1, \dots, m$ . The resulting estimator  $\hat{\xi}$  is given by (2.6) with the*

linearization at a starting value  $\xi^* = (\alpha_1^*, \dots, \alpha_m^*, \theta^*)^T$ .

(2) The starting value is an estimator  $\xi^*$  such that  $|\xi^* - \xi| = O_p(n^{-\delta})$  for some  $\delta > 1/4$ .

Here  $|\cdot|$  is the  $L_\infty$  norm.

(3) The function  $F(x)$  in differential equation (1.1) has bounded  $p$ -th order derivative.

(4)  $n \rightarrow \infty$ ,  $h \rightarrow 0$ ,  $nh \rightarrow \infty$  and  $m \rightarrow \infty$ .

(5) The kernel function  $K \geq 0$  is compactly supported and bounded. Denote the moments of  $K$  by  $\mu_j(K) = \int K(u)u^j du$ . Then  $\mu_0(K) = \int K(u)du = 1$ , and all odd-order moments  $\mu_j(K) = 0$  vanish.

(6) The observation time points  $t_1, \dots, t_n$  and fitted time points  $t_1^*, \dots, t_m^*$  follow a distribution with densities  $f(t)$  and  $f_g(t)$ ,  $t \in [a_0, b_0]$ , respectively. Over the time interval  $t \in [a_0, b_0]$ ,  $f(t) > 0$  and  $f_g(t) > 0$  are bounded with continuous derivatives  $f'(t)$  and  $f'_g(t)$ .

(7) The weight function  $\omega(t) \geq 0$  is bounded over the time interval  $t \in [a_0, b_0]$ .

Then conditional on the observation time points  $t_1, \dots, t_n$ , fitted time points  $t_1^*, \dots, t_m^*$  and  $\xi^*$ , the differential equation parameter estimator  $\hat{\theta}$  has conditional bias

$$\text{Bias}(\hat{\theta}) = o_p(n^{-1/2}) + O_p(h^{p+1}) \quad p \text{ odd}, \quad \text{Bias}(\hat{\theta}) = o_p(n^{-1/2}) + O_p(h^p) \quad p \text{ even},$$

and conditional variance  $\text{var}(\hat{\theta}) = O_p((nmh^3)^{-1} + (nh)^{-1})$  if  $\omega(a_0) \neq 0$  or  $\omega(b_0) \neq 0$ ; and  $\text{var}(\hat{\theta}) = O_p((nmh^3)^{-1} + n^{-1})$  if  $\omega(a_0) = \omega(b_0) = 0$ .

Particularly, when  $\omega(a_0) = \omega(b_0) = 0$  and  $mh^3 \rightarrow \infty$ ,

$$\text{var}(\hat{\theta}) = \frac{\sigma^2}{n} A_F^{-1} [B_F - (C_F + C_F^T)] A_F^{-1}, \quad (2.7)$$

with  $A_F = \int [F_\theta * F_{\theta T} * \omega * f * f_g](t) dt$ ,  $B_F = \int [(\omega * f_g * F_\theta)' * (\omega * f_g * F_{\theta T})' * f](t) dt$  and  $C_F = \int [(f' + f * F_X) * \omega * f_g * F_\theta * \{\omega * f_g * F_{\theta T}\}'](t) dt$ .

Here and in the following we use the shorthand notations  $[f * g](t) = f(t)g(t)$ ,  $F_X(t) = [\frac{\partial}{\partial X} F(X; \theta)](t) = \frac{\partial}{\partial X} F(X; \theta)|_{X=X(t)} = D_X(X; \theta)|_{X=X(t)}$ ,  $F_\theta(t) = [\frac{\partial}{\partial \theta} F(X; \theta)](t) = \frac{\partial}{\partial \theta} F(X; \theta)|_{X=X(t)}$  and  $F_{\theta T}(t) = [F_\theta(t)]^T$ . The proof outline of Theorem 1 is given in the Appendix and details are provided in the online supplementary materials.

We have used the random design for time points  $t_1, \dots, t_n$  and  $t_1^*, \dots, t_m^*$  in the Theorem by assuming that they follow a random distribution with densities  $f(t)$  and  $f_g(t)$ , respectively. We can also consider a fixed design so that  $\int_{t_0}^{t_i} f(t) dt = (i - 1)/(n - 1)$  and  $\int_{t_0}^{t_k^*} f_g(t) dt = (k - 1)/(m - 1)$  for  $i = 1, \dots, n$  and  $k = 1, \dots, m$ . The proof for the fixed design case is similar but more tedious. The local polynomial regression is design

adaptive for  $p$  odd, that is, the asymptotic bias and variance are the same under both the random design and the fixed design (Fan and Gijbels 1996, p68). We expect that this is true under our model setting. In either case, the parameter estimator  $\hat{\theta}$  converges at the parametric  $n^{-1/2}$  rate when  $h = o(n^{-1/2p})$  and  $m^{-1}h^{-3} = o(1)$ . Notice that the function  $\hat{\alpha}_k$  is still estimated at a nonparametric rate which is slower than  $n^{-1/2}$ . This result is similar to those obtained in Brunel (2008), Liang & Wu (2008, 2010), and Fang, Wu & Zhu (2011).

The result of this theorem is also similar to the one-step maximum likelihood approximation (Theorem 4.3 of Lehmann & Casella 1998) in some sense. The one-step Newton-Raphson iteration of the likelihood equation starting at a  $n^{-1/2}$ -consistent estimator results in a more efficient estimator. Here we show that one Gauss-Newton iteration for maximizing (2.5) starting at  $n^{-\delta}$  rate estimator ( $\delta > 1/4$ ) could result in a new  $n^{-1/2}$  rate estimator for  $\theta$ . Our one-iteration estimator (2.6) is a linear estimator, which is used to improve the Liang-Wu's pseudo-least squares (PsLS) estimator (Liang & Wu 2008, 2010). The initial estimator for  $\alpha_k$ 's can be taken as the smoothing estimator without using the differential equation information. For example, one iteration starting from the Liang-Wu's PsLS estimator  $\hat{\theta}$  and a local polynomial estimator for  $\alpha_k$ 's results in a linear estimator with a  $n^{-1/2}$  rate for  $\theta$ . The linearity may also be useful for future extension of our method to mixed-effects differential equation models for longitudinal data (Fang, Wu & Zhu, 2011).

*Remark.* From the theorem, we can see that, for a small enough  $h = o(n^{-1/(2p)})$ , the bias for the PsLS estimator and our proposed estimator is in the same order of  $o_p(n^{-1/2})$ , so the variance dominates the mean squared error of the two estimators. For  $\omega(a_0) = \omega(b_0) = 0$  and if  $m$  is chosen to be large enough so that  $mh^3 \rightarrow \infty$ , then we have an explicit expression (2.7) for the variance of our proposed estimator. For simplicity, considering the case of uniformly distributed  $t_i$ s and  $t_k^*$ s on time interval  $[0, 1]$ , then  $f(t) = f_g(t) = 1$ , and the variance  $var(\hat{\theta})$  of our estimator becomes

$$\frac{\sigma^2}{n} A_F^{-1} \left( \int [(\omega * F_\theta)' * (\omega * F_{\theta T})' - F_X * \omega * \{F_\theta * (\omega * F_{\theta T})' + (\omega * F_\theta)' * F_{\theta T}\}](t) dt \right) A_F^{-1},$$

where  $A_F$  is now  $\int [F_\theta * F_{\theta T} * \omega](t) dt$ . Compared to our estimator, the variance of Liang-Wu PsLS estimator (Liang and Wu 2010) has an extra term  $\frac{\sigma^2}{n} A_F^{-1} \int [(\omega * F_X * F_\theta)' * (\omega * F_X * F_{\theta T})'](t) dt A_F^{-1}$ , which is clearly a positive semi-definite matrix. Thus, our estimator has a smaller asymptotic variance compared to that of the PsLS estimator.

This extra term in the variance of Liang-Wu's PsLS estimator corresponds to the error propagated from the first stage estimator  $\hat{X}(t)$ .

*Remark.* As long as  $h^p$  is of smaller order than  $n^{-1/2}$ , the order  $p$  of the polynomial is not very important. As  $p$  increases, there are more terms in (2.5) and the computational burden increases. Therefore, a small value of  $p = 1$  or  $p = 2$  would be preferred in practice. In the numerical studies below, we used  $p = 2$  which is the same as that in Liang-Wu's method (2008) for fair comparisons.

### 3 Numerical Studies

In this section, first we compare the performance of the proposed method with Liang-Wu's method (2008), the method of Ramsay, Hooker, Campbell & Cao (2007) and the nonlinear least squares estimator by Monte Carlo simulations. In addition, we apply the proposed method to a real data set on immune cell trafficking for influenza infection to illustrate the usefulness of the proposed method. We measure the performance of the estimators by their average relative error (ARE) in simulation studies as

$$ARE = \sum_{i=1}^r \left| \frac{\hat{\theta}_i}{\theta} - 1 \right|$$

with  $\hat{\theta}_i$  as the estimate for  $\theta$  in the  $i$ th simulation runs with  $i = 1, 2, \dots, r$ . The computational cost and convergence are also considered in evaluating different estimation methods.

Since the Liang-Wu's pseudo-least squares estimator and the proposed new estimator in this paper are computationally efficient, they can be used as the starting point for the nonlinear least squares estimator. So this hybrid strategy may enjoy both computational efficiency of the pseudo-least squares estimator or the new estimator and high estimation accuracy of the nonlinear least squares estimator. We will also evaluate the performance of the hybrid approaches in our simulation studies.

**Example 1.** In this simulation example, we simulated the data from the FitzHugh-Nagumo system of differential equations that were originally used to model the behavior of spike potentials in the giant axon of squid neurons in FitzHugh (1961) and Nagumo, Arimoto, & Yoshizawa (1962). This model was also used for simulation studies by Ramsay, Hooker, Campbell & Cao (2007) and Liang & Wu (2008). We use this model to investigate the finite-sample behavior of the proposed method and other existing

methods. The FitzHugh-Nagumo system can be written as

$$\begin{aligned}\frac{d}{dt}X_1 &= (X_1 + X_2 - \frac{X_1^3}{3})c, \\ \frac{d}{dt}X_2 &= -\frac{X_1 - a + bX_2}{c},\end{aligned}\tag{3.1}$$

with true parameter values  $\theta = (a, b, c) = (0.34, 0.2, 3)$  in our simulations. We assume that  $X_1$  and  $X_2$  are measured over a grid of  $n = 51$  equally-spaced time points, every 0.4 time interval in the range of  $t = [0, 20]$  with measurement errors as in equation (2.1) with  $(\sigma_1, \sigma_2)$  taking as  $(0.1, 0.1)$ ,  $(0.1, 0.3)$ ,  $(0.3, 0.1)$  or  $(0.3, 0.3)$  for the measurement standard errors for  $X_1$  and  $X_2$  respectively. Thus, we obtained  $n = 51$  data points. For each simulated data set, we apply the proposed estimation method and other existing methods to obtain the following estimates: the nonlinear least squares estimator  $\hat{\theta}^{NLS}$ , Ramsay et al.'s collocation estimator  $\hat{\theta}^{col}$ , the Liang-Wu's pseudo-least squares estimators  $\hat{\theta}^{PLS}$  with  $m = n$  grid points. For fair comparisons, all these estimators used a common starting value of  $\theta = (a, b, c)$  with  $a$ ,  $b$  and  $c$  independently generated uniformly between 0 and twice the true value. That is, the starting  $\theta$  follows a uniform distribution on a cube centered at true value  $(0.34, 0.2, 3)$  and one corner at  $(0, 0, 0)$ . The proposed new estimator  $\hat{\theta}^{new}$  with  $m = n$  is intended as an improvement of the pseudo-least squares estimators  $\hat{\theta}^{PLS}$ , and was calculated by (2.6) starting at  $\hat{\theta}^{PLS}$ . For the hybrid approach, we also find the nonlinear least squares estimator  $\hat{\theta}_{PLS}^{NLS}$ ,  $\hat{\theta}_{new}^{NLS}$  and  $\hat{\theta}_{col}^{NLS}$  using respectively the Liang-Wu's pseudo-least squares estimator  $\hat{\theta}^{PLS}$ , the proposed new estimator  $\hat{\theta}^{new}$  and Ramsay et al.'s collocation estimator  $\hat{\theta}^{col}$  as the starting points. For the Liang-Wu pseudo-least squares estimator and the proposed estimator, the local quadratic polynomial smoothing was used and the piecewise linear weight function suggested in Brunel (2008) was used:  $w(t) = 1$  for  $1 \leq t \leq 19$ ;  $w(t) = t$  for  $0 \leq t \leq 1$ ;  $w(t) = 20 - t$  for  $19 \leq t \leq 20$ . The Ramsay, Hooker, Campbell & Cao (2007)'s collocation estimator  $\hat{\theta}^{col}$  was implemented using the R package *CollocInfer* (Hooker, Xiao & Ramsay, 2010) with 51 equally-spaced knots between  $t = 0$  and  $t = 20$ . The smoothing parameter for the collocation estimator was chosen as those in the FitzHugh-Nagumo system demo example in the package. For our new estimator and the Liang-Wu's pseudo-least squares estimator, we used the bandwidth recommended by Liang & Wu (2008):  $\hat{h}_{opt} \times n^{-3/35}(\log n)^{-1/16}$ . Here  $\hat{h}_{opt}$  is the optimal bandwidth for the local polynomial fitting without the differential equation constraints, and we calculated it from the R package 'lokern' (Maechler, 2010). All these methods are coded in R and run on the same computer together on the same simulated data sets for fair

comparisons.

Table 1 summarizes the average relative errors and computing times of the various estimators based on  $r = 400$  simulation runs. From these simulation results, we observe the following patterns: 1) The Liang-Wu's pseudo-least squares estimator is always most computationally efficient and always converges, but its estimation accuracy in the sense of average relative errors always has a large gap to improve compared to the best estimate. 2) The proposed new estimator improves the average relative errors of the Liang-Wu's pseudo-least squares estimator for most cases with a small price of computational cost as we intended. In particular, when the newly proposed estimator is used to initiate the nonlinear least squares estimate, it produces the best estimate in terms of the average relative errors for all the cases among all the methods. However, when the Liang-Wu's estimator is used to initiate the nonlinear least squares estimate, it may produce convergence problems and diverges in a substantial proportion of cases. 3) For the standard nonlinear least squares estimator with random starting points within the twice of the magnitudes of true parameter values, it is very unstable, and in many cases it either converges to the local minima or fails to converge. So it has a high computational cost and poor estimation accuracy as we expected. However, the nonlinear least squares estimator can be significantly improved in the sense of computational cost and estimation accuracy if the proposed new estimator is used as the initial estimate. 4) The Ramsay, Hooker, Campbell & Cao (2007)'s collocation estimator significantly improves the nonlinear least square estimator in the sense of convergence if the same random starting points are used. However, its estimation accuracy (in terms of the average relative error) is much worse than the best nonlinear least squares estimator using the proposed estimator as the starting point and its computational cost is highest in most cases among all the methods in the simulation studies. But the comparison of computational cost to Ramsay, Hooker, Campbell & Cao (2007)'s method may need to be taken with a grain of salt as it is affected by the actual implementation procedure, in particular the selection of the penalty parameter. 5) Additional simulation results from a different nonlinear differential equation model are included in the online supplementary materials, which shows similar results. In the supplementary Table 3, we also reported the standard deviation (STD) of the estimators in Table 1 here. We can see that the trend and conclusions for the STD are similar to those for the AREs, i.e., comparing to the Liang-Wu's pseudo-least squares (PLS) estimator, our new estimator has a lower

$(\sigma_1, \sigma_2)$	parameter	Estimators							
		$\hat{\theta}^{NLS}$	$\hat{\theta}^{col}$	$\hat{\theta}^{PLS}$	$\hat{\theta}^{new}$	$\hat{\theta}_{PLS}^{NLS}$	$\hat{\theta}_{new}^{NLS}$	$\hat{\theta}_{col}^{NLS}$	
(0.1,0.1)	ARE	a	2.58	7.70	4.26	5.42	1.96	1.75	1.75
		b	14.1	58.2	19.64	20.71	12.5	11.95	11.89
		c	2.32	6.07	26.82	21.16	0.69	0.37	0.37
	diverge	36.75	6.00	0	0	4.25	0.50	1.75	
	time	11.59	12.07	0.17	0.24	8.17	7.01	20.12	
(0.1,0.3)	ARE	a	6.73	10.34	6.85	8.06	3.72	2.49	2.49
		b	42.1	69.7	52.78	49.28	33.6	28.9	29.0
		c	9.08	7.73	33.95	22.31	2.44	0.55	0.56
	diverge	39.75	5.25	0	0	14.75	1.25	3.75	
	time	10.93	12.89	0.18	0.25	9.11	8.14	21.01	
(0.3,0.1)	ARE	a	5.21	13.61	10.30	8.60	5.26	4.97	5.03
		b	22.9	73.0	29.49	27.43	24.4	23.18	23.53
		c	1.91	7.71	34.08	21.70	1.42	1.05	1.06
	diverge	34.50	5.75	0	0	8.75	2.00	3.75	
	time	13.08	14.00	0.18	0.25	12.62	11.66	25.11	
(0.3,0.3)	ARE	a	6.12	13.58	11.44	10.73	5.68	5.44	5.49
		b	35.0	82.6	55.02	53.28	35.8	36.1	35.8
		c	4.21	9.04	43.33	24.44	2.44	1.41	1.49
	diverge	39.75	6.25	0	0	20.00	7.00	6.50	
	time	12.75	14.73	0.18	0.25	12.06	10.99	26.18	

Table 1: Performance (ARE) of different estimators for Example 1 with  $n = 51$  observations:  $\hat{\theta}^{NLS}$ =nonlinear least squares estimate using a random starting point;  $\hat{\theta}^{col}$ =Ramsay et al's collocation estimate using the same starting point;  $\hat{\theta}^{PLS}$ =pseudo-least squares estimate using the same starting point;  $\hat{\theta}^{new}$ =the proposed new estimate started from  $\hat{\theta}^{PLS}$ ;  $\hat{\theta}_{PLS}^{NLS}$ =nonlinear least squares estimate started from  $\hat{\theta}^{PLS}$ ;  $\hat{\theta}_{new}^{NLS}$ =nonlinear least squares estimate started from  $\hat{\theta}^{new}$ ;  $\hat{\theta}_{col}^{NLS}$ =nonlinear least squares estimate started from  $\hat{\theta}^{col}$ .

STD in all three parameters  $a$ ,  $b$  and  $c$  for the cases  $(\sigma_1, \sigma_2) = (0.3, 0.1)$  and  $(0.3, 0.3)$ ; while for the other two cases, the two methods produce mixed performance in terms of both STD and ARE (for some parameters, the PLS estimator is better and for some other parameters, our new estimate is better). However, the NLS estimator using our new estimator as the initial value always performs better for all the cases in both STD and ARE, compared to those using the PLS estimator as the initial value. Also included in the online supplementary materials are simulation results for evaluating the data augmentation size  $m$ . While the proposed new estimator's performance remains similar for larger  $m$  in most cases, increasing  $m$  does lead to improvement of AREs in a few cases. However, for the NLS estimator  $\theta_{new}^{NLS}$  using the proposed estimator as a starting point, the performance improvement is not significant. Therefore, we would recommend using  $m = n$  in practice when using the proposed estimator as the starting point for the NLS estimator.

**Example 2.** To further illustrate the usefulness of the proposed method, we apply the method to a differential equation model for the growth and migration of influenza virus-specific effector CD8+ T cells among lymph node ( $T_E^m$ ), spleen ( $T_E^s$ ), and lung ( $T_E^l$ ) of mice. The mechanistic differential equation model can be written as (Wu, Kumar, Miao, Holden-Wiltse, Mosmann, Livingstone, Belz, Perelson, Zand & Topham, 2011),

$$\begin{aligned} \frac{d}{dt}T_E^m &= [\rho_m D^m(t - \tau) - \delta_m]T_E^m - (\gamma_{ms} + \gamma_{ml})T_E^m, \\ \frac{d}{dt}T_E^s &= [\rho_s D^s(t - \tau) - \delta_s]T_E^s - \gamma_{sl}T_E^s + \gamma_{ms}T_E^m, \\ \frac{d}{dt}T_E^l &= \gamma_{ml}T_E^m + \gamma_{sl}T_E^s - \delta_l T_E^l, \end{aligned} \quad (3.2)$$

where  $D^m$  denotes the number of mature dendritic cells in the mediastinal lymph node (MLN),  $D^s$  the number of mature dendritic cells in spleen;  $\tau$  is the time delay of the effects of dendritic cells on CD8+ T cell proliferation;  $\rho_m$  and  $\rho_s$  are the proliferation rates of CD8+ T cells stimulated by per dendritic cell in MLN and spleen, respectively;  $\delta_m$ ,  $\delta_s$  and  $\delta_l$  are the disappearance rates in MLN, spleen and lung, respectively;  $\gamma_{ms}$  is the migration rate from MLN to spleen,  $\gamma_{ml}$  the migration rate from MLN to lung, and  $\gamma_{sl}$  the migration rate from spleen to lung. For this differential equation system, a total of  $n = 77$  data points at 9 distinct time points for each of the three state variables,  $(T_E^m, T_E^s, T_E^l)$ , are available (see Figure 1). The data for  $D^m$  are also available. In the analysis, data of  $D^s$  are not available and is assumed to follow a similar pattern as  $D^m$  as argued in Wu, Kumar, Miao, Holden-Wiltse, Mosmann, Livingstone, Belz, Perelson, Zand & Topham (2011). The smoothed estimates of  $D^m$  were used in the analysis.

Parameters	Estimation Methods			
	PsLS	DCLP	NLS	DCLP-NLS
$T_E^m(5)$	$4.23E+3$	$4.23E+3$	$3.96E+3$	$3.96E+3$
$T_E^s(5)$	$3.33E+3$	$3.33E+3$	$3.64E+3$	$3.66E+3$
$T_E^l(5)$	$13.1E+3$	$13.1E+3$	$13.1E+3$	$13.1E+3$
$\rho_m$	$1.95E-5$	$1.46E-5$	$1.66E-5$	$1.66E-5$
$\rho_s$	$2.18E-5$	$4.78E-5$	$4.48E-5$	$4.47E-5$
$\delta_l$	$1.53E-29$	3.96	3.96	3.97
$\gamma_{ms}$	$1.41E-1$	$1.38E-1$	$1.57E-1$	$1.57E-1$
$\gamma_{ms}$	$4.17E-5$	$6.11E-1$	$4.95E-1$	$4.96E-1$
<b>residual sum of squares</b>	112.4	18.48	15.77	15.77
<b>average time</b>	0.32	0.78	10.81	5.52

Table 2: The estimated parameter values by different procedures for CD8+ T cells data analysis. PsLS denotes the pseudo-least squares estimate; DCLP denotes the proposed differential equation constrained local polynomial estimate; NLS denotes the nonlinear least squares estimate; DCLP-NLS denotes the nonlinear least squares estimate started from the proposed estimate.

More details about the experimental design and biological background of this study can be found in Wu, Kumar, Miao, Holden-Wiltse, Mosmann, Livingstone, Belz, Perelson, Zand & Topham (2011).

To stabilize the measurement error variance, a logarithm transformation is applied. That is, we let  $X = (X_1, X_2, X_3)^\tau = (\log(T_E^m), \log(T_E^s), \log(T_E^l))^\tau$ . The differential equations can be re-expressed as

$$\begin{aligned}
 \frac{d}{dt} X_1 &= \rho_m D^m(t - \tau) - \delta_m - \gamma_{ms} - \gamma_{ml}, \\
 \frac{d}{dt} X_2 &= \rho_s D^s(t - \tau) - \delta_s - \gamma_{sl} + \gamma_{ms} \exp(X_1 - X_2), \\
 \frac{d}{dt} X_3 &= \gamma_{ml} \exp(X_1 - X_3) + \gamma_{sl} \exp(X_2 - X_3) - \delta_l.
 \end{aligned}
 \tag{3.3}$$

The differential equation model (3.3) is fitted to data from day 5 to day 14 since the influenza-specific CD8+ T cells are not produced yet from Days 0-5. The time delay is set to  $\tau = 3.08$  days, and parameters  $\delta_m$ ,  $\delta_s$  and  $\gamma_{ml}$  were set to zero according to Wu, Kumar, Miao, Holden-Wiltse, Mosmann, Livingstone, Belz, Perelson, Zand & Topham (2011).

We apply our proposed estimation method with a piece-wise linear weight function:  $w(t) = 1$  for  $6 \leq t \leq 13$ ;  $w(t) = t - 5$  for  $5 \leq t \leq 6$ ;  $w(t) = 14 - t$  for  $13 \leq t \leq 14$  to the real data set as suggested by Brunel (2008). For comparisons, we also obtained

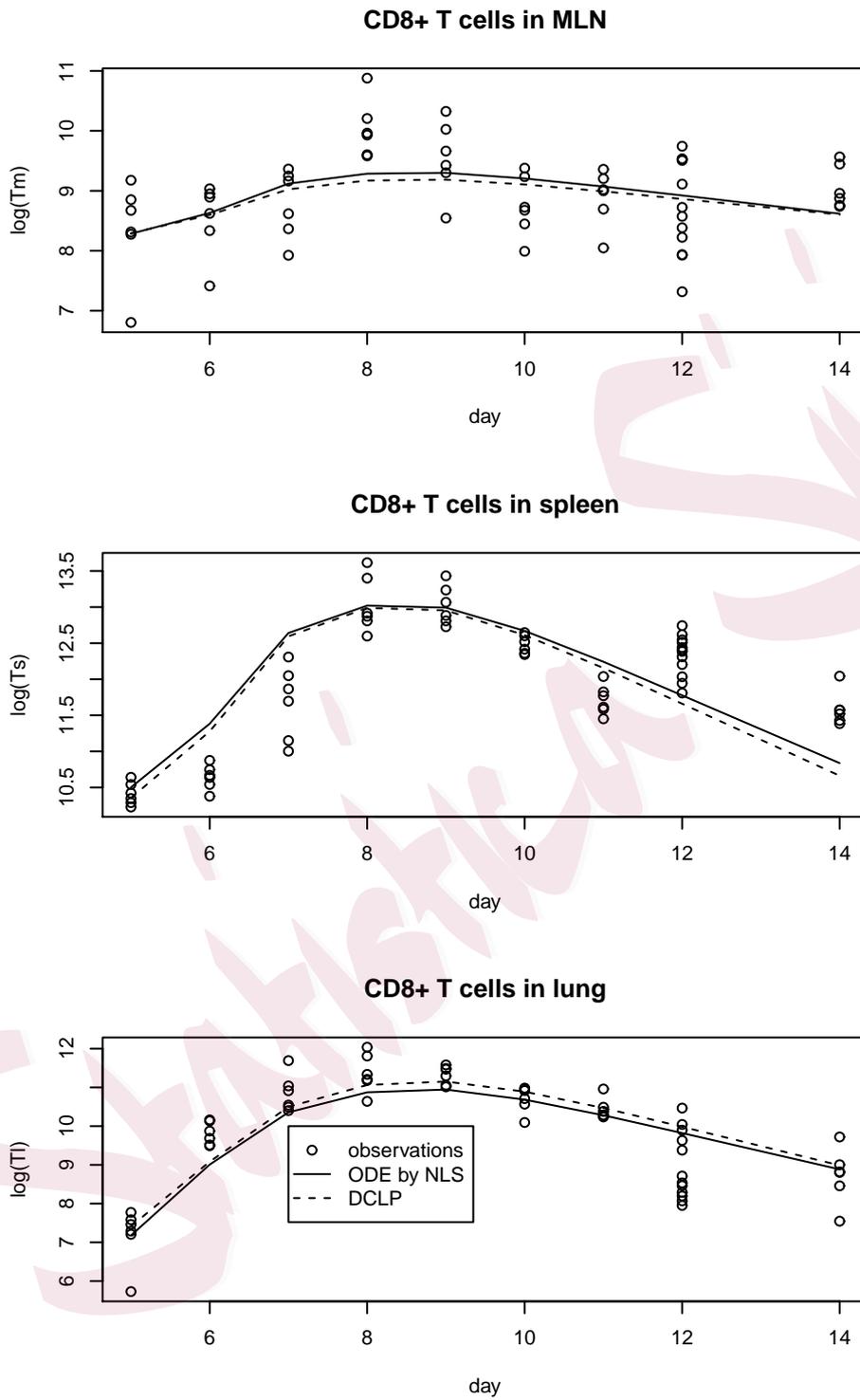


Figure 1: Data of influenza-specific CD8+ T cells in MLN, spleen and lung and the corresponding fitted curves.

the Liang-Wu's pseudo-least squares estimates and the nonlinear least squares estimates for this data set. A grid search was used to obtain the proposed differential equation constrained local polynomial estimates and the nonlinear least squares estimates. We report the results of parameter estimates for these estimation methods in Table 2 and fitted curves in Figure 1.

From these results, we can see that the proposed new differential equation constrained local polynomial estimates of kinetic parameters are much closer to the nonlinear least squares estimates compared to that of the Liang-Wu's pseudo-least squares estimates. Both the Liang-Wu's method and our new method can save computational times significantly. The residual sum of squares of our new method and the nonlinear least squares estimates is close and small while the Liang-Wu's pseudo-least squares estimate has a very large residual sum of squares. When we used the Liang-Wu's pseudo-least squares estimates as the starting point for the nonlinear least squares estimates, we failed to reach the convergence. When we used our new estimate as the starting point for the nonlinear least squares estimate, we quickly achieved the convergence in approximately half the time that the original nonlinear least squares algorithm took. Thus, this real data analysis example also demonstrates the benefits of the proposed differential equation constrained local polynomial approach.

#### 4 Concluding Remarks

In this paper, we intend to propose a new estimation method for differential equation parameters based on the differential equation constraint local polynomial regression with a goal for improving the Liang-Wu's pseudo-least squares estimate. We investigated the asymptotic properties and finite-sample behaviors of the proposed method. Our simulation studies and real data analysis show that the proposed new estimator is clearly better than the Liang-Wu's pseudo-least squares estimator in estimation accuracy with a small price of computational cost. Due to their computational efficiency, the pseudo-least squares and differential equation constrained local polynomial estimates could be used as the starting point for the more refined nonlinear least squares estimate and our simulation results also show that our new estimate is also better for this purpose. Our simulation results also demonstrate that the Ramsay, Hooker, Campbell & Cao (2007)'s collocation method is more stable and can improve the estimation accuracy of the non-

linear least squares estimate significantly, but it cannot achieve the estimation accuracy of the nonlinear least squares estimate without using our proposed estimator as the starting point, and its computational cost is highest among all the methods in our simulation studies.

Lu, Liang, Li & Wu (2011) show that the computationally efficient method such as the pseudo-least squares estimate is very useful to deal with high-dimensional differential equation models in which case the nonlinear least squares method often fails due to high computational cost and instability in computational implementations. We also expect that the proposed differential equation constrained local polynomial approach can improve the performance of the pseudo-least squares estimates in the high-dimensional case, which is a worthy future research topic.

A limitation of the proposed method is that its estimation accuracy has not reached to that of the nonlinear least squares estimate. There is still a space to improve the proposed differential equation constrained local polynomial estimate. Another limitation of the proposed method, similar to the Liang-Wu's pseudo-least squares estimator, is its requirement to measure all state variables. If there are any latent (unobserved) variables, the proposed method cannot be directly applied. However, the proposed method can be adapted to deal with the latent state variables by paying more for computational cost. More careful investigations are needed to evaluate the trade-off between the additional computational cost and the benefits. We also notice that, so far there is no good method to optimally select the bandwidth ( $h$ ) and the data augmentation size ( $m$ ), which is still an open question. We have followed the recommendations in Liang and Wu (2008) for bandwidth selection and it works well in our numerical studies. The selection of  $m$  for data-augmentation is not very critical based on our simulation results. In theory, we can select  $m$  as large as possible as long as we can afford the additional computational cost. However, the formal theoretical investigation on selection of optimal  $h$  and  $m$  is definitely another worthy future research topic.

The proposed differential equation constrained local polynomial estimator is a linearized estimator in contrast to a nonlinear estimator such as Liang-Wu's pseudo-least squares estimator. Hence it is possible to extend the approach to population differential equation models (ODE). Longitudinal dynamic (random coefficient) ODE models have been suggested by Putter et al. (2002), Huang and Wu (2006), and Huang, Liu and Wu (2006), in which the hierarchical Bayesian approach was used to estimate popula-

tion dynamic parameters in HIV dynamic models from longitudinal clinical data. Li et al (2002) proposed a spline-enhanced population model to study pharmacokinetics using a random time-varying coefficient ODE model. Guedj, Thiebaut, and Commenges (2007) used the maximum likelihood approach directly to estimate unknown parameters in random coefficient ODE models. Fang, Wu and Zhu (2011) extended the two-stage estimation method to random coefficient ODE models for longitudinal data. However, the extension of the differential equation constrained local polynomial estimator to the population mixed-effects ODE model is not trivial and remains an open research topic.

## Acknowledgement

We appreciate helpful discussions with Dr. Hongqi Xue. This work is partially supported by NIH grants, HHSN272201000055C, RO1 AI087135, and the University of Rochester CTSI(RR024160) Pilot Award.

## Appendix

### Proof of Theorem 1:

We analyze the order of estimation errors similar to the usual derivations of local polynomial regression. (For example, see section 3.7 in Fan and Gijbels 1996.) The order of some common quantities would be useful. Let  $S_{k,j} = \sum_{i=1}^n K_h(t_i - t_k^*)(t_i - t_k^*)^j$ . Then

$$S_{k,j} = nh^j f(t_k^*) \mu_j(K) [1 + o_p(1)] \quad j \text{ even}, \quad S_{k,j} = nh^{j+1} f'(t_k^*) \mu_{j+1}(K) [1 + o_p(1)] \quad j \text{ odd}, \quad (\text{A.1})$$

where  $f(t)$  is the density at  $t$  and  $\mu_j(K) = \int K(u)u^j du$ .

To consider properties of the estimator  $\hat{\xi} = (J^T W J)^{-1} J^T W \tilde{Y}$  in (2.6), we first study the matrix  $(J^T W J)^{-1}$  and  $J^T W$ . Since  $G_{i,k}(\xi)$  only depends on  $(\alpha_k, \theta)$ , the

Jacobian matrix  $J$  is sparse with many zero elements:

$$J = \begin{pmatrix} \widetilde{DX}_{1,1} & \dots & 0 & \widetilde{D\theta}_{1,1} \\ \vdots & \dots & \vdots & \vdots \\ \widetilde{DX}_{n,1} & \dots & 0 & \widetilde{D\theta}_{n,1} \\ \vdots & \ddots & \vdots & \vdots \\ 0 & \dots & \widetilde{DX}_{1,m} & \widetilde{D\theta}_{1,m} \\ \vdots & \dots & \vdots & \vdots \\ 0 & \dots & \widetilde{DX}_{n,m} & \widetilde{D\theta}_{n,m} \end{pmatrix},$$

where  $\widetilde{DX}_{i,k} = 1 + \sum_{j=1}^p \frac{(t_i - t_k^*)^j}{j!} D_{X,k}^{(j-1)}$  and  $\widetilde{D\theta}_{i,k} = \sum_{j=1}^p \frac{(t_i - t_k^*)^j}{j!} D_{\theta^T,k}^{(j-1)}$  with

$$D_{X,k}^{(j)} = D_X^{(j)}(\alpha_k^*; \theta^*), \quad D_{\theta^T,k}^{(j)} = D_{\theta^T}^{(j)}(\alpha_k^*; \theta^*) = \left( \frac{\partial}{\partial \theta_1} F^{(j)}(\alpha; \theta), \dots, \frac{\partial}{\partial \theta_q} F^{(j)}(\alpha; \theta) \right)_{\alpha=\alpha_k^*, \theta=\theta^*}.$$

Since  $p$  is fixed,  $\widetilde{DX}_{i,k}$  and  $\widetilde{D\theta}_{i,k}$  are sums of fixed number of terms. Since by (A.1), the kernel sums of  $(t_i - t_k^*)^j$  is at most of order  $O_p(nh^j)$ , the error analysis later often only need to focus on the lowest power term in  $\widetilde{DX}_{i,k}$  and  $\widetilde{D\theta}_{i,k}$ . That is, 1 and  $(t_i - t_k^*)D_{\theta^T,k}^{(0)}$  respectively.

Direct calculation shows that

$$J^T W J = \begin{pmatrix} D_{m \times m} & L_{m \times q} \\ L_{q \times m}^T & C_{q \times q} \end{pmatrix}, \quad (\text{A.2})$$

where the subscripts of the four sub-matrices denote their dimensions. The matrix  $D$  is a  $m \times m$  diagonal matrix with entries

$$D_k = \sum_{i=1}^n K_h(t_i - t_k^*) \omega(t_k^*) (\widetilde{DX}_{i,k})^2, \quad k = 1, \dots, m. \quad (\text{A.3})$$

The  $k$ -th row of the  $L$  matrix is

$$L_k = \sum_{i=1}^n K_h(t_i - t_k^*) \omega(t_k^*) \widetilde{DX}_{i,k} \widetilde{D\theta}_{i,k}, \quad (\text{A.4})$$

and

$$C = \sum_{k=1}^m \sum_{i=1}^n K_h(t_i - t_k^*) \omega(t_k^*) \widetilde{D\theta}_{i,k}^T \widetilde{D\theta}_{i,k}. \quad (\text{A.5})$$

**Lemma 1**  $D_k = n\omega(t_k^*)f(t_k^*) + o_p(n)$ ,

$$L_k = nh^2\mu_2(K)\omega(t_k^*)[f'(t_k^*)D_{\theta^T,k}^{(0)} + f(t_k^*)D_{X,k}^{(0)}D_{\theta^T,k}^{(0)}] + o_p(nh^2),$$

and  $C = nmh^2\mu_2(K)A_F + o_p(nmh^2)$ .

The definition of  $A_F$  is given under (2.7). **Proof of Lemma 1:** The proof of Lemma 1 comes from direct calculations using  $\widetilde{DX}_{i,k} = 1 + \sum_{j=1}^p (t_i - t_k^*)^j \frac{D_{X,k}^{(j-1)}}{j!}$ ,  $\widetilde{D\theta}_{i,k} = \sum_{j=1}^p (t_i - t_k^*)^j \frac{D_{\theta^T,k}^{(j-1)}}{j!}$  and (A.1). Notice that that  $\widetilde{DX}_{i,k}$  has  $p + 1$  terms that each is of the form of powers  $(t_i - t_k^*)^j$  multiplied by a bounded quantity. So  $D_k$  by (A.3) is sum of  $(p + 1)^2$  terms each of the form  $S_{k,j} = \sum_{i=1}^n K_h(t_i - t_k^*)(t_i - t_k^*)^j$  multiplied by a bounded quantity. Specifically,

$$D_k = [S_{k,0} + \sum_{j=1}^p S_{k,j} \left( \frac{D_{X,k}^{(j-1)}}{j!} \right) + \sum_{l=1}^p \left( \frac{D_{X,k}^{(l-1)}}{l!} \right) (S_{k,l} + \sum_{j=1}^p S_{k,l+j} \frac{D_{X,k}^{(j-1)}}{j!})] \omega(t_k^*).$$

Since the asymptotic is done for fixed  $p$ ,  $m \rightarrow \infty$  and  $n \rightarrow \infty$ , asymptotically  $D_k$  corresponds to the term with highest order among the  $(p + 1)^2$  terms. The leading term is  $S_{k,0}\omega(t_k^*) = n\omega(t_k^*)f(t_k^*) + o_p(n)$  by (A.1). The rest of terms are of order  $S_{k,j}$  for some  $j \geq 1$  so are of order  $O_p(nh^j)$  or  $O_p(nh^{j+1})$ . Either way, they are at most of order  $O_p(nh^2) = o_p(n)$ . Hence the sum  $D_k = n\omega(t_k^*)f(t_k^*) + o_p(n)$  is of order  $O_p(n)$ .

Similarly, looking at the leading terms in  $\widetilde{DX}_{i,k}\widetilde{D\theta}_{i,k}$  and  $\widetilde{D\theta}_{i,k}^T\widetilde{D\theta}_{i,k}$  gives the results for  $L_k$  and  $C$ . More detailed analysis can be found in the online supplemental materials.

This finishes the proof of Lemma 1.

It is easy to check by the block matrix algebra that

$$(J^T W J)^{-1} = \begin{pmatrix} D_{m \times m} & L_{m \times q} \\ L_{q \times m}^T & C_{q \times q} \end{pmatrix}^{-1} = \begin{pmatrix} D^{-1} + D^{-1}LV^{-1}L^TD^{-1} & -D^{-1}LV^{-1} \\ -V^{-1}L^TD^{-1} & V^{-1} \end{pmatrix} \quad (\text{A.6})$$

with  $V = C - L^TD^{-1}L$ . The order of quantities in (A.6) is described in the following lemma whose proof is provided in the online supplemental materials.

**Lemma 2**  $L^TD^{-1}L = O_p(mnh^4)$ ,  $V^{-1} = C^{-1}[1 + O_p(h^2)] = O_p(\frac{1}{nmh^2})$ ,  $D^{-1}LV^{-1} = O_p(\frac{1}{mn})$  and  $D^{-1}LV^{-1}L^TD^{-1} = O_p(\frac{h^2}{n})$ .

Using the results in Lemma 1 and 2,

$$(J^T W J)^{-1} = \begin{pmatrix} D_{m \times m}^{-1} + o_p(\frac{1}{n}) & O_p(\frac{1}{mn})_{m \times q} \\ O_p(\frac{1}{mn})_{q \times m} & C_{q \times q}^{-1} + o_p(\frac{1}{nmh^2}) \end{pmatrix}. \quad (\text{A.7})$$

Here and in the following, we write the matrix is of an order such as  $O((mn)^{-1})_{m \times q}$  when all its element is of the order.

**Remark:** For a  $d$ -dimensional  $X$ , the order analysis of the matrices would all remain the same. The  $D_{m \times m}$  matrix would become  $D_{md \times md}$  with diagonal block matrices  $D_k$  of size  $d \times d$ . And  $L_k$  would be matrices of size  $d \times q$ . As  $d$  is fixed, the multiplication of matrices with dimension  $d$  instead of 1 does not change the order. So the whole proof can be extended to  $d$ -dimensional  $X$  straightforwardly.

#### A.1. Asymptotic bias

The bias of  $\hat{\xi}$  given  $t_1, \dots, t_n, t_1^*, \dots, t_m^*, \xi^*$  is

$$\begin{aligned} \text{Bias}(\hat{\xi}) &= (J^T W J)^{-1} J^T W E(\tilde{Y}) - \xi_0 = (J^T W J)^{-1} J^T W \{E(Y - G(\xi^*) + J\xi^*) - J\xi_0\} \\ &= (J^T W J)^{-1} J^T W \{E(Y) - G(\xi^*) - J(\xi_0 - \xi^*)\} \end{aligned}$$

Denote  $J = (J_{1,1}^T, J_{2,1}^T, \dots, J_{n,1}^T, J_{1,2}^T, \dots, J_{n,m}^T)^T$ . Hence the elements in  $E(Y) - G(\xi^*) - J(\xi_0 - \xi^*)$  are those  $E(Y_i) - G_{i,k}(\xi^*) - J_{i,k}(\xi_0 - \xi^*)$ 's. With Taylor expansion of  $E(Y_i) = X(t_i)$  at time point  $t = t_k^*$ , we have

$$X(t_i) = X(t_k^*) + \sum_{j=1}^p \frac{(t_i - t_k^*)^j}{j!} X^{(j)}(t_k^*) + (t_i - t_k^*)^{p+1} \frac{X^{(p+1)}(\tilde{t}_{i,k})}{(p+1)!} = G_{i,k}(\xi_0) + (t_i - t_k^*)^{p+1} \frac{X^{(p+1)}(\tilde{t}_{i,k})}{(p+1)!},$$

where  $\tilde{t}_{i,k}$  is a point between  $t_k^*$  and  $t_i$ . Since  $G_{i,k}(\xi_0) - G_{i,k}(\xi^*) - J_{i,k}(\xi_0 - \xi^*) = O_p(|\xi_0 - \xi^*|^2) = O_p(n^{-2\delta})$ , we have

$$E(Y_i) - G_{i,k}(\xi^*) - J_{i,k}(\xi_0 - \xi^*) = (t_i - t_k^*)^{p+1} \frac{X^{(p+1)}(\tilde{t}_{i,k})}{(p+1)!} + O_p(n^{-2\delta}). \quad (\text{A.8})$$

Denote  $T_j = ((t_1 - t_1^*)^j, \dots, (t_n - t_1^*)^j, (t_1 - t_2^*)^j, \dots, (t_n - t_m^*)^j)^T$ . Similar to the analysis in the proof of Lemma 1, we evaluate the order of  $J^T W T_j$  by focusing on the term with the lowest power of  $(t_i - t_k^*)$  as the higher power terms lead to a smaller order kernel sum. The first  $m$  elements in  $J^T W T_j$  are of the form  $\sum_{i=1}^n K_h(t_i - t_k^*)(t_i - t_k^*)^j \omega(t_k^*) \widetilde{D}X_{i,k}$ ,  $k = 1, \dots, m$ . The lowest power term in  $\widetilde{D}X_{i,k}$  is 1 (i.e.,  $(t_i - t_k^*)^0$ ) so that those  $m$  elements are of the same order as  $S_{k,j} = \sum_{i=1}^n K_h(t_i - t_k^*)(t_i - t_k^*)^j$  which is  $O_p(nh^j)$  for  $p$  even, and  $O_p(nh^{j+1})$  for  $p$  odd by (A.1). The last  $q$  elements in  $J^T W T_j$  are  $\sum_{k=1}^m [\sum_{i=1}^n K_h(t_i - t_k^*)(t_i - t_k^*)^j \omega(t_k^*) \widetilde{D}\theta_{i,k}]$ . Again, the lowest power term in  $\widetilde{D}\theta_{i,k}$  is  $(t_i - t_k^*)$  so that those last  $q$  elements are of the same order as  $\sum_{k=1}^m [\sum_{i=1}^n K_h(t_i - t_k^*)(t_i - t_k^*)^{j+1}] = \sum_{k=1}^m S_{k,j+1}$ . That is, they are of order  $O_p(mnh^{j+2})$  for  $p$  even, and

$O_p(mnh^{j+1})$  for  $p$  odd by (A.1). In summation,

$$J^T W T_j = \begin{pmatrix} O_p(nh^j)_{m \times 1} \\ O_p(mnh^{j+2})_{q \times 1} \end{pmatrix} \quad \text{for } j \text{ even}; \quad \begin{pmatrix} O_p(nh^{j+1})_{m \times 1} \\ O_p(mnh^{j+1})_{q \times 1} \end{pmatrix} \quad \text{for } j \text{ odd}.$$

From (A.8),  $E(Y) - G(\xi^*) - J(\xi_0 - \xi^*) = T_{p+1}O_p(1) + T_0O_p(n^{-2\delta})$ . Plug-in the orders of  $J^T W T_{p+1}$  and  $J^T W T_0$ , we have that  $J^T W \{E(Y) - G(\xi^*) - J(\xi_0 - \xi^*)\}$  is

$$\begin{pmatrix} O_p(n(h^{p+1} + n^{-2\delta}))_{m \times 1} \\ O_p(mnh^2(h^{p+1} + n^{-2\delta}))_{q \times 1} \end{pmatrix} \quad \text{for } p \text{ odd}; \quad \begin{pmatrix} O_p(n(h^{p+2} + n^{-2\delta}))_{m \times 1} \\ O_p(mnh^2(h^p + n^{-2\delta}))_{q \times 1} \end{pmatrix} \quad \text{for } p \text{ even}.$$

Combining this with (A.6) and Lemma 2, the bias in estimating  $\theta$  is

$$\text{Bias}(\hat{\theta}) = O_p(n^{-2\delta}) + O_p(h^{p+1}) \quad p \text{ odd}, \quad \text{Bias}(\hat{\theta}) = O_p(n^{-2\delta}) + O_p(h^p) \quad p \text{ even}.$$

Since  $\delta > -1/4$ ,  $\text{Bias}(\hat{\theta}) = o_p(n^{-1/2})$  for a small  $h = o(n^{-1/2p})$ .

#### A.2. Asymptotic variance

For the variance of  $\hat{\xi}$  given  $t_1, \dots, t_n, t_1^*, \dots, t_m^*, \xi^*$ , notice that  $\text{var}(\hat{\xi}) = (J^T W J)^{-1} J^T W \text{var}(\tilde{Y}) W J (J^T W J)^{-1}$ . Denote  $\Sigma = \text{var}((Y_1, \dots, Y_n)^T) = \text{diag}\{\underbrace{\sigma^2, \dots, \sigma^2}_n\}$ . So  $\Sigma_Y = \text{var}(\tilde{Y}) = \text{var}(Y)$  are simply  $m \times m$  blocks of  $\Sigma$ . Thus,

$$J^T W \text{var}(\tilde{Y}) W J = \begin{pmatrix} D_{m \times m}^* & L_{m \times q}^* \\ (L^*)_{q \times m}^T & C_{q \times q}^* \end{pmatrix}, \quad (\text{A.9})$$

where the  $(k, j)$ -th element in  $D^*$  is

$$D_{k,j}^* = \sigma^2 \omega(t_k^*) \omega(t_j^*) \left[ \sum_{i=1}^n K_h(t_i - t_k^*) K_h(t_i - t_j^*) \widetilde{D} \widetilde{X}_{i,k} \widetilde{D} \widetilde{X}_{i,j} \right], \quad \text{for } k, j = 1, \dots, m, \quad (\text{A.10})$$

the  $k$ -th row in  $L^*$  is

$$L_k^* = \sigma^2 \omega(t_k^*) \left[ \sum_{j=1}^m \omega(t_j^*) \sum_{i=1}^n K_h(t_i - t_k^*) K_h(t_i - t_j^*) \widetilde{D} \widetilde{X}_{i,k} \widetilde{D} \theta_{i,j} \right], \quad \text{for } k = 1, \dots, m, \quad (\text{A.11})$$

and

$$C^* = \sum_{k=1}^m \sum_{j=1}^m \sigma^2 \omega(t_k^*) \omega(t_j^*) \left[ \sum_{i=1}^n K_h(t_i - t_k^*) K_h(t_i - t_j^*) \widetilde{D} \theta_{i,k}^T \widetilde{D} \theta_{i,j} \right]. \quad (\text{A.12})$$

**Lemma 3**

$$D_{k,k}^* = O_p(nh^{-1}), \quad D_{k,j}^* = o_p(n) \quad \text{for } k \neq j. \quad (\text{A.13})$$

$$L_k^* = nmh^2\sigma^2\mu_2(K)[\omega * f * \{\omega * f_g * F_{\theta T}\}'](t_k^*) + o_p(nmh^2). \quad (\text{A.14})$$

When  $\omega(a_0) \neq 0$  or  $\omega(b_0) \neq 0$ ,  $C^* = O_p(nmh + nm^2h^3)$ ; when  $\omega(a_0) = \omega(b_0) = 0$ ,  $C^* = O_p(nmh + nm^2h^4)$ . Particularly, when  $\omega(a_0) = \omega(b_0) = 0$  and  $mh^3 \rightarrow \infty$ ,

$$C^* = nm^2h^4\sigma^2[\mu_2(K)]^2B_F + o_p(nm^2h^4). \quad (\text{A.15})$$

The proof of Lemma 3 is given in the online supplemental materials. Using (A.6), we directly calculate  $\text{var}(\hat{\theta})$  as

$$V^{-1}L^TD^{-1}D^*D^{-1}LV^{-1} - V^{-1}(L^*)^TD^{-1}LV^{-1} - V^{-1}L^TD^{-1}L^*V^{-1} + V^{-1}C^*V^{-1}. \quad (\text{A.16})$$

Then the order can be calculated using the results in Lemmas 1, 2 and 3. For the case of  $\omega(a_0) = \omega(b_0) = 0$ , the first term in (A.16) is of smaller order  $O(\frac{1}{nmh})$  and ignored. When  $mh^3 \rightarrow \infty$ , we will see from the other three terms  $\text{var}(\hat{\theta}) = O_p(\frac{1}{n})$ . Using Lemma 3, the last term in (A.16) becomes  $V^{-1}C^*V^{-1} = \frac{\sigma^2}{n}A_F^{-1}B_FA_F^{-1} + o(\frac{1}{n})$ . Using Lemmas 1 and 2, the third term is  $-V^{-1}L^TD^{-1}L^*V^{-1} = -\frac{\sigma^2}{n}A_F^{-1}C_FA_F^{-1} + o_p(\frac{1}{n})$ . The second term in (A.16) is the transpose of the third term. Combining them together, we have

$$\text{var}(\hat{\theta}) = \frac{\sigma^2}{n}A_F^{-1}[B_F - (C_F + C_F^T)]A_F^{-1} + o_p(\frac{1}{n}).$$

For the second case of  $\omega(a_0) \neq 0$  or  $\omega(b_0) \neq 0$ , similar calculation using Lemma 3 shows that the variance of  $\hat{\theta}$  is of order  $O_p(\frac{1}{nmh^3} + \frac{1}{nh})$ .

More details are provided in the online supplemental materials.

Statistica Sinica

## Bibliography

- BRUNEL, N. J-B.(2008). Parameter estimation of ODEs via nonparametric estimators. *Electronic Journal of Statistics* **2**, 124-1267.
- CHEN, J. & WU, H. (2008). Efficient Local Estimation for Time-varying Coefficients in Deterministic Dynamic Models with Applications to HIV-1 Dynamics. *Journal of the American Statistical Association* **103**, 369-384.
- CHEN, T., HE, H. L. & CHURCH, G. M. (1999). Modeling gene expression with differential equations. *Pacific Symposium on Biocomputing*, **4**, 29-40.
- FAN, J. & GIJBELS, I. (1996). *Local Polynomial Modelling and Its Applications*. Chapman and Hall/CRC.
- FANG, Y., WU, H. & ZHU, L. (2011). A Two-Stage Estimation Method for Random Coefficient Differential Equation Models with Application to Longitudinal HIV Dynamic Data, *Statistica Sinica*, **21**, 1145-1170.
- FITZHUGH, R. (1961). Impulses and Physiological States in Models of Nerve Membrane. *Biophysical Journal*, **1**, 445C466.
- HOOKE, G., XIAO, L. & RAMSAY, J. O. (2010). CollocInfer: Collocation Inference for Dynamic Systems. R package version 0.1.2. <http://CRAN.R-project.org/package=CollocInfer>
- Huang, Y., Liu, D., and Wu, H. (2006). Hierarchical Bayesian methods for estimation of parameters in a longitudinal HIV dynamic system. *Biometrics*, **62**, 413-423.
- Huang, Y. and Wu, H. (2006). A Bayesian approach for estimating antiviral efficacy in HIV dynamic models. *Journal of Applied Statistics*, **33**, 155-174.

- Guedj, J., Thiébaud, R., and Commenges, D. (2007). Maximum likelihood estimation in dynamical models of HIV. *Biometrics*, **63**, 1198-1206.
- MAECHLER M. (2010). lokern: Kernel Regression Smoothing with Local or Global Plug-in Bandwidth. R package version 1.1-1. <http://CRAN.R-project.org/package=lokern>
- LEHMANN, E. L. & CASELLA, G. (1998). *Theory of Point Estimation*. Springer.
- Li, L., Brown, M. B., Lee, K. H., and Gupta, S. (2002). Estimation and inference for a spline-enhanced population pharmacokinetic model. *Biometrics* **58**, 601-611.
- LIANG, H. & WU, H. (2008). Parameter Estimation of Differential Equation Models Using a Framework of Measurement Error in Regression Models. *Journal of the American Statistical Association*, **103**, 1570-1583.
- LIANG, H. & WU, H. (2010). Correction on “Parameter Estimation of Differential Equation Models Using a Framework of Measurement Error in Regression Models (JASA, 103, 1570-1583)”. *Journal of the American Statistical Association*, **105**, 1636.
- LU, T., LIANG, H., LI, H. & WU, H. (2011). High Dimensional ODEs Coupled with Mixed-Effects Modeling Techniques for Dynamic Gene Regulatory Network Identification, *Journal of the American Statistical Association*, *Journal of the American Statistical Association*, **106**, 1242-1258.
- NAGUMO, J. S., ARIMOTO, S. & YOSHIZAWA, S. (1962). An Active Pulse Transmission Line Simulating a Nerve Axon. *Proceedings of the IRE*, **50**, 2061-2070.
- POYTON, A. A., VARZIRI, M.S., MCAULEY, K.B., MCLELLAN, P.J. & RAMSAY, J. O. (2006). Parameter estimation in continuous-time dynamic models using principal differential analysis. *Computers and Chemical Engineering*, **30**, 698-708.
- Putter, H., Heisterkamp, S. H., Lange, J. M., and De Wolf, F. (2002). A Bayesian approach to parameter estimation in HIV dynamical models. *Stat. Med.*, **21**, 2199-2214.
- RAMSAY, J. O., HOOKER, G., CAMPBELL, D. & CAO, J. (2007). Parameter estimation for differential equations: a generalized smoothing approach (with discussion). *J. R. Statist. Soc. B*, **69**, 741-796.

WU, H., KUMAR, A., MIAO, H., HOLDEN-WILTSE, J., MOSMANN, T.R., LIVINGSTONE, A. M., BELZ, G.T., PERELSON, A. S., ZAND, M. & TOPHAM, D. J. (2011) Modeling of Influenza-Specific CD8+ T Cells During the Primary Response Indicates that the Spleen Is a Major Source of Effectors. *Journal of Immunology*. **187**, 4474-4482.

Statistica Sinica