

Efficient Estimation in Expectile Regression Using Envelope Models

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Abstract: As a generalization of the classical linear regression, expectile regression (ER) explores the relationship between the conditional expectile of a response variable and a set of predictor variables. ER with respect to different expectile levels can provide a comprehensive picture of the conditional distribution of the response variable given the predictors. We adopt an efficient estimation method called the envelope model ([7]) in ER, and construct a novel envelope expectile regression (EER) model. Estimation of the EER parameters can be performed using the generalized method of moments (GMM). We establish the consistency and derive the asymptotic distribution of the EER estimators. In addition, we show that the EER estimators are asymptotically more efficient than the ER estimators. Numerical experiments and real data examples are provided to demonstrate the efficiency gains attained by EER compared to ER, and the efficiency gains can further lead to improvements in prediction.

Keywords and phrases: Sufficient dimension reduction, envelope model, expectile regression, generalized method of moments.

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1. Introduction

The classical linear regression is a commonly used method when we want to study the relationship between a response variable Y and a p -dimensional predictor vector \mathbf{X} . It depicts the linear dependence between the conditional mean of Y and \mathbf{X} . However, it assumes that the conditional distribution of Y given \mathbf{X} is homoscedastic, which is not always satisfied in real datasets. While the condition mean gives important information on the conditional distribution of Y and \mathbf{X} , it does not provide a complete picture of the distribution, especially for skewed distributions arising from price or income data. As a result, quantile regression (QR), which can overcome the limitations, gained considerable interest in recent years. QR studies the conditional quantile of Y given \mathbf{X} . It is a distribution free method and does not impose any assumption on the conditional distribution of the response Y . Investigation of multiple quantile levels gives us a comprehensive description of the distribution of Y conditional on \mathbf{X} .

An alternative to QR is expectile regression (ER). The idea of expectile was firstly studied in [31]. The π th expectile of Y , denoted by $f_\pi(Y)$, is defined as

$$f_\pi(Y) = \arg \min_{f \in \mathbb{R}} \mathbb{E}[\phi_\pi(Y - f)], \quad \pi \in (0, 1),$$

where $\phi_\pi(z) = |\pi - I(z < 0)|z^2$ is called the asymmetric least squares loss function. When $\pi = 0.5$, $\phi_\pi(z)$ is the least squares loss and $f_\pi(Y) = \mathbb{E}[Y]$. ER studies the conditional expectile of the response Y given the predictors \mathbf{X} . It has been explored in many statistics and econometric literature. [52] proposed a local linear polynomial estimator of the conditional expectiles with a one-dimensional predictor and established the corresponding asymptotic properties. [24] developed the conditional autoregressive expectile models (CARE), which investigated the impact of previous asset returns on the conditional expectile-based Value at Risk (EVaR) of current asset returns, and allowed different effects from positive and negative returns. They established the asymptotic normality of the CARE estimators, and extended the results in [31] to stationary and weakly dependent data. [49] proposed a varying-coefficient expectile model to estimate the conditional EVaR of asset returns. This approach allows the coefficients to change with an effect modifying risk factor and provides a more flexible way to model the data. They showed that the varying-coefficient expectile model yields more stable estimators and more accurate prediction intervals compared to the CARE models. In recent years, many advances have been taken

place on model selection in expectile regression. [18] studied the sparse expectile regression under high dimensional settings where the penalty functions include the Lasso and nonconvex penalties. [42] introduced a variety of model selection methods in semiparametric expectile regression. [27] provided asymptotic distributions of penalized expectile regression with SCAD and adaptive LASSO penalties for both *i.i.d.* and non-*i.i.d.* random errors. In addition, semiparametric ([36, 40, 39]) and nonparametric ([51, 50, 15, 21]) expectile estimation methods have been proposed in literature.

There are a lot of connections between QR and ER. Similar as QR, ER does not impose any distributional assumption on the response Y as well, and it can provide us with complete information on the conditional distribution of Y . Therefore, both QR and ER are able to overcome the previously mentioned limitation of the classical linear regression. Moreover, [52] pointed out there exists a one-to-one mapping between expectiles and quantiles. Hence ER can be interpreted as a flexible QR. In addition, expectiles can be used to estimate a quantile-based risk measure called expected shortfall [45]. However, at the same time, we should not ignore some different properties between QR and ER. Recall that the α th quantile of the response Y , denoted by $q_\alpha(Y)$, is defined as $q_\alpha(Y) = \arg \min_{q \in \mathbb{R}} E\{\tau_\alpha(Y - q)\}$, where $\tau_\alpha(z) = |\alpha - I(z < 0)||z|$ is the check loss function. Therefore quantiles minimize the expected check loss τ_α while expectiles minimize the expected asymmetric least squares loss ϕ_π . The check loss function is not differentiable while the asymmetric least squares loss function is differentiable because of the quadratic term. The difference in the loss functions give QR and ER their respective advantages. The main advantage of QR is that its results are easier to be interpreted and it is more robust to outliers than ER; while the main advantage of ER over QR is that ER is more computationally friendly, especially in a semiparametric model, as pointed out by [40]. Moreover, ER is more sensitive to the extreme values in datasets because ER takes the distance between an observation and the estimated expectile into account while QR only considers whether an observation is greater or less than the estimated quantile. This characteristic makes ER more desirable in many applications. One example is on the risk measures in the fields of econometrics and finance. Value at Risk (VaR) is a popular measure for evaluating portfolio risk based on quantiles. It provides crucial information about the potential loss, however, it is insensitive to the severity of more extreme realization since it does not depend on the tail shape of the distribution ([10, 14]). [24] proposed the risk measure EVaR and indicated EVaR can reflect the magnitude of the extreme losses for the underlying risk. Therefore, given a set of risk factors, studying conditional EVaR rather than conditional VaR may lead to a more proper response to a catastrophic loss. At last, for the special case with $\alpha = 0.5$, QR degenerates to the conditional median regression. For the special case with $\pi = 0.5$, ER degenerates to the standard linear regression.

In this paper we propose a new ER method, the *envelope expectile regression* (EER), for efficient estimation of the parameters in ER. It is motivated by the fact that some linear combinations of the predictors may be irrelevant to the

distribution of the response. Our method takes this data structure into account, which results in more efficient estimation compared with the standard ER. We call those irrelevant combinations *the immaterial part* and the remaining part *the material part*. To identify the material part and the immaterial part, we employ a nascent technique called the *envelope* method. The immaterial part is then excluded from the subsequent analysis, leading to efficiency gains in estimation. To be noted, the immaterial part is different from the subset of inactive predictors in the popular penalized variable selection methods. We can see that both the penalized variable selection methods and the envelope method reduce the number of free parameters in the model but they are based on different assumptions. The penalized variable selection methods assume a subset of individual predictor variables are irrelevant to regression and thus have coefficients zero; while the envelope method assumes some linear combinations of the predictors are irrelevant to regression but all predictors can have nonzero coefficients. There are different application scenarios for the penalized variable selection methods and the envelope method. For example, suppose that a biologist wants to analyze the relationship between genes and a certain disease, given a dataset with measurements on the severity of the disease (response) and the expression levels of different genes (predictors). If the biologist believes only a few genes in the dataset have effects on the disease, then he would apply the penalized variable selection methods to identify those genes. On the other hand, if the biologist feels like each gene in the dataset is related to the disease in some way, then he would use dimension reduction methods such as principal component regression, partial least squares or the envelope model to find the linear combination(s) of the genes that affects the disease. The envelope method was first proposed in ([7]) under the context of multivariate linear regression for efficient estimation. It is then applied to many contexts in multivariate analysis including partial least squares ([5]), generalized linear models ([8]), elliptical linear regression ([16]), reduced rank regression ([6]), variable selection ([44]), bayesian analysis ([23]), matrix and tensor valued response ([11, 26]), and quantile regression ([12]). In this paper, we apply the idea of envelope method to expectile regression. The parameters are estimated by the generalized method of moments (GMM; [20]). And we demonstrate that the EER estimators have smaller asymptotic variance than the ER estimators both theoretically and numerically.

This paper is organized as follows. We derive the EER model in Section 2. Estimation of the EER model is discussed in Section 3. In Section 4, we investigate the theoretical properties of the EER estimators, and prove that the EER estimators are asymptotically more efficient than the ER estimators. We use simulations to demonstrate the performance of the EER model in estimation and prediction in Section 5. State murder rate data and S&P 500 index data are analyzed in Section 6 as examples. The extension to semiparametric settings is discussed in Section 7. Section 8 concludes the paper with a discussion. All technical proofs are deferred to the supplementary material.

The following notations will be used in our discussion. Let $\mathbb{R}^{p \times u}$ be the set of all $p \times u$ matrices. For any matrix $\mathbf{A} \in \mathbb{R}^{p \times u}$, $\text{span}(\mathbf{A})$ represents the subspace

spanned by the columns of \mathbf{A} . Let $\mathbf{P}_{\mathbf{A}}$ be the projection matrix onto $\text{span}(\mathbf{A})$ and $\mathbf{Q}_{\mathbf{A}} = \mathbf{I}_p - \mathbf{P}_{\mathbf{A}}$ be the projection matrix onto its orthogonal complement $\text{span}(\mathbf{A})^\perp$. For any subspace \mathcal{S} in \mathbb{R}^p , $\mathbf{P}_{\mathcal{S}}$ represents the projection matrix onto \mathcal{S} and $\mathbf{Q}_{\mathcal{S}}$ represents the projection matrix onto \mathcal{S}^\perp . Let “vec” represent the vectorization operator that vectorizes a matrix columnwise and “vech” represent the half-vectorization operator that vectorizes the lower triangle of a symmetric matrix. We use $\|\cdot\|$ to represent Frobenius norm and \dagger to denote the Moore-Penrose inverse. We write \xrightarrow{d} for convergence in distribution and \xrightarrow{p} for convergence in probability. Moreover, a subspace \mathcal{R} in \mathbb{R}^p is a reducing subspace of a matrix $\mathbf{M} \in \mathbb{R}^{p \times p}$ if and only if \mathbf{M} can be decomposed as $\mathbf{M} = \mathbf{P}_{\mathcal{R}}\mathbf{M}\mathbf{P}_{\mathcal{R}} + \mathbf{Q}_{\mathcal{R}}\mathbf{M}\mathbf{Q}_{\mathcal{R}}$.

2. Envelope Expectile Regression

2.1. Expectile Regression

Consider a univariate response Y and a p -dimensional predictor vector \mathbf{X} . The standard ER considers the π th conditional expectile of Y as a linear function of \mathbf{X}

$$f_\pi(Y|\mathbf{X}) = \mu_\pi + \boldsymbol{\beta}_\pi^T \mathbf{X}, \quad (2.1)$$

where $f_\pi(Y|\mathbf{X})$ represents the π th conditional expectile of Y given \mathbf{X} , μ_π is the intercept and $\boldsymbol{\beta}_\pi \in \mathbb{R}^p$ contains the coefficients. When $\pi = 0.5$, $f_{0.5}(Y|\mathbf{X})$ degenerates to the conditional mean $E(Y|\mathbf{X})$ and ER degenerates to the standard linear regression.

To get the ER estimators, we use a property of the conditional expectile discussed in [31] that

$$(\mu_\pi, \boldsymbol{\beta}_\pi) = \arg \min_{\mu \in \mathbb{R}, \boldsymbol{\beta} \in \mathbb{R}^p} E\{\phi_\pi(Y - \mu - \boldsymbol{\beta}^T \mathbf{X})|\mathbf{X}\},$$

where $\phi_\pi(z) = |\pi - I(z < 0)|z^2$ is an asymmetric squared loss function. Given the random samples $\{(Y_i, \mathbf{X}_i) : i = 1, \dots, n\}$ of (Y, \mathbf{X}) , the ER estimators $\hat{\mu}_\pi$ and $\hat{\boldsymbol{\beta}}_\pi$ can be obtained by solving

$$(\hat{\mu}_\pi, \hat{\boldsymbol{\beta}}_\pi) = \arg \min_{\mu_\pi \in \mathbb{R}, \boldsymbol{\beta}_\pi \in \mathbb{R}^p} \frac{1}{n} \sum_{i=1}^n \phi_\pi(Y_i - \mu_\pi - \boldsymbol{\beta}_\pi^T \mathbf{X}_i).$$

Taking the first derivative with respect to $(\mu_\pi, \boldsymbol{\beta}_\pi^T)^T$, the minimizer should satisfy the following estimating equations

$$\frac{1}{n} \sum_{i=1}^n \mathbf{W}_i (Y_i - \mu_\pi - \boldsymbol{\beta}_\pi^T \mathbf{X}_i) |I(Y_i < \mu_\pi + \mathbf{X}_i^T \boldsymbol{\beta}_\pi) - \pi| = 0, \quad (2.2)$$

where $\mathbf{W}_i = (1, \mathbf{X}_i^T)^T$.

2.2. Envelope Expectile Regression

EER is derived from the motivation that certain linear combinations of the predictors are irrelevant to some conditional expectiles, e.g. $E(Y|\mathbf{X})$, of the response. For example, a stock index may be related to only a few combinations of the economic factors, while these combinations are uncorrelated with the other combinations that are not responsible for the variation in the index. Following this observation, we suppose that there exists a subspace \mathcal{S}_π in the full predictor space \mathbb{R}^p such that the π th conditional expectile of the response Y is related to the predictor vector \mathbf{X} only through $\mathbf{P}_{\mathcal{S}_\pi}\mathbf{X}$. Specifically, we assume that

$$(a) \quad f_\pi(Y|\mathbf{X}) = f_\pi(Y|\mathbf{P}_{\mathcal{S}_\pi}\mathbf{X}) \quad \text{and} \quad (b) \quad \text{Cov}(\mathbf{Q}_{\mathcal{S}_\pi}\mathbf{X}, \mathbf{P}_{\mathcal{S}_\pi}\mathbf{X}) = 0. \quad (2.3)$$

The conditions in (2.3) imply that $\mathbf{Q}_{\mathcal{S}_\pi}\mathbf{X}$ does not provide any information to the π th conditional expectile of Y neither from itself nor from its association with $\mathbf{P}_{\mathcal{S}_\pi}\mathbf{X}$. We call $\mathbf{P}_{\mathcal{S}_\pi}\mathbf{X}$ the material part of \mathbf{X} and $\mathbf{Q}_{\mathcal{S}_\pi}\mathbf{X}$ the immaterial part of \mathbf{X} .

Remark 1. *Conditions in (2.3) incorporate the idea of sufficient dimension reduction in expectile regression. When $\pi = 0.5$ and the conditions in (2.3) are superimposed, model (2.1) is the envelope-based partial least squares (PLS) regression of Y on \mathbf{X} ([5]). It is well known that PLS regression improves prediction performance over ordinary least squares regression (e.g., [1, 4, 22]). For general $\pi \in (0, 1)$, conditions in (2.3) establish a general asymmetric envelope-based PLS regression framework, which improves estimation efficiency and enhances prediction performance over ER both theoretically and numerically as evidenced in Sections 4-6.*

Under the parameterization of ER (2.1), condition (2.3a) is equivalent to (i) $\beta_\pi \in \mathcal{S}_\pi$, and condition (2.3b) holds if and only if (ii) \mathcal{S}_π is a reducing subspace of $\Sigma_{\mathbf{X}}$ ([7]), where $\Sigma_{\mathbf{X}}$ is the covariance matrix of \mathbf{X} . If we find such a subspace \mathcal{S}_π , we can identify the immaterial part $\mathbf{Q}_{\mathcal{S}_\pi}\mathbf{X}$ when evaluating the relationship between $f_\pi(Y|\mathbf{X})$ and \mathbf{X} . Then estimation efficiency gains can be achieved by accounting for the immaterial variation in subsequent analysis. There may exist more than one subspace satisfying (i) and (ii). For instance, the full space \mathbb{R}^p always satisfies (i) and (ii). To achieve the most efficiency gains, we focus on the smallest subspace (i.e., the subspace with smallest dimension) that satisfies (i) and (ii), such that we can identify all the immaterial information. We call this subspace the $\Sigma_{\mathbf{X}}$ -envelope of β_π , and denote it by $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$ or \mathcal{E}_π if it appears in subscripts. The dimension of the envelope subspace $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$ is denoted by u_π ($0 \leq u_\pi \leq p$). [7] discussed the existence and uniqueness of the envelope subspace $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$.

Before deriving the EER model, we first discuss the parameterization of the envelope subspace $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$. The envelope subspace can be determined by its basis. However, there can be many bases of $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$. To make the parameters identifiable, we define one representative basis Γ_π for each envelope subspace: Take an arbitrary basis \mathbf{G}_π of the envelope subspace $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$. Since the dimension of the envelope subspace $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$ is u_π , \mathbf{G}_π has rank u_π and we can

find u_π rows in \mathbf{G}_π that constitute a nonsingular matrix. If there are multiple combinations of such u_π rows, we take the rows with smallest indices. Without loss of generality, we assume that the first u_π rows of \mathbf{G}_π constitute a nonsingular matrix, and we write \mathbf{G}_π as $(\mathbf{G}_{1\pi}^T, \mathbf{G}_{2\pi}^T)^T$, where $\mathbf{G}_{1\pi} \in \mathbb{R}^{u_\pi \times u_\pi}$ and $\mathbf{G}_{2\pi} \in \mathbb{R}^{(p-u_\pi) \times u_\pi}$. Define the representative basis $\mathbf{\Gamma}_\pi = \mathbf{G}_\pi \mathbf{G}_{1\pi}^{-1} \equiv (\mathbf{I}_{u_\pi}, \mathbf{A}^T)^T$, where $\mathbf{A} \in \mathbb{R}^{(p-u_\pi) \times u_\pi}$. It can be shown that the representing basis defined as above is unique for $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$. Therefore we have established a one-to-one correspondence between \mathbf{A} and the envelope subspace $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$. And if we know \mathbf{A} , we can completely decide $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$. The following lemma shows that the basis of the orthogonal complement of $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$ can also be expressed as a function of \mathbf{A} , which is useful for establishing the EER model. The proof is given in Section 1.1 of the supplementary material.

Lemma 1. *If $\mathbf{\Gamma}_\pi = (\mathbf{I}_{u_\pi}, \mathbf{A}^T)^T$ is a basis of $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$, then $\mathbf{\Gamma}_{0\pi} = (-\mathbf{A}, \mathbf{I}_{p-u_\pi})^T$ is a basis of $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)^\perp$.*

Using the representative bases $\mathbf{\Gamma}_\pi$ and $\mathbf{\Gamma}_{0\pi}$, we reparametrize β_π and $\Sigma_{\mathbf{X}}$ to derive the EER model. Since $\beta_\pi \in \mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$, there exists a u_π -dimensional coordinate vector η_π such that $\beta_\pi = \mathbf{\Gamma}_\pi \eta_\pi$. Because $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$ is a reducing subspace of $\Sigma_{\mathbf{X}}$, $\Sigma_{\mathbf{X}}$ can be decomposed as $\Sigma_{\mathbf{X}} = \mathbf{P}_{\mathcal{E}_\pi} \Sigma_{\mathbf{X}} \mathbf{P}_{\mathcal{E}_\pi} + \mathbf{Q}_{\mathcal{E}_\pi} \Sigma_{\mathbf{X}} \mathbf{Q}_{\mathcal{E}_\pi}$, where $\mathbf{P}_{\mathcal{E}_\pi} \Sigma_{\mathbf{X}} \mathbf{P}_{\mathcal{E}_\pi}$ is the variance of the material part $\mathbf{P}_{\mathcal{E}_\pi} \mathbf{X}$ and $\mathbf{Q}_{\mathcal{E}_\pi} \Sigma_{\mathbf{X}} \mathbf{Q}_{\mathcal{E}_\pi}$ is the variance of the immaterial part $\mathbf{Q}_{\mathcal{E}_\pi} \mathbf{X}$. With $\mathbf{\Gamma}_\pi$ and $\mathbf{\Gamma}_{0\pi}$ defined in Lemma 1, the coordinate form of the EER model is as follows

$$\begin{aligned} f_\pi(Y|\mathbf{X}) &= \mu_\pi + \eta_\pi^T \mathbf{\Gamma}_\pi^T \mathbf{X} \\ \Sigma_{\mathbf{X}} &= \mathbf{\Gamma}_\pi \mathbf{\Omega}_\pi \mathbf{\Gamma}_\pi^T + \mathbf{\Gamma}_{0\pi} \mathbf{\Omega}_{0\pi} \mathbf{\Gamma}_{0\pi}^T. \end{aligned} \quad (2.4)$$

In the EER model, we can see the predictor vector \mathbf{X} affects the conditional expectile $f_\pi(Y|\mathbf{X})$ only through its linear combinations $\mathbf{\Gamma}_\pi^T \mathbf{X}$. The number of the linear combinations is u_π because $\mathbf{\Gamma}_\pi$ has u_π columns. Therefore, u_π represents the number of relevant linear combinations of the predictors, rather than the number of active predictors in penalized models. The u_π -dimensional vector η_π carries the coordinates of β_π relative to $\mathbf{\Gamma}_\pi$. The positive definite matrix $\mathbf{\Omega}_\pi \in \mathbb{R}^{u_\pi \times u_\pi}$ carries the coordinates of $\Sigma_{\mathbf{X}}$ relative to $\mathbf{\Gamma}_\pi$ and the positive definite matrix $\mathbf{\Omega}_{0\pi} \in \mathbb{R}^{(p-u_\pi) \times (p-u_\pi)}$ carries the coordinates of $\Sigma_{\mathbf{X}}$ relative to $\mathbf{\Gamma}_{0\pi}$. The parameter vector in the EER model is then $\zeta = (\mu_\pi, \eta_\pi^T, \text{vec}(\mathbf{A})^T, \text{vech}(\mathbf{\Omega}_\pi)^T, \text{vech}(\mathbf{\Omega}_{0\pi})^T, \boldsymbol{\mu}_{\mathbf{X}}^T)^T$, where $\boldsymbol{\mu}_{\mathbf{X}}$ is the mean of \mathbf{X} . Here, we include $\boldsymbol{\mu}_{\mathbf{X}}$ in the parameter vector since it is involved in the estimating equations in Section 3. The total number of parameters is $1 + u_\pi + p + p(p+1)/2$. The parameter count is as follows: μ_π has 1 parameter, η_π has u_π parameters, \mathbf{A} has $u_\pi(p-u_\pi)$ parameters, $\mathbf{\Omega}_\pi$ and $\mathbf{\Omega}_{0\pi}$ are both symmetric matrices, and they have $u_\pi(u_\pi+1)/2$ and $(p-u_\pi)(p-u_\pi+1)/2$ parameters respectively, and $\boldsymbol{\mu}_{\mathbf{X}}$ has p parameters. When $u_\pi = p$, the EER model degenerates to the ER model (2.1). The parameters in ER are μ_π and β_π . In this paper, we also consider $\Sigma_{\mathbf{X}}$ and $\boldsymbol{\mu}_{\mathbf{X}}$ as parameters in ER to make it comparable with the EER model when asymptotic covariance of the estimators is concerned. The parameter vector in ER is $\theta = (\mu_\pi, \beta_\pi^T, \text{vech}(\Sigma_{\mathbf{X}})^T, \boldsymbol{\mu}_{\mathbf{X}}^T)^T$, and the total number of parameters is 1

+ $2p + p(p + 1)/2$. Comparing the number of parameters, we can see that the EER model reduces the number of parameters by $p - u_\pi$.

3. Estimation

In this section, we derive the EER estimators. Since there is no distributional assumption on the predictors or the response, the maximum likelihood estimation is not applicable for the EER model. Instead we adopt the generalized method of moments (GMM; [20]) to obtain the EER estimators.

We first present the estimating equations under ER, and then reparameterize the equations for the estimation of the EER model. The estimating equations under ER are constructed from (2.2) and the moment conditions of \mathbf{X} :

$$e_n(\boldsymbol{\theta}) = \begin{pmatrix} \frac{1}{n} \sum_{i=1}^n \mathbf{W}_i(Y_i - \mu_\pi - \mathbf{X}_i^T \boldsymbol{\beta}_\pi) |I(Y_i < \mu_\pi + \mathbf{X}_i^T \boldsymbol{\beta}_\pi) - \pi| \\ \text{vech}(\boldsymbol{\Sigma}_{\mathbf{X}}) - \text{vech}(\mathbf{S}_{\mathbf{X}}) \\ \boldsymbol{\mu}_{\mathbf{X}} - \bar{\mathbf{X}} \end{pmatrix} = 0, \quad (3.1)$$

where $\bar{\mathbf{X}} = \sum_{i=1}^n \mathbf{X}_i/n$ is the sample mean and $\mathbf{S}_{\mathbf{X}} = \sum_{i=1}^n (\mathbf{X}_i - \boldsymbol{\mu}_{\mathbf{X}})(\mathbf{X}_i - \boldsymbol{\mu}_{\mathbf{X}})^T/n$ is the sample covariance matrix given $\boldsymbol{\mu}_{\mathbf{X}}$. Let $\boldsymbol{\theta}$ denote the ER estimator by solving (3.1).

Under the EER model (2.4), we have $\boldsymbol{\beta}_\pi = \boldsymbol{\Gamma}_\pi \boldsymbol{\eta}_\pi$ and $\boldsymbol{\Sigma}_{\mathbf{X}} = \boldsymbol{\Gamma}_\pi \boldsymbol{\Omega}_\pi \boldsymbol{\Gamma}_\pi^T + \boldsymbol{\Gamma}_{0\pi} \boldsymbol{\Omega}_{0\pi} \boldsymbol{\Gamma}_{0\pi}^T$. Then we can build a map ψ between the EER parameter vector $\boldsymbol{\zeta}$ and the ER parameter vector $\boldsymbol{\theta}$, i.e., $\psi(\boldsymbol{\zeta}) = \boldsymbol{\theta}$. Then we can reparameterize (3.1) to get the estimating equations for the EER model:

$$e_n^*(\boldsymbol{\zeta}) = \begin{pmatrix} \frac{1}{n} \sum_{i=1}^n \mathbf{W}_i(Y_i - \mu_\pi - \mathbf{X}_i^T \boldsymbol{\Gamma}_\pi \boldsymbol{\eta}_\pi) |I(Y_i < \mu_\pi + \mathbf{X}_i^T \boldsymbol{\Gamma}_\pi \boldsymbol{\eta}_\pi) - \pi| \\ \text{vech}(\boldsymbol{\Gamma}_\pi \boldsymbol{\Omega}_\pi \boldsymbol{\Gamma}_\pi^T + \boldsymbol{\Gamma}_{0\pi} \boldsymbol{\Omega}_{0\pi} \boldsymbol{\Gamma}_{0\pi}^T) - \text{vech}(\mathbf{S}_{\mathbf{X}}) \\ \boldsymbol{\mu}_{\mathbf{X}} - \bar{\mathbf{X}} \end{pmatrix} = 0. \quad (3.2)$$

Note that in the EER model (2.4), other than $\boldsymbol{\zeta} = (\mu_\pi, \boldsymbol{\eta}_\pi^T, \text{vec}(\mathbf{A})^T, \text{vech}(\boldsymbol{\Omega}_\pi)^T, \text{vech}(\boldsymbol{\Omega}_{0\pi})^T, \boldsymbol{\mu}_{\mathbf{X}}^T)^T$, the dimension of the envelope subspace u_π is also an important parameter. Here we first discuss the estimation of $\boldsymbol{\zeta}$ assuming u_π is known. In the estimating equation (3.2), there are $1 + 2p + p(p + 1)/2$ equations and $1 + u_\pi + p + p(p + 1)/2$ unknown parameters. As a result, it is possible that no solution exists. We then apply the GMM approach ([20]) to obtain the EER estimator $\hat{\boldsymbol{\zeta}}$. Let $\mathbf{Z} = (\mathbf{X}^T, Y)^T$ and $\mathbf{W} = (1, \mathbf{X}^T)^T$. Define $s(\mathbf{Z}; \boldsymbol{\theta})$ to be the population version of the moment conditions in (3.1):

$$s(\mathbf{Z}; \boldsymbol{\theta}) = \begin{pmatrix} s_1(\mathbf{Z}; \boldsymbol{\theta}) \\ s_2(\mathbf{Z}; \boldsymbol{\theta}) \\ s_3(\mathbf{Z}; \boldsymbol{\theta}) \end{pmatrix} = \begin{pmatrix} \mathbf{W}(Y - \mu_\pi - \mathbf{X}^T \boldsymbol{\beta}_\pi) |I(Y < \mu_\pi + \mathbf{X}^T \boldsymbol{\beta}_\pi) - \pi| \\ \text{vech}(\boldsymbol{\Sigma}_{\mathbf{X}}) - \text{vech}\{(\mathbf{X} - \boldsymbol{\mu}_{\mathbf{X}})(\mathbf{X} - \boldsymbol{\mu}_{\mathbf{X}})^T\} \\ \boldsymbol{\mu}_{\mathbf{X}} - \mathbf{X} \end{pmatrix}. \quad (3.3)$$

The estimation procedure can be summarized in the following steps.

Step 1 : Get the intermediate estimator $\hat{\boldsymbol{\zeta}}^*$ by minimizing $e_n^*(\boldsymbol{\zeta})^T e_n^*(\boldsymbol{\zeta})$.

Step 2 : Compute the scale matrix

$$\hat{\Delta} = \left[\frac{1}{n} \sum_{i=1}^n s(\mathbf{Z}_i; \psi(\hat{\zeta}^*)) s(\mathbf{Z}_i; \psi(\hat{\zeta}^*))^T \right]^{-1}.$$

Step 3 : Obtain the GMM estimator $\hat{\zeta}$ by minimizing $e_n^*(\zeta)^T \hat{\Delta} e_n^*(\zeta)$.

Once we obtain $\hat{\zeta}$, the EER estimators of β_π and $\Sigma_{\mathbf{X}}$ are $\hat{\beta}_\pi = \hat{\Gamma}_\pi \hat{\eta}_\pi$ and $\hat{\Sigma}_{\mathbf{X}} = \hat{\Gamma}_\pi \hat{\Omega}_\pi \hat{\Gamma}_\pi^T + \hat{\Gamma}_{0\pi} \hat{\Omega}_{0\pi} \hat{\Gamma}_{0\pi}^T$. And we use $\hat{\theta}$ to denote the EER estimator of θ : $\hat{\theta} = (\hat{\mu}_\pi, \hat{\beta}_\pi^T, \text{vech}(\hat{\Sigma}_{\mathbf{X}})^T, \hat{\mu}_{\mathbf{X}}^T)^T$. An analysis of the computational burden of the estimation procedure is given in Section 8 of the Supplement.

Remark 2. In some envelope literature, e.g., [7], a different parameterization is adopted for Γ_π : Γ_π is required to be a semi-orthogonal matrix, i.e., $\Gamma_\pi^T \Gamma_\pi = \mathbf{I}_{u_\pi}$. In this case, $\text{span}(\Gamma_\pi)$ is a point on a $p \times u_\pi$ Grassmann manifold, where a $p \times u_\pi$ Grassmann manifold is the set of all u_π -dimensional subspaces in a p -dimensional space. The above procedure can still be used to estimate the parameters, except that Step 1 and Step 3 involve Grassmann manifold optimization, which could be complicated and slow in sizable problems. For more information about Grassmann manifold optimization algorithms, please refer to [13, 28].

Now we discuss the selection of u_π . Similar to other dimension reduction based methods (such as principal component analysis, partial least squares or reduced rank regression), the model selection for the EER model (2.4) is essentially the selection of the dimension u_π . In existing envelope models and methods, the dimension u_π is usually chosen by AIC, BIC or log-likelihood ratio testing. Since AIC, BIC as well as log-likelihood ratio testing all requires a likelihood function, they are not applicable in the context of the EER model. As a result, we adopt a nonparameteric method, robust cross validation (RCV; [32]), for the selection of u_π . Following [17, page 244], we use the ‘‘one-standard error’’ rule with RCV to choose the most parsimonious model which has about the same predictive accuracy as the best model. RCV is performed in the following steps:

Step 1 : Randomly split the data into K folds. Usually K takes the value of 5 or 10. Successively use the k th fold for testing and the remaining folds for training, $k = 1, \dots, K$.

Step 2 : For each possible u_π ($0 \leq u_\pi \leq p$), compute the mean expectile loss $\text{RCV}(u_\pi) = \frac{1}{n} \sum_{i=1}^n \phi_\pi(Y_i - \hat{\mu}_{\pi, -k(i)} - \hat{\beta}_{\pi, -k(i)}^T \mathbf{X}_i)$, where $\hat{\mu}_{\pi, -k(i)}$ and $\hat{\beta}_{\pi, -k(i)}$ are the EER estimators using the data excluding the k th fold that contains the i th observation.

Step 3 : Instead of choosing the \tilde{u}_π which achieves the smallest mean expectile loss $\text{RCV}(\tilde{u}_\pi)$, we select the smallest \hat{u}_π whose mean expectile loss is less than one standard error above $\text{RCV}(\tilde{u}_\pi)$.

We provide an implementation of the EER model in the R package `expectEnv` which is available at <https://github.com/chentuo1993/expectEnv>. Using the

GMM approach, this package provides the EER estimator for parameters ζ and θ . It also implements RCV for the selection of the envelope dimension u_π and computes bootstrap standard deviations for the EER estimators.

4. Theoretical Results

In this section, we prove the EER estimator $\hat{\theta}$ is asymptotically more efficient than or as efficient as the ER estimator $\tilde{\theta}$. We first derive the asymptotic distribution of $\tilde{\theta}$. [31] proved that the ER estimator $\tilde{\beta}_\pi$ is consistent and asymptotically normal under some regularity conditions. We extend this result from $\tilde{\beta}_\pi$ to $\tilde{\theta}$ in Theorem 1. Next we establish the asymptotic distribution of $\hat{\theta}$ in Theorem 2, and show that the EER estimator is at least as efficient as the ER estimator by comparing the asymptotic covariance matrices. Because of the non-smoothness in the estimating equations and the EER model does not impose any assumptions on the distribution of Y , the traditional likelihood approach in envelope literature cannot be applied to derive the asymptotic distribution of $\hat{\theta}$. Furthermore, there is over-parameterization in the estimating equation in the sense that the number of equations are greater than the number of the parameters. The theoretical tools we use to overcome these issues are Theorem 2.1 in [30] and Proposition 4.1 in [38]. To simplify the notations, we use $\text{avar}(\sqrt{n}\hat{\theta})$ to denote the asymptotic covariance matrix of $\hat{\theta}$ and $\text{avar}(\sqrt{n}\tilde{\theta})$ to denote the asymptotic covariance matrix of $\tilde{\theta}$ hereafter.

First we derive the asymptotic distribution of the ER estimator, and the following regularity conditions are assumed.

- (C1) For each sample size n , $\mathbf{Z}_i = (\mathbf{X}_i^T, Y_i)^T$ is independent and identically distributed. Let f_1 be the conditional density of Y given \mathbf{X} and f_2 be the marginal density function of \mathbf{X} . Then \mathbf{Z}_i has a probability density function $f_1(Y_i|\mathbf{X}_i)f_2(\mathbf{X}_i)$ with respect to a measure $\mu_{\mathbf{Z}}$. Also, the conditional density $f_1(Y|\mathbf{X})$ is continuous in Y for almost all \mathbf{X} .
- (C2) There is a constant $d \geq 0$ and a measurable function $\alpha(\mathbf{Z})$ that satisfy $f_1(Y|\mathbf{X}) \leq \alpha(\mathbf{Z})$, $\int |\mathbf{Z}|^{4+d} \alpha(\mathbf{Z}) f_2(\mathbf{X}) d\mu_{\mathbf{Z}} \leq +\infty$, and $\int \alpha(\mathbf{Z}) f_2(\mathbf{X}) d\mu_{\mathbf{Z}} \leq +\infty$.
- (C3) $E(\mathbf{X}\mathbf{X}^T)$ is nonsingular.
- (C4) Let θ_0 be the true value of θ . The expectation $E_{\theta_0}[s(\mathbf{Z}; \theta)]$ is twice differentiable at θ_0 with $\left. \frac{\partial E_{\theta_0}[s(\mathbf{Z}; \theta)]}{\partial \theta^T} \right|_{\theta=\theta_0}$ having full rank and a finite Frobenius norm. The matrix $E_{\theta_0}[s(\mathbf{Z}; \theta_0)s(\mathbf{Z}; \theta_0)^T]$ is positive definite and has a finite Frobenius norm.
- (C5) The support Θ of θ is a compact set and θ_0 is an interior point of Θ .

Conditions (C1)–(C3) are the conditions used in Theorem 3 of [31] to prove the consistency of the expectile estimator $\tilde{\beta}_\pi$ in ER. Let $\theta_1 = (\mu_\pi, \beta_\pi^T)^T$, $\theta_2 = (\text{vech}(\Sigma_{\mathbf{X}})^T, \mu_{\mathbf{X}}^T)^T$, and let θ_{10} and θ_{20} be the true value of θ_1 and θ_2 respectively. Then the dependence of s_1 , s_2 and s_3 in (3.3) on θ can be specified as $s_1(\mathbf{Z}; \theta) = s_1(\mathbf{Z}; \theta_1)$, $s_2(\mathbf{Z}; \theta) = s_2(\mathbf{Z}; \theta_2)$ and $s_3(\mathbf{Z}; \theta) = s_3(\mathbf{Z}; \theta_2)$.

Theorem 1. Assume the above regularity conditions (C1)–(C5) are satisfied, then we have

$$\sqrt{n}(\tilde{\boldsymbol{\theta}} - \boldsymbol{\theta}_0) \xrightarrow{d} \mathcal{N}(0, \mathbf{C}^{-1} \mathbf{G} \mathbf{C}^{-1}),$$

where

$$\mathbf{C} = \frac{\partial \mathbb{E}_{\boldsymbol{\theta}_0}[s(\mathbf{Z}; \boldsymbol{\theta})]}{\partial \boldsymbol{\theta}^T} \Big|_{\boldsymbol{\theta}=\boldsymbol{\theta}_0} = \begin{pmatrix} -\mathbb{E}_{\boldsymbol{\theta}_0} \left[\mathbf{W} \mathbf{W}^T | I(Y < \mathbf{W}^T \boldsymbol{\theta}_{10}) - \pi | \right] & 0 & 0 \\ 0 & \mathbf{I}_{p(p+1)/2} & 0 \\ 0 & 0 & \mathbf{I}_p \end{pmatrix}$$

and

$$\mathbf{G} = \mathbb{E}_{\boldsymbol{\theta}_0}[s(\mathbf{Z}; \boldsymbol{\theta}_0)s(\mathbf{Z}; \boldsymbol{\theta}_0)^T] = \begin{pmatrix} \mathbf{G}_{1,1} & 0 & 0 \\ 0 & \mathbf{G}_{2,2} & \mathbf{G}_{2,3} \\ 0 & \mathbf{G}_{2,3}^T & \mathbf{G}_{3,3} \end{pmatrix}$$

with

$$\mathbf{G}_{1,1} = \mathbb{E}_{\boldsymbol{\theta}_0} \left[\mathbf{W} \mathbf{W}^T (Y - \mathbf{W}^T \boldsymbol{\theta}_{10})^2 | I(Y < \mathbf{W}^T \boldsymbol{\theta}_{10}) - \pi |^2 \right],$$

$$\mathbf{G}_{2,2} = \text{Var}_{\boldsymbol{\theta}_0} \{ \text{vech}[(\mathbf{X} - \boldsymbol{\mu}_0)(\mathbf{X} - \boldsymbol{\mu}_0)^T] \},$$

$$\mathbf{G}_{2,3} = \mathbb{E}_{\boldsymbol{\theta}_0} \{ \text{vech}[(\mathbf{X} - \boldsymbol{\mu}_0)(\mathbf{X} - \boldsymbol{\mu}_0)^T](\boldsymbol{\mu}_0 - \mathbf{X})^T \},$$

$$\mathbf{G}_{3,3} = \text{Var}_{\boldsymbol{\theta}_0}(\mathbf{X}).$$

The proof of Theorem 1 is given in Section 1.2 of the supplementary material. Since both \mathbf{C} and \mathbf{G} are block diagonal, $\tilde{\boldsymbol{\theta}}_1 = (\tilde{\boldsymbol{\mu}}_\pi, \tilde{\boldsymbol{\beta}}_\pi^T)^T$ is asymptotically independent of $\tilde{\boldsymbol{\theta}}_2 = (\text{vech}(\tilde{\boldsymbol{\Sigma}}_{\mathbf{X}})^T, \tilde{\boldsymbol{\mu}}_{\mathbf{X}}^T)^T$. Theorem 1 provides the asymptotic distribution for all the parameters, and the asymptotic distribution of $\tilde{\boldsymbol{\beta}}_\pi$ agrees with the results in Theorem 3 of [31]. Now we established the asymptotic distribution of the EER estimator.

Theorem 2. Suppose that the EER model (2.4) holds. Under the regularity conditions (C1)–(C5), if $\frac{\partial \mathbb{E}_{\boldsymbol{\theta}_0}[s(\mathbf{Z}; \boldsymbol{\theta})s(\mathbf{Z}; \boldsymbol{\theta})^T]}{\partial \boldsymbol{\theta}^T} \Big|_{\boldsymbol{\theta}=\boldsymbol{\theta}_0}$ has a finite Frobenius norm and the support of $\boldsymbol{\zeta}$ is compact, then

$$\sqrt{n}(\hat{\boldsymbol{\theta}} - \boldsymbol{\theta}_0) \xrightarrow{d} \mathcal{N}(0, \boldsymbol{\Psi}(\boldsymbol{\Psi}^T \mathbf{C} \mathbf{G}^{-1} \mathbf{C} \boldsymbol{\Psi})^\dagger \boldsymbol{\Psi}^T),$$

where $\boldsymbol{\Psi} = \partial \psi(\boldsymbol{\zeta}) / \partial \boldsymbol{\zeta}$.

The proof of Theorem 2 is given in Section 1.3 of the supplementary material. Theorem 2 indicates that the EER estimator is \sqrt{n} -consistent, and is asymptotically normal. Since we have the explicit form of the asymptotic covariance matrices of the ER estimator and the EER estimator, we can compare the efficiency of the two estimators.

Corollary 1. Under the same conditions in Theorem 2, $\text{avar}(\sqrt{n}\hat{\boldsymbol{\theta}}) \leq \text{avar}(\sqrt{n}\tilde{\boldsymbol{\theta}})$.

The proof of Corollary 1 is given in Section 1.4 of the supplementary material. Corollary 1 asserts that the EER estimator is asymptotically more efficient than or as efficient as the ER estimator.

5. Simulations

In this section, we demonstrate the estimation efficiency gains of the EER model via numerical experiments. In the simulations and examples in Section 6, we consider a set of expectile levels of the distribution, and estimation is performed for one level at a time instead of all levels simultaneously. Schnabel and Eilers [37] describes how to determine the complete distribution from a set of expectiles using penalized least squares.

We consider the following simulation settings:

$$Y_i = 3 + \alpha_1^T \mathbf{X}_i + (8 + \alpha_2^T \mathbf{X}_i)\epsilon_i, \quad \text{for } i = 1, \dots, n.$$

We set $p = 12$ and $u_\pi = 2$. Both α_1 and α_2 were p -dimensional vectors. All elements in α_1 were 4. The first $p/2$ elements in α_2 were 0.1 and the rest $p/2$ elements were 0. Four types of error distribution were used to generate ϵ : standard normal distribution $\epsilon \sim \mathcal{N}(0, 1)$, student's t -distribution with 4 degrees of freedom $\epsilon \sim t_4$, mixed normal distribution $\epsilon \sim 0.9\mathcal{N}(0, 1) + 0.1\mathcal{N}(1, 5)$, and exponential distribution with mean 1, i.e., $\epsilon \sim \text{Exp}(1)$.

Based upon the settings, the π th conditional expectile of Y had the following form

$$f_\pi(Y|\mathbf{X}) = 3 + \alpha_1^T \mathbf{X} + (8 + \alpha_2^T \mathbf{X})f_\pi(\epsilon) = 3 + 8f_\pi(\epsilon) + (\alpha_1 + \alpha_2 f_\pi(\epsilon))^T \mathbf{X},$$

where $f_\pi(\epsilon)$ represented the π th expectile of the error distribution. The slope coefficients are contained in $\beta_\pi = \alpha_1 + \alpha_2 f_\pi(\epsilon)$ and the intercept is $\mu_\pi = 3 + 8f_\pi(\epsilon)$. The predictor vector \mathbf{X} followed a normal distribution with mean 0 and covariance matrix $\Sigma_{\mathbf{X}} = \Phi \Lambda \Phi^T + \Phi_0 \Lambda_0 \Phi_0^T$, where Λ was a $u_\pi \times u_\pi$ diagonal matrix with diagonal elements 100 and 9, and Λ_0 was a $(p - u_\pi) \times (p - u_\pi)$ identity matrix. The matrix $\Phi \in \mathbb{R}^{p_1 \times u_\pi}$ was a semi-orthogonal matrix with the first $p/2$ rows being $(\sqrt{6}/6, 0)$ and the remaining $p/2$ rows being $(0, \sqrt{6}/6)$. And the matrix $\Phi_0 \in \mathbb{R}^{p \times (p - u_\pi)}$ was a semi-orthogonal matrix that satisfied $\Phi^T \Phi_0 = 0$. Since $\alpha_1 = \Phi \cdot (4\sqrt{6}, 4\sqrt{6})^T$ and $\alpha_2 = \Phi \cdot (\sqrt{6}/10, 0)^T$, we have $\beta_\pi \in \text{span}(\Phi) = \mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$. Thus $f_\pi(Y|\mathbf{X})$ and \mathbf{X} satisfied the EER model (2.4).

We varied the sample size n from 50 to 800. For each sample size, 100 replications were generated. For each replication, we computed the EER estimator, the ER estimator, the boosting estimator (componentwise gradient boosting expectile regression assuming each predictor has a linear effect) as well as the sparse ER estimator ([18]) of β_π . The boosting estimator was computed by R package `expectreg` [41] with its default settings, i.e. the maximum number of boosting iterations is 4000, and cross validation is used to determine the optimal amount of boosting iterations between 1 and 4000. The sparse ER estimator was computed by R package `SALES` [19]. We use the default choice of the tuning parameter, i.e. the largest value whose cross validation error is within one standard error of the minimum cross validation error. An alternative choice of the tuning parameter is the value that gives the minimum cross validation error. Results based on this alternative choice is included in Section 10 of the supplementary

materials. For each element in β_π , we computed the sample standard deviation for the 100 EER estimators, 100 ER estimators, 100 boosting estimators and 100 sparse ER estimators. We took expectile levels 0.10, 0.25, 0.50, 0.75 and 0.90 as examples. The results of a randomly chosen element in β_π with $\pi = 0.50$ and $\pi = 0.90$ are summarized in Figure 1. They reflect the mean and the upper tail properties of the response. The results of other expectile levels are given in Section 7 of the supplementary materials.

Figure 1 shows substantial efficiency gains achieved by the EER model in the estimation of β_π . With all error distributions and expectile levels π , the sample standard deviations of the EER estimators are much smaller than the sample standard deviations of the ER estimators, the boosting estimators and the sparse ER estimators for all sample sizes. Take the first plot as an example (normal errors and $\pi = 0.5$), with sample size 200, the standard deviation of the EER estimator is already smaller than the asymptotic standard deviation of the ER estimator. The asymptotic standard deviation of the ER estimator in this case is 0.27 and the asymptotic standard deviation of the EER estimator is 0.12. We notice that the boosting estimator and the sparse ER estimator have about the same sample standard deviations as the ER estimator. This is because the variation from the immaterial part affects these two methods in a similar way as to ER. The sample standard deviations of the ER estimators and EER estimators both approach their corresponding asymptotic standard deviations as n increases, which confirms the asymptotic distributions established in Theorem 1 and Theorem 2.

Moreover, we computed the bootstrap standard deviations of the four estimators for each component in β_π using the paired bootstrap method with 100 bootstrap repetitions. For demonstration purpose, we use the normal error as an example and include the results for $\pi = 0.5$ and $\pi = 0.9$ in Figure 2. The results indicate that the bootstrap standard deviation is a good approximation to the actual sample standard deviation. Therefore we use the bootstrap standard deviation as an estimator of the actual standard deviation in data analysis in Section 6.

Now we compared the prediction performance of the EER model with the ER model, the boosting model and the sparse ER model. Under the same simulation settings, we fixed the sample size at $n = 800$ and generated 300 replications. For each replication, 400 samples were used for training and another 400 samples were used for testing. With each of the four models, we first fitted the model to the training samples. Then for each (\mathbf{X}_i, Y_i) in the testing samples, we computed $\hat{f}_\pi(Y_i|\mathbf{X}_i)$, the predicted π th conditional expectile of Y_i given \mathbf{X}_i . The prediction performance is measured by the root mean square error (RMSE) defined as

$$\text{RMSE} = \sqrt{\frac{1}{400} \sum_{i=1}^{400} [f_\pi(Y_i|\mathbf{X}_i) - \hat{f}_\pi(Y_i|\mathbf{X}_i)]^2}.$$

Table 1 and Figures 3 – 6 summarize the RMSEs under the four models with different error distributions. The EER estimator shows a superior prediction

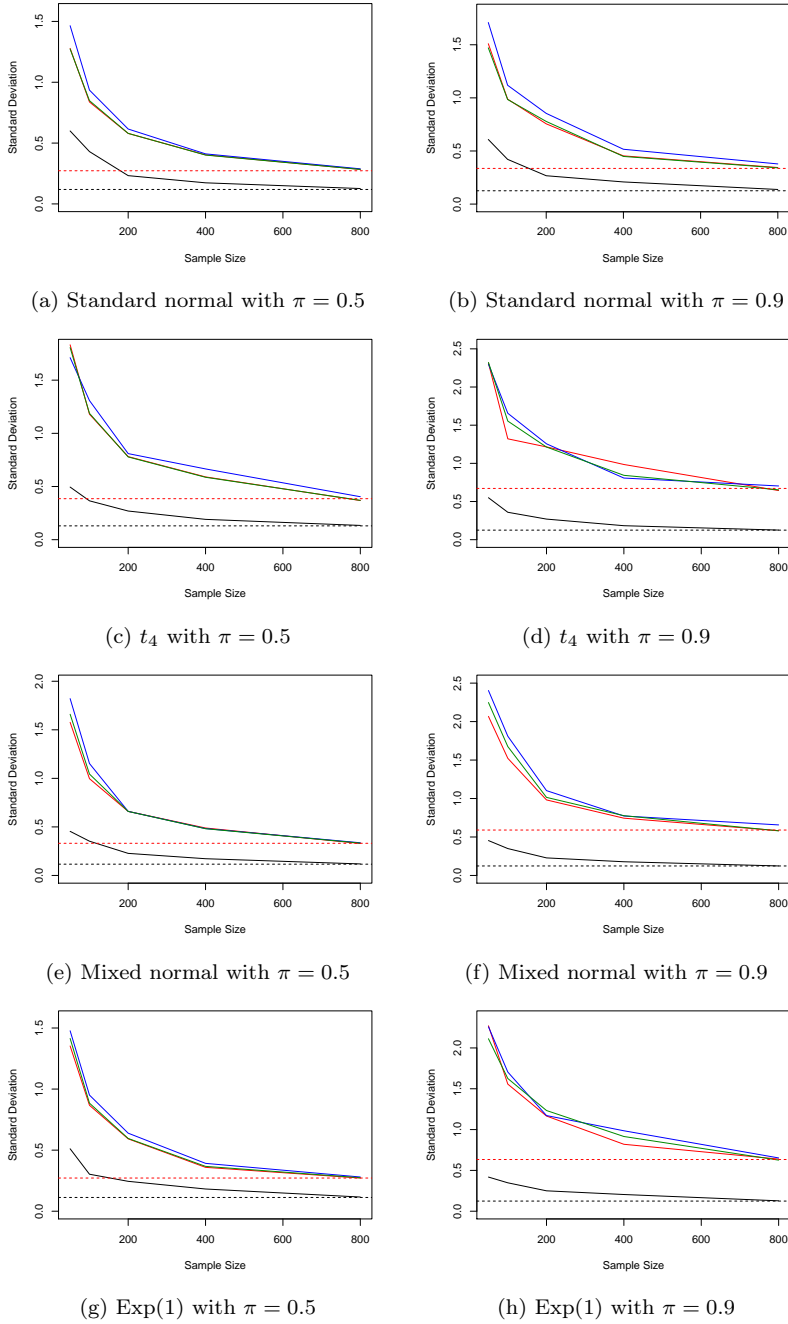


Fig 1: Comparison of the sample standard deviations. Red lines mark the ER estimator. Blue lines mark the sparse ER estimator. Green lines mark the boosting estimator. Black lines mark the EER estimator. The horizontal lines mark the asymptotic standard deviations of the ER estimator (the upper line in each panel) and the EER estimator (the lower line in each panel).

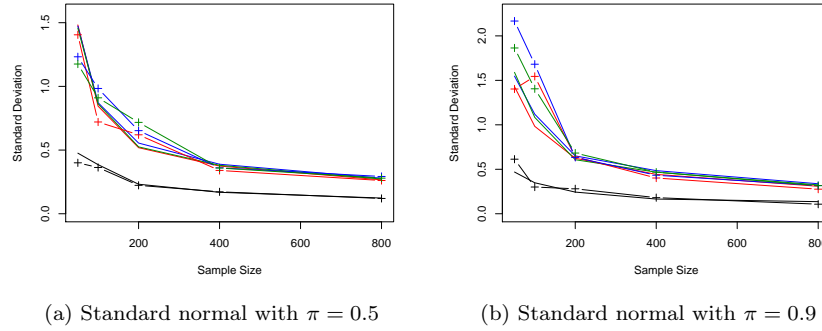


Fig 2: Sample standard deviations and bootstrap standard deviations. Red lines mark the ER estimator. Blue lines mark the sparse ER estimator. Green lines mark the boosting estimator. Black lines mark the EER estimator. Lines with “+” mark the bootstrap standard deviations for the corresponding estimators.

performance in all the cases. The boosting estimator and the ER estimator has comparable performance. Among the four models, the prediction performance of the sparse ER model is the worst because it tends to fit an overly sparse model, which introduces bias to its estimator. Take $\pi = 0.5$ as an example, the EER reduces the average RMSE by 37.7% to 46.3% compared to ER estimator, 37.7% to 46.5% compared to the boosting estimator, and 67.4% to 76.4% compared to the sparse ER estimator.

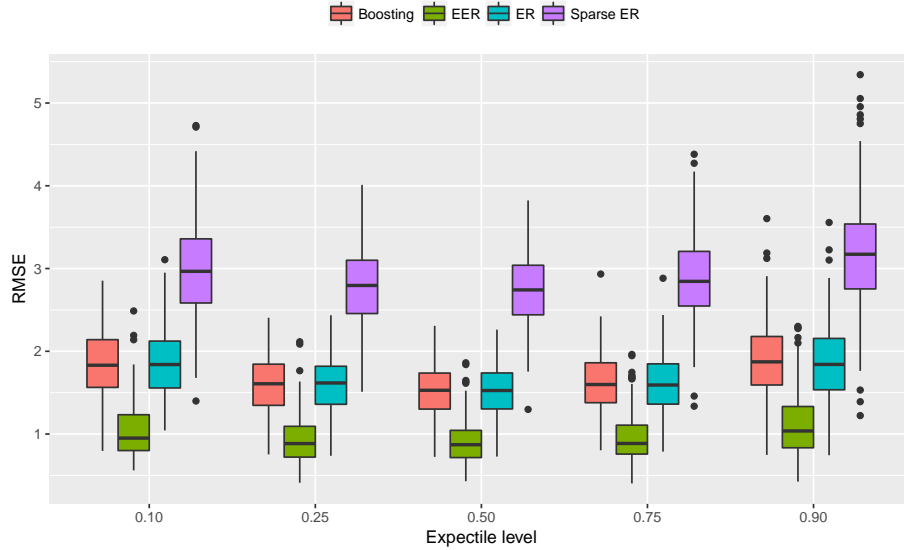


Fig 3: Boxplots of the RMSEs under the four models with $\epsilon \sim \mathcal{N}(0, 1)$.

TABLE 1
 Comparison of the RMSEs, averaged over 300 replications.
 (a) $\epsilon \sim \mathcal{N}(0, 1)$

	EER	ER	Boosting	Sparse ER
$\pi = 0.10$	1.04	1.85	1.86	2.99
$\pi = 0.25$	0.93	1.60	1.60	2.79
$\pi = 0.50$	0.90	1.52	1.52	2.76
$\pi = 0.75$	0.95	1.61	1.62	2.88
$\pi = 0.90$	1.10	1.87	1.91	3.17

(b) $\epsilon \sim t_4$

	EER	ER	Boosting	Sparse ER
$\pi = 0.10$	1.84	3.49	3.50	6.62
$\pi = 0.25$	1.28	2.46	2.47	5.27
$\pi = 0.50$	1.15	2.14	2.15	4.88
$\pi = 0.75$	1.31	2.44	2.45	5.39
$\pi = 0.90$	1.85	3.46	3.51	7.00

(c) $\epsilon \sim 0.9\mathcal{N}(0, 1) + 0.1\mathcal{N}(1, 5)$

	EER	ER	Boosting	Sparse ER
$\pi = 0.10$	1.20	2.21	2.22	3.95
$\pi = 0.25$	1.05	1.87	1.87	3.69
$\pi = 0.50$	1.05	1.86	1.87	3.88
$\pi = 0.75$	1.24	2.21	2.22	4.59
$\pi = 0.90$	1.75	3.14	3.18	6.05

(d) $\epsilon \sim \text{Exp}(1)$

	EER	ER	Boosting	Sparse ER
$\pi = 0.10$	0.70	0.76	0.79	1.96
$\pi = 0.25$	0.77	1.07	1.07	2.58
$\pi = 0.50$	0.96	1.54	1.54	3.40
$\pi = 0.75$	1.34	2.27	2.28	4.51
$\pi = 0.90$	1.99	3.37	3.40	6.03

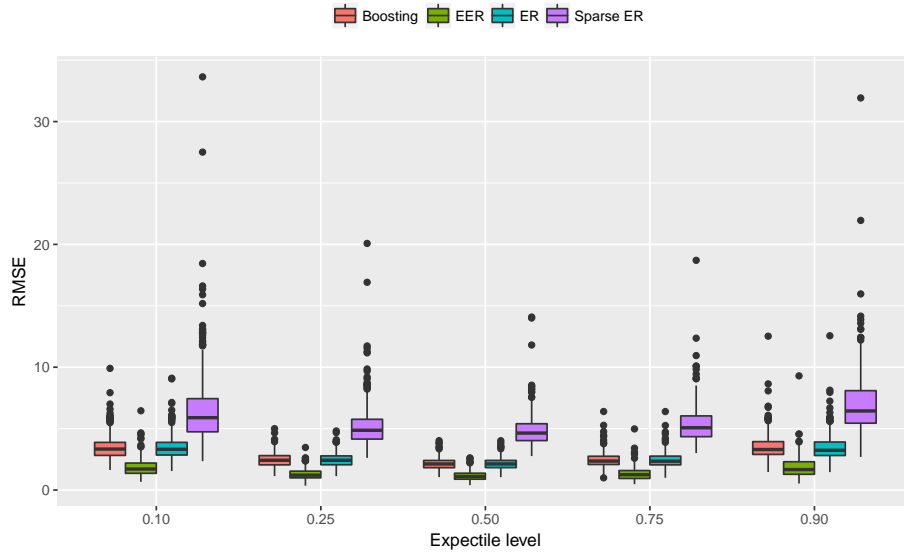


Fig 4: Boxplots of the RMSEs under the four models with $\epsilon \sim t_4$.

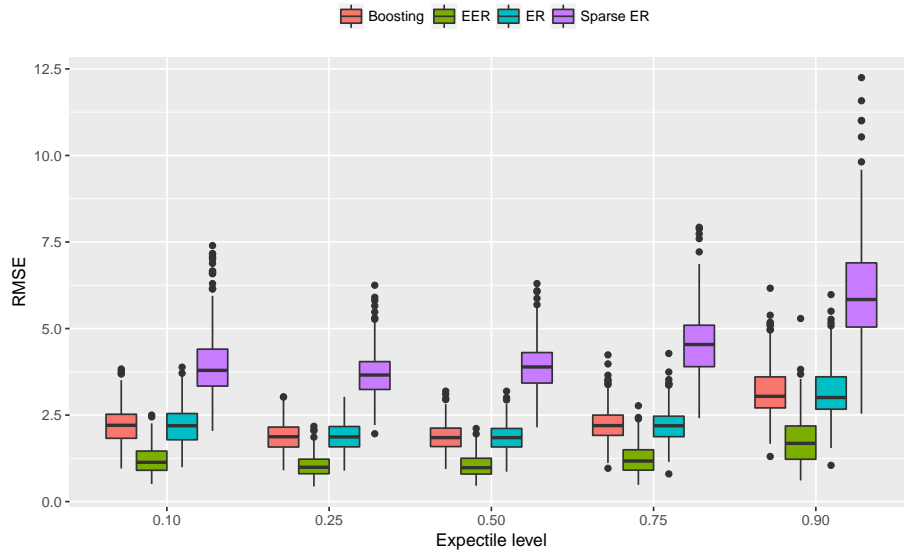


Fig 5: Boxplots of the RMSEs under the four models with $\epsilon \sim 0.9\mathcal{N}(0, 1) + 0.1\mathcal{N}(1, 5)$.

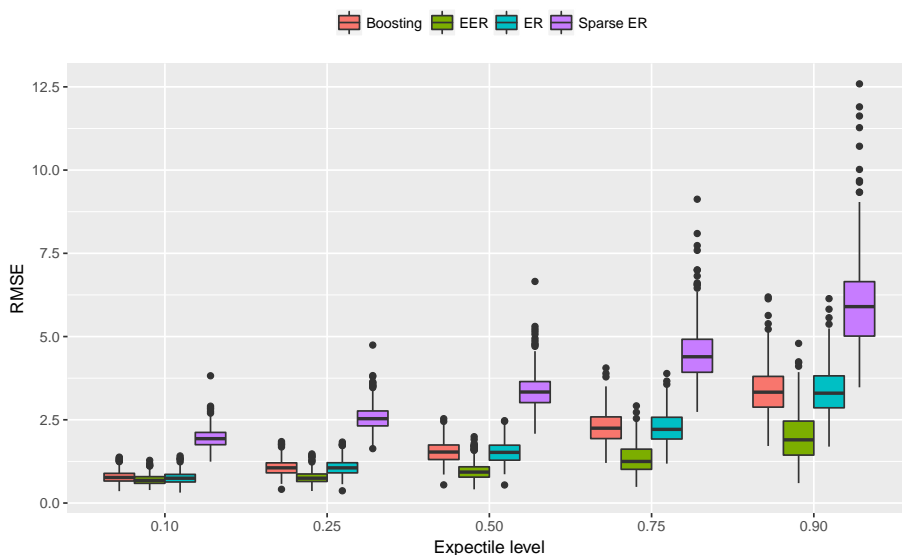


Fig 6: Boxplots of the RMSEs under the four models with $\epsilon \sim \text{Exp}(1)$.

As shown in Section 8 of the supplementary material, the computation complexity of the EER estimator is a polynomial function of the number of the predictors p . Here we outline a comparison on the run time of the ER estimator, the EER estimator, the sparse ER estimator and the boosting estimator under the same setting that produced Table 1 with $\epsilon \sim \mathcal{N}(0, 1)$ and $\pi = 0.10$. Table 2 displays the average run time for each estimator in the 300 replications. We notice that the ER estimator takes the least time to compute. Compare to the ER estimator, the EER estimator takes about three times longer to compute (assuming u_π is known), the sparse ER estimator takes about 30 times longer to compute and the boosting estimator takes about 700 times longer to compute. The trend is similar for other error distributions and expectile levels.

TABLE 2
The run time (in seconds) of the EER estimator, the ER estimator, the boosting estimator and the sparse ER estimator given $\epsilon \sim \mathcal{N}(0, 1)$ and $\pi = 0.10$ in the prediction performance comparison simulation.

ER	EER	Sparse ER	Boosting
0.042	0.13	1.32	29.44

In addition, we examined the performance of RCV in the selection of u_π . In the same settings that generated Figure 1, we applied RCV to choose the envelope dimension u_π . We performed 100 replications for each sample size. The fraction that RCV selects the true dimension $u_\pi = 2$ is summarized in Table 3. RCV shows a stable performance in the selection of u_π . With a small sample size 25, RCV selects the true dimension more than 75% of the time.

And its accuracy increases to 90% when sample size reaches 50. When RCV does not pick the correct dimension, it always overestimates the dimension. A bigger u_π yields a more conservative model, the resulting EER estimator loses some efficiency (compared to the EER model with the correct u_π), but it keeps all the material information and does not introduce bias. Therefore we use RCV to choose u_π in data analysis examples.

In this section, the data is generated from an EER model, and the EER estimator can achieve efficiency gains in estimation and better prediction performance over the boosting model and the sparse ER model. However, the results can be different if the data were generated from a different underlying structure. Since the boosting model and the sparse ER model are variable selection methods, if the sparsity structure rather than the envelope structure is present, or in other words, some predictors are inactive and have coefficients zero, these two models can be more efficient than the EER model by making use of the sparsity structure. A simulation under such setting is included in Section 2 of the supplementary materials. Therefore we cannot conclude if the EER estimator is more efficient or less efficient than the boosting estimator and the sparse ER estimator in general. It depends on the underlying relationship between the response and the predictors, if the envelope structure holds or the sparsity structure holds. If a particular predictor has coefficient zero, then the sparsity structure holds. If β_π is contained in the subspace spanned by some eigenvectors of $\Sigma_{\mathbf{X}}$, then the envelope structure holds. A potentially interesting scenario is that the data have both the envelope structure and the sparsity structure at the same time. A simulation under such setting is included in Section 3 of the supplementary materials. For completion, a simulation with no immaterial part is included in Section 4 of the supplement. In such case, any non-degenerate EER model ($u_\pi < p$) does not hold. However we can still expect to have a smaller mean squared error (MSE) from an approximate EER estimator in some cases due to the bias-variance tradeoff. Since quantiles and expectiles have a one-to-one mapping [52], we also computed the envelope quantile regression estimator [12] and compared it with the EER estimator using the same simulation setting in this section and the S&P 500 data. Details are included in Section 9 of the supplement.

TABLE 3
The fraction that RCV selects the true u_π with different error distributions.

(a) $\epsilon \sim \mathcal{N}(0, 1)$					
	$\pi = 0.10$	$\pi = 0.25$	$\pi = 0.50$	$\pi = 0.75$	$\pi = 0.90$
$n = 25$	76%	80%	81%	79%	86%
$n = 50$	90%	92%	90%	94%	91%
$n = 100$	98%	99%	100%	100%	98%
$n = 200$	100%	100%	100%	100%	100%
$n = 400$	100%	100%	100%	100%	100%
$n = 800$	100%	100%	100%	100%	100%
(b) $\epsilon \sim t_4$					
	$\pi = 0.10$	$\pi = 0.25$	$\pi = 0.50$	$\pi = 0.75$	$\pi = 0.90$
$n = 25$	85%	87%	83%	84%	79%
$n = 50$	94%	92%	96%	94%	93%
$n = 100$	96%	98%	98%	100%	96%
$n = 200$	100%	100%	100%	100%	100%
$n = 400$	100%	100%	100%	100%	100%
$n = 800$	100%	100%	100%	100%	100%
(c) $\epsilon \sim 0.9\mathcal{N}(0, 1) + 0.1\mathcal{N}(1, 5)$					
	$\pi = 0.10$	$\pi = 0.25$	$\pi = 0.50$	$\pi = 0.75$	$\pi = 0.90$
$n = 25$	85%	89%	89%	92%	90%
$n = 50$	98%	98%	95%	92%	92%
$n = 100$	95%	100%	98%	97%	97%
$n = 200$	100%	100%	100%	100%	100%
$n = 400$	100%	100%	100%	100%	99%
$n = 800$	100%	100%	100%	100%	100%
(d) $\epsilon \sim \text{Exp}(1)$					
	$\pi = 0.10$	$\pi = 0.25$	$\pi = 0.50$	$\pi = 0.75$	$\pi = 0.90$
$n = 25$	79%	82%	78%	83%	81%
$n = 50$	91%	96%	96%	94%	91%
$n = 100$	100%	100%	100%	100%	99%
$n = 200$	100%	100%	100%	100%	98%
$n = 400$	100%	100%	100%	100%	100%
$n = 800$	100%	100%	100%	100%	100%

6. Data Analysis

6.1. *state.x77*

The dataset “state.x77” (contained in `datasets` package in R) contains eight measurements including population, average income, illiteracy, life expectancy, murder rate, high-school graduates percentage, land area and frost level for the 50 states in the United States of America. The dataset has been used in [35] as an example of multiple linear regression. Following [35], we took the murder rate as response, and population, average income, illiteracy rate and frost levels as the predictors. The density plot of murder rate indicated it was a bimodal distribution. In addition, we fitted a standard linear regression model on the dataset and checked for the homoscedasticity by Breush-Pagan test [2]. A p-value of 0.03 showed evidence of heteroskedasticity. Therefore, ER is more appropriate to fit the dataset compared to the standard linear regression. We fitted the data using the ER with π varied in different levels at 0.10, 0.25, 0.50, 0.75 and 0.90. Because the four predictors are in quite different scales, they were scaled to have unit standard deviation before the analysis. The analysis results of non-scaled predictors are given in Section 5 of the supplementary materials. RCV was used to select the dimensions of the envelope subspace for different expectile levels. Then the ER estimator, the boosting estimator, the sparse ER estimator and the EER estimator of β_π were computed. For each component in β_π , we calculated the bootstrap standard deviations for the four estimators with 200 bootstrap repetitions. Since the boosting method and the sparse ER model are variable selection methods, some of the four predictors were selected as active and others were selected as inactive. The bootstrap standard deviation comparison of these methods with EER method is only performed on the selected active variables. The dimension selection results for the EER model, the estimated regression coefficients and the efficiency comparison are summarized in Table 4 and Table 5.

TABLE 4
The estimated regression coefficients for the standardized predictors given by the EER model, the ER model, the boosting model and the sparse ER model.

	EER				ER			
	Population	Income	Illiteracy	Frost	Population	Income	Illiteracy	Frost
$\pi = 0.10$	0.949	-0.349	1.382	-1.056	1.288	-0.698	2.010	-0.480
$\pi = 0.25$	0.755	-0.683	1.372	-1.345	1.152	-0.403	2.227	-0.280
$\pi = 0.50$	0.667	-0.648	1.327	-1.298	0.999	0.040	2.525	0.030
$\pi = 0.75$	0.954	-0.023	1.451	-1.074	0.921	0.338	2.729	0.271
$\pi = 0.90$	0.379	-0.404	0.820	-0.829	0.791	0.441	2.794	0.410
	Boosting				Sparse ER			
	Population	Income	Illiteracy	Frost	Population	Income	Illiteracy	Frost
$\pi = 0.10$	1.272	-0.689	1.965	-0.520	0.506	0.000	1.444	-0.490
$\pi = 0.25$	1.029	-0.253	2.187	-0.262	0.361	0.000	1.763	-0.153
$\pi = 0.50$	0.846	0.000	2.313	0.000	0.000	0.000	1.472	0.000
$\pi = 0.75$	0.529	0.000	2.045	0.000	0.000	0.000	1.293	0.000
$\pi = 0.90$	0.283	0.000	1.780	0.000	0.000	0.000	0.563	0.000

TABLE 5

Dimension selection results and efficiency comparison among the EER estimator, the ER estimator, the boosting estimator and the sparse ER estimator. Columns 3-4 contain the bootstrap standard deviation ratios of the ER estimator versus the EER estimator. Columns 5-6 contain the bootstrap standard deviation ratios of the sparse ER estimator versus the EER estimator. And columns 7-8 contain the bootstrap standard deviation ratios of the boosting estimator versus the EER estimator.

\hat{u}_π	π	ER to EER		Sparse ER to EER		Boosting to EER		
		Range	Average	Range	Average	Range	Average	
	$\pi = 0.10$	2	1.21 – 1.75	1.50	1.32 – 1.66	1.49	1.04 – 1.46	1.26
	$\pi = 0.25$	1	1.20 – 2.72	1.91	1.01 – 1.88	1.45	1.13 – 2.00	1.55
	$\pi = 0.50$	1	1.35 – 2.77	2.08	1.56 – 1.56	1.56	1.21 – 1.50	1.36
	$\pi = 0.75$	2	1.05 – 1.58	1.32	1.22 – 1.22	1.22	1.10 – 1.23	1.17
	$\pi = 0.90$	1	1.90 – 3.00	2.32	2.26 – 2.26	2.26	1.70 – 2.01	1.86

From Table 4 we can see the estimated regression coefficients given by the EER model and the ER model differ from each other. Take the predictor income (the per capita income) as an example, the ER model suggests that income has a negative effect on murder rate at lower quantiles ($\pi = 0.10$ and $\pi = 0.25$) and has a positive effect to murder rate at high quantile levels ($\pi = 0.50$, $\pi = 0.75$ and $\pi = 0.90$) while the EER models indicates that income has a negative effect on murder rate for all quantile levels. The results from the EER model seems to be more meaningful.

Table 5 shows the efficiency gains of the EER estimator over the other estimators at all investigated expectile levels. Taking $\pi = 0.90$ for example, RCV selected $u_\pi = 1$. The ratios of the bootstrap standard deviations of the ER estimator versus the EER estimator range from 1.92 to 2.99 with an average of 2.31. To achieve the same efficiency gains under the ER, we need to increase the sample size to $2.31^2 \approx 5.3$ times the original sample size. Similar efficiency gains are also noticed when comparing with the sparse ER estimator and the boosting estimator. The efficiency gains also lead to different interpretations of the data. For instance, with $\pi = 0.5$, population and illiteracy rate are significant predictors with positive coefficients under ER. Income and frost level (number of days with minimum temperature below freezing) are not significant. Sparse ER model selects illiteracy rate as the only active predictor with a positive coefficient. The boosting method selects population and illiteracy rate as active predictors with positive coefficients, and income and frost level as inactive predictors. However, under the EER model, all predictors are significant. While population and illiteracy rate have positive coefficients, income and frost level have negative association with murder rate. This indicates that the EER estimator can detect weaker signal from the data from improved estimation efficiency. The predictive performance of the EER estimator is slightly worse than the ER estimator because the RCV tends to select a parsimonious model, which may have a larger predicted expectile loss than the ER model. The details are included in Section 6 of the supplementary materials.

6.2. S&P 500 Index

Now we provide another example using the S&P 500 Index data to show that the efficiency gains from the EER model can lead to a better prediction performance. The data [25] contains 351 quarterly economic observations from January, 1927 to December, 2014. The response was the equity premium, which is the return on the S&P 500 Index minus the return on treasury bill. We used 11 quarterly predictors following [3, 48, 9, 29, 33, 25]. The predictors included dividend yield (the difference between the log of dividends and the log of lagged prices), earnings–price ratio (the difference between the log of earnings and the log of prices), book–to–market ratio (the ratio of book value to market value for the Dow Jones Industrial Average), net equity expansion (ratio of 12-month moving sums of net issues by NYSE-listed stocks divided by the total market capitalization of NYSE stocks), stock variance (the sum of squared daily returns on the S&P 500), treasury bill rate (the 3-month rate), term spread (difference between the long-term yield on government bonds and the treasury bill rate), long–term rate of return for government bonds, default yield spread (difference between BAA– and AAA–rated corporate bond yields), default return spread (the difference between the return on long-term corporate bonds and the return on long–term government bonds) and inflation. Some of the predictors were stock characteristics and the others reflected operation conditions of the selected companies. All of them had significant impacts on S&P 500 Index. Since an investigation on the dataset showed strong evidence of heteroskedasticity, the standard linear regression is not adequate to explore the relationship between the response and the predictors. Moreover, there were two extreme values in the response. Hence instead of a QR, we conducted an ER on the dataset which is more sensitive to the extreme values and could make more efficient use of the information in the dataset.

We fitted the data using the model:

$$f_{\pi}(Y_{t+1}|\mathbf{X}_t) = \mu_{\pi} + \beta_{\pi}^T \mathbf{X}_t,$$

where Y_{t+1} was the equity premium at time $t + 1$ and \mathbf{X}_t was the predictor vector at time t . Both the ER and the EER model were applied to predict the conditional expectile of the response $f_{\pi}(Y_{t+1}|\mathbf{X}_t)$ in a moving window with a size of 80 quarters, i.e., use observations $\{(\mathbf{X}_t, Y_{t+1}), t = t_0 - 80, \dots, t_0 - 1\}$ to predict $f_{\pi}(Y_{t_0+1}|\mathbf{X}_{t_0})$, where $t_0 = 81, \dots, 351$. We used 80 as the size of the moving window because [25] showed that a 20-years estimation window delivers better results than alternative estimation windows. The predicted expectile loss is an important measure of the prediction performance in ER. Once we have $\hat{\mu}_{\pi}$ and $\hat{\beta}_{\pi}$, the predicted expectile loss is computed as $\phi_{\pi}(Y_{t_0+1} - \hat{\mu}_{\pi} - \hat{\beta}_{\pi}^T \mathbf{X}_{t_0})$.

We took expectile levels $\pi = 0.10, 0.25, 0.50, 0.75$ and 0.90 as examples. The results of the predicted expectile losses are summarized in Table 6. Table 6 shows that the EER model has smaller mean predicted expectile loss compared to the ER model with all the expectile levels. In addition, the EER model also has smaller standard deviations of the predicted expectile losses, which indicates

TABLE 6

Mean and standard deviations of the predicted expectile losses under the ER and the EER model with different expectile levels. The 2nd and 3rd columns give the mean of the predicted expectile losses (in the unit of 10^{-3}). The 5th and 6th columns give the standard deviations of the predicted expectile losses (in the unit of 10^{-3}). The 4th and 7th columns represent percentage reduction of the EER model compared to the ER on the relative quantity.

	Mean			Standard Deviation		
	ER	EER	Reduction	ER	EER	Reduction
$\pi = 0.10$	3.34	2.26	32.26%	7.56	4.98	34.05%
$\pi = 0.25$	4.67	2.98	36.19%	10.15	5.88	42.05%
$\pi = 0.50$	5.32	3.10	41.74%	13.82	5.61	59.39%
$\pi = 0.75$	4.59	2.70	41.16%	15.19	4.19	72.43%
$\pi = 0.90$	3.58	2.29	36.06%	17.44	11.70	32.90%

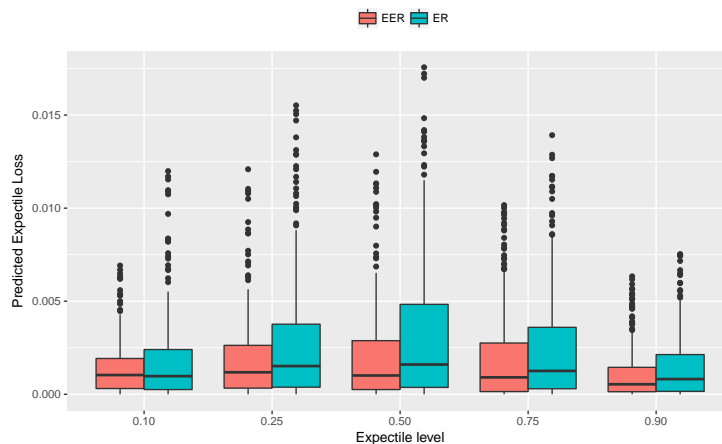


Fig 7: Boxplots of the trimmed predicted expectile losses.

that the EER model can give a more stable prediction. Figure 7 contains boxplots to graphically display the location and spread of the predicted expectile losses. Since there were some outliers, we trimmed the largest 5% of the predicted expectile losses under both the ER and the EER model for each expectile level to improve visibility. Both Table 6 and Figure 7 demonstrate substantial advantage of the EER model in prediction performance over the ER model.

We should note that the response in this example is actually weakly autocorrelated as revealed from its autocorrelation function (ACF) plot and the partial autocorrelation function (PACF) plot. Therefore an EER model that accommodates time dependent data is more suitable for the analysis of this dataset. [24] extended the asymptotic results of [31] to allow for stationary and weakly dependent data in the ER model. In some applications, the time effects can also be formulated by a mixed regression, and mixed expectile models are studied in

[46]. The development of an EER model for time dependent data or for mixed expectile model is a potentially interesting future research direction and can have applications in financial, medical or meteorological datasets.

7. Extension to Semiparametric Settings

As an anonymous reviewer pointed out that one advantage to expectiles is the possibility to have very flexible semiparametric predictors. In this section, we consider a semiparametric ER model where the response is related to a combination of linear predictors $\mathbf{X} \in \mathbb{R}^{p_1}$ and nonlinear predictors $\mathbf{Z} \in \mathbb{R}^{p_2}$

$$f_\pi(Y|\mathbf{X}, \mathbf{Z}) = \mu_\pi + \beta_\pi^T \mathbf{X} + g(\mathbf{Z}), \quad (7.1)$$

where g is a smooth function, μ_π is the intercept and β_π contains coefficients for the linear predictors. We assume that $E(g(\mathbf{Z})) = 0$ for identification.

To impose the envelope structure on β_π , we use the technique in partial envelope model [43] and consider the $\Sigma_{\mathbf{X}}$ -envelope of β_π , denoted by $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$. Let u_π denote its dimension, and let $\Gamma_\pi \in \mathbb{R}^{p \times u_\pi}$ and $\Gamma_{0\pi} \in \mathbb{R}^{p \times (p-u_\pi)}$ be orthonormal bases of $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)$ and $\mathcal{E}_{\Sigma_{\mathbf{X}}}(\beta_\pi)^\perp$ respectively. Then β_π can be written as $\beta_\pi = \Gamma_\pi \eta_\pi$, where $\eta_\pi \in \mathbb{R}^{u_\pi}$ contains the coordinates of β_π with respect to Γ_π . The covariance matrix $\Sigma_{\mathbf{X}}$ can be written as $\Sigma_{\mathbf{X}} = \Gamma_\pi \Omega_\pi \Gamma_\pi^T + \Gamma_{0\pi} \Omega_{0\pi} \Gamma_{0\pi}^T$, where Ω_π contains the coordinates of $\Sigma_{\mathbf{X}}$ with respect to Γ_π and $\Omega_{0\pi}$ contains the coordinates of $\Sigma_{\mathbf{X}}$ with respect to $\Gamma_{0\pi}$. Then the semiparametric EER model is formulated as

$$f_\pi(Y|\mathbf{X}, \mathbf{Z}) = \mu_\pi + \eta_\pi^T \Gamma_\pi^T \mathbf{X} + g(\mathbf{Z}), \quad \Sigma_{\mathbf{X}} = \Gamma_\pi \Omega_\pi \Gamma_\pi^T + \Gamma_{0\pi} \Omega_{0\pi} \Gamma_{0\pi}^T. \quad (7.2)$$

Note that the envelope structure is only imposed on the linear predictor \mathbf{X} , not the entire predictor vector $(\mathbf{X}^T, \mathbf{Z}^T)^T$. This means that the conditional expectile depends on \mathbf{X} only through $\Gamma_\pi^T \mathbf{X}$, i.e. $f_\pi(Y|\mathbf{X}, \mathbf{Z}) = f_\pi(Y|\Gamma_\pi^T \mathbf{X}, \mathbf{Z})$, and the variation of \mathbf{X} can be decomposed into the variation of the material part and variation of the immaterial part, i.e., $\Sigma_{\mathbf{X}} = \text{var}(\mathbf{P}_\Gamma \mathbf{X}) + \text{var}(\mathbf{Q}_\Gamma \mathbf{X})$. By linking β_π to the material part, the semiparametric EER model (7.2) is expected to improve the efficiency in the estimation of β_π . To impose envelope structure on $g(\mathbf{Z})$ involves completely different scopes and techniques, and we leave it as an important future research direction. To estimate from model (7.2), we use the following iterative algorithm. We denote the estimated linear part as \hat{Y}_1 , and the estimated nonlinear part as \hat{Y}_2 .

- Step 1** : Initialize $\hat{Y}_2 = 0$.
- Step 2** : Fit the EER model with $Y - \hat{Y}_2$ as the response and \mathbf{X} as the predictors. Use the EER estimators to update \hat{Y}_1 .
- Step 3** : Fit the ER additive model [40] with Y being the response, $\hat{\Gamma}_\pi^T \mathbf{X}$ being the linear predictors and \mathbf{Z} being the nonlinear predictors, where $\hat{\Gamma}_\pi$ was calculated from Step 2. Update \hat{Y}_2 .
- Step 4** : Repeat Step 2 – Step 3 until the convergence of $\hat{Y}_1 + \hat{Y}_2$.

To illustrate the performance of semiparametric EER, we consider the following simulation settings:

$$Y_i = 3 + \boldsymbol{\alpha}_1^T \mathbf{X}_i + h(\mathbf{Z}_i) + (8 + \boldsymbol{\alpha}_2^T \mathbf{X}_i)\epsilon_i, \quad \text{for } i = 1, \dots, n,$$

where $\mathbf{X} \in \mathbb{R}^{p_1}$ contains the linear predictors and $\mathbf{Z} \in \mathbb{R}^{p_2}$ contains the nonlinear predictors. We set $p_1 = 12$ and $p_2 = 4$. The nonlinear predictors $\mathbf{Z} = (Z_1, Z_2, Z_3, Z_4)$ followed the multivariate normal distribution $\mathcal{N}_4(0, \mathbf{I}_4)$ and the function $h(\mathbf{Z}) = \exp(Z_1) + \sin(Z_2) + \cos(Z_3) + Z_4^3$. The error ϵ was generated from the standard normal distribution.

Based upon the settings, the π th conditional expectile of Y has the following form

$$\begin{aligned} f_\pi(Y|\mathbf{X}, \mathbf{Z}) &= 3 + \boldsymbol{\alpha}_1^T \mathbf{X} + h(\mathbf{Z}) + (8 + \boldsymbol{\alpha}_2^T \mathbf{X})f_\pi(\epsilon) \\ &= 3 + 8f_\pi(\epsilon) + \mathbb{E}(h(\mathbf{Z})) + (\boldsymbol{\alpha}_1 + \boldsymbol{\alpha}_2 f_\pi(\epsilon))^T \mathbf{X} + h(\mathbf{Z}) - \mathbb{E}(h(\mathbf{Z})), \end{aligned}$$

where $f_\pi(\epsilon)$ represents the π th expectile of the standard normal distribution. For the linear part, the slope coefficients are contained in $\boldsymbol{\beta}_\pi = \boldsymbol{\alpha}_1 + \boldsymbol{\alpha}_2 f_\pi(\epsilon)$ and the intercept is $\mu_\pi = 3 + 8f_\pi(\epsilon) + \mathbb{E}(h(\mathbf{Z}))$. The parameters $\boldsymbol{\alpha}_1$, $\boldsymbol{\alpha}_2$ and the predictor vector \mathbf{X} were generated in the same way as in Section 5. So the linear part follows an envelope structure. The nonlinear part was given by $g(\mathbf{Z}) = h(\mathbf{Z}) - \mathbb{E}(h(\mathbf{Z}))$.

We varied the sample size n from 50 to 800. For each sample size, 100 replications were generated. For each replication, we computed the semiparametric ER estimator of $\boldsymbol{\beta}_\pi$ by the ER additive model ([41]), and computed the semiparametric EER estimator of $\boldsymbol{\beta}_\pi$ using the preceding algorithm. Then for each component of $\boldsymbol{\beta}_\pi$, we computed the sample standard deviations for the semiparametric ER estimator and the semiparametric EER estimator based on the results from the 100 replications. We took expectile levels $\pi = 0.50$ and $\pi = 0.90$ as examples. The results of a randomly chosen component in $\boldsymbol{\beta}_\pi$ are summarized in Figure 8. The sample standard deviations of the semiparametric EER estimators are much smaller than the sample standard deviations of the semiparametric ER estimators. For example, when $n = 200$, the standard deviation of the semiparametric ER estimator is 0.61 while the standard deviation of the semiparametric EER estimator is 0.23 for $\pi = 0.5$. For $\pi = 0.9$, the standard deviations are 0.72 and 0.29 for the semiparametric ER estimator and the semiparametric EER estimator. This indicates that the envelope structure also yields substantial efficiency gains in estimation of $\boldsymbol{\beta}_\pi$ under the semiparametric ER context.

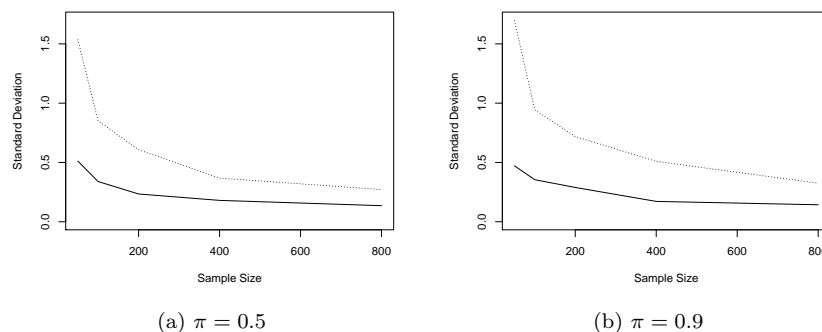


Fig 8: Sample standard deviations. Dashed lines mark the semiparametric ER estimator. Solid lines mark the semiparametric EER estimator.

The baseball salary data was studied in [47] to determine whether the salary of a baseball player is affected by his offensive performance. The data contains the salary information from the 1992 season for 337 Major League Baseball (MLB) non-pitchers who played at least one game during both the 1991 and 1992 seasons. It also provides 12 offensive statistics for each player from the 1991 season including batting average, on-base percentage, number of runs, hits, doubles, triples, home runs, batted in, walks, strike-outs, stolen bases and errors. We took the salary as response and the 12 offensive statistics as predictors. By exploring the scatter plots of the response versus each predictors, we identified five predictors for which the association with the response is not sufficiently explained by a linear relationship. The five predictors were batting average, on-base percentage, number of triples, strike-outs and errors. Therefore, they were used as nonlinear predictors and the remains were used as linear predictors. Before the analysis, each predictor was scaled to have unit standard deviation. We fitted the semiparametric ER model and the semiparametric EER model with $\pi = 0.5$ and 0.9 to the data. RCV suggested $u_\pi = 1$ for $\pi = 0.5$. For each element in β_π , we calculated the bootstrap standard deviations for both the semiparametric ER estimator and the semiparametric EER estimator with 100 bootstrap samples. The ratios of bootstrap standard deviations of the semiparametric ER estimator versus the semiparametric EER estimator range from 3.99 to 13.65 with an average of 9.34. For $\pi = 0.9$, $u_\pi = 2$ was selected by RCV for the EER model. The ratios of the bootstrap standard deviations range from 1.32 to 10.87 with an average of 5.44. The results indicate the semiparametric EER model achieves substantial efficiency gains compared to the semiparametric ER model. To get the same average estimation efficiency under the semiparametric ER model, we need to increase the sample size to $9.34^2 \approx 87$ times the original sample size with $\pi = 0.5$ and $5.44^2 \approx 30$ times the original sample size with $\pi = 0.9$.

8. Discussion and Future Work

In this paper, we develop the EER model as an efficient estimation method for the ER. We estimate the parameters using GMM and established the asymptotic distribution of the estimators. Efficiency gains are demonstrated both theoretically and numerically. A potentially interesting extension is to perform variable selection to the predictors and develop a sparse EER model. In many applications such as the S&P 500 Index data, some predictors do not affect certain conditional expectile(s) of the response and have coefficients zero. It is of practical interest to identify those predictors, especially in high dimensional settings. Another direction is to derive the EER model with censored response. Censored data are often encountered in medical studies and econometrics when the variable of interest is only observed under certain conditions, such as top coding in income surveys, e.g. “\$500,000 and above” ([34]). An EER model that can accommodate censored response will be applicable in such settings.

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