Fused LASSO as Non-Crossing Quantile Regression*

Tibor Szendrei [†]
Department of Economics, Heriot-Watt University, UK.
National Institute of Economic and Social Research, UK.

Arnab Bhattacharjee Department of Economics, Heriot-Watt University, UK. National Institute of Economic and Social Research, UK.

Mark E. Schaffer Department of Economics, Heriot-Watt University, UK.

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Abstract

Growth-at-Risk is vital for empirical macroeconomics but is often suspect to quantile crossing due to data limitations. While existing literature addresses this through post-processing of the fitted quantiles, these methods do not correct the estimated coefficients. We advocate for imposing non-crossing constraints during estimation and demonstrate their equivalence to fused LASSO with quantile-specific shrinkage parameters. By re-examining Growth-at-Risk through an interquantile shrinkage lens, we achieve improved left-tail forecasts and better identification of variables that drive quantile variation. We show that these improvements have ramifications for policy tools such as Expected Shortfall and Quantile Local Projections.

Keywords: Interquantile shrinkage, Crossing Quantile Curves, High-Dimensional Econometrics, Growth-at-Risk.

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[†]Corresponding author: t.szendrei@niesr.ac.uk

1 Introduction

Growth-at-Risk (GaR) has become a key measure of economic vulnerability since the work of Adrian et al. (2019). The key idea is to think about GDP growth through the lens of value-at-risk, and use quantile regression of Koenker and Bassett (1978) to uncover nonlineary macro-financial linkages. The need for GaR was highlighted by the global financial crisis which showed how downside risks, or lower quantiles of the density of GDP growth, evolve with the state of financial market.

GaR has become a popular tool for policy makers and researchers alike. Figueres and Jarociński (2020) apply the method to euro area data and show that the macro-financial linkages driving skewness are not specific to the US. Kohns and Szendrei (2024) use shrinkage methods to fit GDP densities with over 200 covariates and find that accounting for multiple sources of risk leads to better density fits of GDP growth. Iseringhausen et al. (2023) use the same dataset and look at the skewness of the fitted quantile estimates, which encompasses the aggregate impact of the considered covariates. They show that aggregate skewness is highly pro-cyclical. Mitchell et al. (2024) use quantiles estimated on US GaR and show that multimodality in the fitted densities is just as important as skewness.¹. Finally, Carriero et al. (2025) has looked at specifications for quantile regression for empirical macroeconomics, with GaR being one of the models being investigated.

While GaR is undoubtedly an important policy tool, it is a macroeconometric approach which in turn has data constraints. This data scarcity is particularly a problem for tail estimation (Koenker, 2005), which is the main goal for a GaR model. One big concern when applying quantile regression in data scarce settings is the notion of quantile crossing, i.e. the situation where the fitted quantiles are not monotonically increasing. This yields improper densities and the literature has tackled this problem using one of two methods. One approach is to fit some distribution one the estimated quantiles in each time period, as done in Adrian et al. (2019) or Korobilis (2017). Another approach is to simply sort the estimated quantiles in each period as advocated by Chernozhukov et al. (2009) and Chernozhukov et al. (2010).²

All these approaches that have been popular in the macroeconometric literature involve some form of post-processing the estimated quantiles. While these two-step methods yield

¹Adrian et al. (2021) and Kohns and Szendrei (2024) also discuss the importance of multimodality stemming from macro-financial linkages

²There is also the novel method of Mitchell et al. (2024) which proposes a nonparametric method to construct densities from sorted quantiles.

proper densities, simply correcting the fitted quantiles does not quantify corresponding changes in the estimated coefficients, i.e., the coefficients estimated in the first step remain uncorrected. This is particularly a problem as quantile coefficients are used to construct quantile IRFS (Chavleishvili and Manganelli, 2024), or quantile local projections (Ruzicka, 2021). As such, for macroeconometric inference it would be desirable to impose non-crossing during estimation of the quantiles. This is the motivation in Szendrei and Varga (2023), who use the non-crossing constraints of Bondell et al. (2010) to identify variables that drive macro-financial nonlinearity in the Euro Area GaR. Szendrei and Varga (2023) also find that imposing non-crossing constraints also improves the forecasted densities of GaR.

While non-crossing constraints are alluring for empirical macroeconomic applications of quantile regression, we do not know how these constraints influence the estimated coefficients. The primary motivation of this paper is to study the impact of such constraints on the model parameters. This is achieved by implementing a set of non-crossing constraints that can be scaled, which in turn makes the non-crossing constraints tighter or looser. Using this adaptive non-crossing constraint we show that non-crossing constraints are equivalent to fused LASSO with quantile specific shrinkage parameters. We propose an estimator using these adaptive non-crossing constraints (Generalised Non-crossing Constraint Quantile Regression), where the hyperparameter regulating interquantile shrinkage is obtained by cross-validation.

We conduct comprehensive Monte Carlo exercises based on those used in the original paper by Bondell et al. (2010), that first proposed non-crossing quantile regression. Through these experiments we show that the proposed estimator is capable of providing further improvements in fit compared to the original BRW estimator. We also consider the variable selection properties of the different estimators and verify that non-crossing constraints are a type of fused shrinkage. The Monte Carlos also reveal how the BRW estimator undershrinks quantile variation while the traditional Fused LASSO, as outlined in Jiang et al. (2013), overshrinks.

We then estimate US GaR using the variables outlined in Adrian et al. (2019). We will investigate the canonical GaR with a interquantile shrinkage framework, i.e. identifying variables that drive nonlinearities. Adrian et al. (2019) alluded to this in their paper, but the estimator they use did not enforce limiting quantile variation during estimation, which influences their downside risk measures. In this GaR exercise our proposed estimator fares much better at distinguishing quantile varying variables than the traditional Fused LASSO. Furthermore, in a small pseudo forecasting exercise we find that the estimator with adaptive non-crossing constraints yields the best left tail forecasts. These findings add to Carriero et al. (2025), namely

that fused shrinkage is also important for GaR. Our results also indicate that it is also important how this shrinkage is induced: adaptive non-crossing constraints yield improvements, while the traditional fused LASSO seem to lead to worse forecasting performance.

These differences in GaR coefficients have an influence on tools used in policy. We show how the choice of imposing interquantile shrinkage, and how to impose it, influences the estimated Expected Shortfall of GDP growth. In particular, Expected Shortfall based on traditional quantile regression is more volatile from one period to the next, while the proposed estimator produces smoother expected shortfall values over time, while still having values during periods of financial stress. We also produce quantile local projections of how GDP responds over time to a unit shock in the financial stress measure. Just like with the expected shortfall values, the LPs of the proposed estimator are smoother than those of the quantile regression LPs. These examples show the policy relevance of implementing interquantile shrinkage in GaR.

This paper is structured as follows. In Section 2 we introduce the quantile regression of Koenker and Bassett (1978) along with the non-crossing constraint of Bondell et al. (2010), before providing the adaptive non-crossing constraints that vary with a hyperparameter α . Using these new constraints, we show that one can rewrite them into the Fused LASSO constraint of Jiang et al. (2013). The section concludes by describing how the hyperparameter is chosen. Section 3 describes the Monte Carlo experiment before presenting the fit and variable selection performance of the different estimators. This is followed by the US Growth-at-Risk application in Section 4. The paper concludes with the key takeaways of the method in Section 5.

2 Methodology

2.1 Quantile Regression

The first building block of the proposed methodology is the quantile regression framework of Koenker and Bassett (1978). Quantile regression is a weighted version of the least absolute deviation regression, and yields lines of best fit that explain different parts of the distribution. The collection of Q estimated quantiles can be used to describe the distribution of a response variable Y conditional on a vector of response variables $X \in \mathbb{R}^K$. Formally, the q^{th} quantile is modelled in regression setting as:

$$Q_{\tau_q} = X^T \beta_{\tau_q}.$$

The collection of the Q quantiles parameters $\beta = \{\beta_{\tau_1}, \beta_{\tau_2}, ..., \beta_{\tau_Q}\}$ describe the conditional distribution. The goal is to estimate the vector of coefficients $\beta_{\tau_q} \in \mathbb{R}^K$ for all quantiles.³ This can be done using quantile regression:

$$\hat{\beta} = \underset{\beta}{\operatorname{argmin}} \sum_{q=1}^{Q} \sum_{t=1}^{T} \rho_{\tau_q} (y_t - x_t^T \beta_{\tau_q})$$

$$\rho_{\tau_q}(u) = u(\tau_q - I(u < 0))$$
(1)

where the second equation is the 'tick-loss' function (Koenker and Bassett, 1978).

Quantile regression is not the only asymmetric estimator which uses the 'tick-loss' function as a way to model different parts of the distribution. Newey and Powell (1987) introduced the concept of expectile regression, i.e., the 'tick-loss' weight applied to the ℓ_2 norm of the residuals. The advantage of expectile regression is that the objective function is differentiable, unlike quantile regression's. Furthermore, when setting $\tau = 0.5$ in an expectile regression, one recovers the OLS estimator, i.e., the conditional mean.

Expectile regression does come with some disadvantages. First, interpretation: while conditional quantiles can be interpreted as splitting the data into τ and $1-\tau$ parts, i.e. the τ^{th} conditional quantile of the sample, the same is not true for expectiles. Instead expectiles are the τ^{th} quantile of some distribution which is related to the original distribution of Y as shown in Jones (1994). Second, since expectile regression is based on the ℓ_2 norm of the residuals, outliers in the dependent variable impacts all expectiles, while outliers only exert undue influence on extreme quantiles in the case of quantile regression. As such, "fan-shaped" estimated quantiles are a clear indication of heteroskedasticity, while the same shape for expectiles might simply be due to outliers in the dependent variable (Newey and Powell, 1987; Sobotka and Kneib, 2012). On account of these disadvantages, in this paper we will focus on quantile regression and leave for future research to explore how our findings translate to expectiles.

Equation (1) gives an estimate for the parameters with which a description of the conditional distribution is obtained, but it is possible that these fitted quantiles cross. Quantile crossing is a violation of monotonocity assumption and often occurs on account of data scarcity or model misspecification (Koenker, 2005). Limits in data availability are frequently encounted in practice, particularly in time-series settings. For forecasting applications, the two main methods for addressing quantile crossing are (1) use the fitted quantiles to fit some distribution for each

 $[\]overline{}^{3}$ Note that with this notation X includes a constant or intercept.

time period as in Adrian et al. (2019) or Korobilis (2017); or (2) sort the estimated quantiles in each period as proposed by Chernozhukov et al. (2009) and Chernozhukov et al. (2010). While these two-step methods yield proper densities, correcting the fitted quantiles does not quantify corresponding changes in the estimated coefficients, i.e., the coefficients estimated in the first step remain uncorrected. This prompted Bondell et al. (2010) to propose an estimator which yields no crossing for the estimated quantiles in-sample. We note that there are other means to estimate non-crossing quantiles like Liu and Wu (2009) who estimate the median first, and sequentially estimates further quantiles, conditional on the previously estimated quantiles not being crossed. While this method will yield non-crossing quantiles, the choice of the first estimated quantile can be arbitary. As such, in this paper we will focus exclusively on noncrossing constraints as done in Bondell et al. (2010) since elements of their constraints have been carried over to other quantile estimators (see Yang and Tokdar (2017) for example).

2.2Non-Crossing Constraints

Non-crossing constraints incorporated into quantile regression are a way to ensure that the estimated quantiles remain monotonically increasing. Imposing them can be done via inequality constraints:

$$\hat{\beta} = \underset{\beta}{\operatorname{argmin}} \sum_{q=1}^{Q} \sum_{t=1}^{T} \rho_q (y_t - x_t^T \beta_{\tau_q})$$

$$s.t. \ x^T \beta_{\tau_q} \ge x^T \beta_{\tau_{q-1}}$$

$$(2)$$

While conceptually simple, the number of constraints in equation (2) can be rather large on account of there being $T \times (Q-1)$ inequality constraints. To address this, Bondell et al. (2010) restrict the domain of the covariates to $\mathcal{D} \in [0,1]^K$ and focus on the worst case scenario in the data,⁴ which reduces the number of constraints to (Q-1).⁵ This simplifies computation a great deal and enables non-crossing constraints to be included without too much additional computational cost. Because quantile regression is invariant to monotone transformations, any affine invertible transformation can be applied: to obtain the coefficients pertaining to the untransformed data, it is enough to apply the inverse transformation to the estimated coefficients (Koenker, 2005).

⁴The situation where the negative difference coefficient's (γ_{j,τ_q}^-) corresponding variables values are 1 and the positive difference coefficients (γ_{j,τ_q}^+) corresponding variables equal 0. ⁵Any domain of interest which has an affine transformation that maps to $\mathcal{D} \in [0,1]^K$ works.

The method of Bondell et al. (2010) recasts the parameters in terms of quantile differences: $(\gamma_{0,\tau_1},...,\gamma_{K,\tau_1})^T = \beta_{\tau_1}$ and $(\gamma_{0,\tau_q},...,\gamma_{K,\tau_q})^T = \beta_{\tau_q} - \beta_{\tau_{q-1}}$ for q=2,...,Q. With this quantile difference reparametrisation, the constraint in equation (2) becomes $x^T\gamma_{\tau_q} \geq 0$. Note that we can without further assumptions decompose the j^{th} difference as $\gamma_{j,\tau_q} = \gamma_{j,\tau_q}^+ - \gamma_{j,\tau_q}^-$, where γ_{j,τ_q}^+ is its positive and $-\gamma_{j,\tau_q}^-$ its negative part. For each γ_{j,τ_q} both parts are non-negative and only one part is allowed to be non-zero. With this reparameterisation, along with the restriction of the data to $\mathcal{D} \in [0,1]^K$, the non-crossing constraint can be redefined as:

$$\gamma_{0,\tau_q} \ge \sum_{j=1}^{K} \gamma_{j,\tau_q}^- \ (q=2,...,Q)$$
(3)

A non-crossing constraint is therefore equivalent to ensuring that the sum of negative shifts do not push the quantile below the change in intercept, which acts as a pure location shifter. Bondell et al. (2010) shows that (3) is a necessary as well as a sufficient condition for non-crossing quantiles.

2.3 Adaptive Non-Crossing Constraints

Throughout the paper we follow the framework and assumptions laid out in Bondell et al. (2010). To derive adaptive non-crossing constraints we first need to recast the non-crossing constraints of equation (2) in a way that does not restrict the domain of interest to $\mathcal{D} \in [0, 1]^K$. This is provided by Lemma 1 below.

Lemma 1. Given that a non-crossing constraint looks at the sum of positive shifters to be larger than the sum of negative shifters, in the worst case scenario, these constraints can be reformulated as:

$$\gamma_{0,\tau_q} + \sum_{j=1}^{K} \min(X_j) \gamma_{j,\tau_q}^+ \ge \sum_{j=1}^{K} \max(X_j) \gamma_{j,\tau_q}^-$$
(4)

where X_j is the j^{th} variable in the design matrix.

Proof. Recall the standard non-crossing constraint formulation from Bondell et al. (2010) needs that in the worst case scenario the sum of positive shifters needs to be larger than the sum of negative shifters:

$$\gamma_{0,\tau_q} + \sum_{j=1}^{K} (Z_j = 0) \cdot \gamma_{j,\tau_q}^+ \ge \sum_{j=1}^{K} (Z_j = 1) \cdot \gamma_{j,\tau_q}^-$$

where $Z_j \in [0,1]$ is a transformed variable of X_j . Assume that $\max(X_j) > \min(X_j)$ for all $j \in \{1,2,...,K\}$, i.e., $Var(X_j) > 0$ for all j. Consider a design matrix variable X_j with domain $[\min(X_j), \max(X_j)]$. X_j can be normalised using the min-max transformation: $Z_{t,j} = \frac{X_{t,j}-\min(X_j)}{\max(X_j)-\min(X_j)}$, where Z_j is the normalised variable. We can express $X_j = \min(X_j) + Z_j \cdot [\max(X_j) - \min(X_j)]$. Instead of expressing the constraint in terms of the transformed variable Z_j , we can express it in terms of the untransformed X_j :

$$\gamma_{0,\tau_q} + \sum_{j=1}^{K} [\min(X_j) + (Z_j = 0) \cdot (\max(X_j) - \min(X_j))] \gamma_{j,\tau_q}^+$$

$$\geq \sum_{j=1}^{K} [\min(X_j) + (Z_j = 1) \cdot (\max(X_j) - \min(X_j))] \gamma_{j,\tau_q}^-$$

We can simplify this equation to recover:

$$\gamma_{0,\tau_q} + \sum_{j=1}^K \min(X_j) \gamma_{j,\tau_q}^+ \ge \sum_{j=1}^K \max(X_j) \gamma_{j,\tau_q}^-$$

Thus there is an equivalence between this equation and the one in Bondell et al. (2010). \Box

This new constraint is a sufficient condition for non-crossing, since if equation (4) is satisfied for the worst case, it is immediately satisfied for every observation. The novelty of equation (4) is that it works on the domain of $\mathbb{D} \in \mathbb{R}$, while the original formulation in Bondell et al. (2010), works for $\mathbb{D} \in [0, 1]$ only.

To study the impact of non-crossing constraints on the estimated coefficients, it is important to be able to tighten (and loosen) these constraints. As such, the next step is to formulate a set of constraints that can become adaptive as a hyperparameter α varies. An intuitively appealing formulation is one that yields the traditional quantile regression estimator when setting $\alpha = 0$ and the Bondell et al. (2010) when $\alpha = 1.6$ This leads to the following adaptive non-crossing constraints:

$$\gamma_{0,\tau_q} + \sum_{j=1}^{K} \left[\bar{X}_j - \alpha(\bar{X}_j - \min(X_j)) \right] \gamma_{j,\tau_q}^+ \ge \sum_{j=1}^{K} \left[\bar{X}_j + \alpha(\max(X_j) - \bar{X}_j) \right] \gamma_{j,\tau_q}^- \tag{5}$$

When $\alpha = 0$, the constraint simplifies to imposing non-crossing quantiles evaluated at the average value of the covariates. Quantile monotonicity at this value is satisfied even by the traditional quantile regression estimator: $\bar{X}^T\beta$ yields the empirical quantiles of Y, which are

⁶Technically, any scalar $\alpha > 0$ would work. We set $\alpha = 1$ to recover the Bondell et al. (2010) for simplicity

monotonically increasing by definition (Koenker, 2005; Koenker and Xiao, 2006). As such imposing equation (5) and setting $\alpha = 0$ will yield the same β coefficients as the traditional quantile regression without constraints.

The constraints in equation (5) equal the non-crossing constraints when $\alpha = 1$ as the equation becomes equation (4). As such using equation (5) as a constraint allows us to recover both the traditional quantile regression estimator of Koenker and Bassett (1978), as well as the non-crossing quantile regression estimator of Bondell et al. (2010).

There are other ways to construct adaptive non-crossing constraints that yield traditional quantile regression as well as Bondell et al. (2010) estimator, specifically through the use of indicator functions that activate the non-crossing constraint. The advantage of equation (5) is that it produces a gradual transition from quantile estimates to non-crossing estimates as α increases. This allows us to study the impacts these constraints have on the estimated coefficients.

Theorem 1. Non-crossing constraints are a type of Fused LASSO, with quantile specific hyperparameters: $k_{\tau_q}^* = \frac{\gamma_{0,\tau_q}}{\alpha}$.

Proof. For simplicity, assume Q=2. We begin with the non-crossing constraint given by equation (5):

$$\gamma_0 \ge \sum_{j=1}^{K} \left[[\bar{X}_j + \alpha(\max(X_j) - \bar{X}_j)] \gamma_j^- - [\bar{X}_j - \alpha(\bar{X}_j - \min(X_j))] \gamma_j^+ \right]$$

where $\gamma_j^- \geq 0$ and $\gamma_j^+ \geq 0$ represent the positive and negative parts of γ_j such that $\gamma_j = \gamma_j^+ - \gamma_j^-$. By Lemma 1, we can rescale the data to $\mathbb{D} \in [-1,1]$. When data is properly standardized and rescaled, we can assume $\bar{X}_j = 0$ for each covariate j. For a symmetric distribution, this standardization is trivial; for asymmetric distributions, one can normalise the data prior to rescaling. After normalising and rescaling, $\bar{X}_j \approx 0$, $max(X_j) = 1$, and $min(X_j) = -1$. With this, the constraint simplifies to:

$$\gamma_0 \ge \sum_{j=1}^K \alpha(\gamma_j^- + \gamma_j^+)$$

Since $\gamma_j^- \ge 0$ and $\gamma_j^+ \ge 0$ by definition, we have $(\gamma_j^- + \gamma_j^+) = |\gamma_j^- + \gamma_j^+|$. Rearranging terms yields:

$$\frac{\gamma_0}{\alpha} \ge \sum_{j=1}^K |\gamma_j^+ + \gamma_j^-|$$

This corresponds exactly to the fused shrinkage formulation in Jiang et al. (2013) with $k^* = \frac{\gamma_0}{\alpha}$.

For Q>2, all γ parameters become quantile specific, leading to quantile specific hyperparameters: $k_{\tau_q}^* = \frac{\gamma_{0,\tau_q}}{\alpha}$. As such, non-crossing constraints lead to fused LASSO shrinkage of the parameters, with quantile specific hyperparameters. This equivalence requires $\alpha>0$, as the relationship is undefined when $\alpha=0$. As $\alpha\to0$, $k_{\tau_q}^*\to\infty$, and we recover the QR coefficients for sufficiently small α .

The results of theorem (1) mean that when α is large enough, the only way to satisfy the constraint is by having $\gamma_{j,\tau_q}^- = \gamma_{j,\tau_q}^+ = 0$, which is equivalent to the Composite Quantile Regression setup of Koenker (1984) and Zou and Yuan (2008).⁷ Because it is possible to recover the traditional quantile regression estimator, the Bondell et al. (2010) estimator, and the composite quantile regression estimator by simply varying α , we refer to the estimator using the constraints of equation (5) as Generalised Non-Crossing Quantile Regression (GNCQR).^{8,9} Formally GNCQR is defined as:

$$\hat{\beta}_{GNCQR}|\alpha = \underset{\beta}{argmin} \sum_{q=1}^{Q} \sum_{t=1}^{T} \rho_{q}(y_{t} - x_{t}^{T}\beta_{\tau_{q}})$$

$$s.t. \ \gamma_{0,\tau_{q}} + \sum_{k=1}^{K} \left[\bar{X}_{k} - \alpha(\bar{X}_{k} - min(X_{k})) \right] \gamma_{k,\tau_{q}}^{+} \geq$$

$$\sum_{k=1}^{K} \left[\bar{X}_{k} + \alpha(max(X_{k}) - \bar{X}_{k}) \right] \gamma_{k,\tau_{q}}^{-}$$

$$(6)$$

Equation (6) makes it clear that α plays a strong role in the estimator. Given that alpha plays a similar role to λ in LASSO, we restrict its possible values to be $\alpha \geq 0$. One might be tempted to restrict potential value of $\alpha \geq 1$, i.e. always enforcing non-crossing quantiles in estimation. We caution against this, because the estimated conditional quantiles are merely an approximation of the true (and unknown) DGP (Koenker and Xiao, 2006). In a situation where we have model misspecification, enforcing non-crossing constraints lead to too much interquantile shrinkage as a consequence of theorem (1).

The advantage of the GNCQR estimator is that it leads to quantile specific shrinkage parameters, while only needing to set the scalar α . In essence, the problem becomes one of model

⁷Note that in the limit there is no unique solution. Because of this, as $\alpha \to \infty$ it becomes numerically challenging to find the solution, leading to numerical instability.

⁸The different types of estimators are defined, with a unified notation, in the appendix.

⁹We follow Powell (2020) in the naming convention, who develop the 'Generalized Quantile Regression'.

selection, but instead of selecting variables from X, the interest is on interquantile shrinkage, i.e. selecting variables whose impact on Y captured by β varies by quantile.

2.4 Hyperparameter selection

A natural candidate for choosing the degree of penalisation is cross-validation. In cross-section applications, this would be the usual k-fold cross-validation or a variant therefore. Of the several types of cross-validation methods available for time series application, we choose the hv-block CV setup of Racine (2000). This block setup has several desirable properties: (1) it has been shown to have a good trade-off between bias and variance in various applications; (2) the required number of computations does not increase with the number of observations to the degree it would with leave-one-out cross-validation, 10 and; (3) when the "h" in the hv-block is set equal to 0, we recover the standard k-fold cross-validation setup, so the method can be used for both cross-section and time-series data. 11

Figure (1) visualises this block setup and which data are removed from the dataset. The reason to remove the data points around the test datasets in time-series data is to ensure that no data leakage occurs when evaluating the performance of the model. When data are dependent, the information from the test dataset can leak into the training set. This can inadvertently lead to overfitting and hence poor generalization. Removing the data around the test dataset can guard against this.¹²

Since the hyperparameter α is a scalar, one can find the optimal hyperparameter using simple grid-search. Hence, the curse of dimensionality that often limits the applicability of grid-search is not present here. Crucially, the grid-search is 'embarrassingly parallel' and this can be utilised to speed up computation times (Bergstra and Bengio, 2012). As such, throughout the paper we will utilise grid-search to obtain the optimal hyperparameter values.

¹⁰See Cerqueira et al. (2020) for a description and comparison of the different types of cross-validation for time-series data

¹¹We use this equivalence in the Monte Carlo setups and applications so that the same cross-validation code can be employed throughout.

¹²For more discussion on hyperparameter selection please see the appendix.

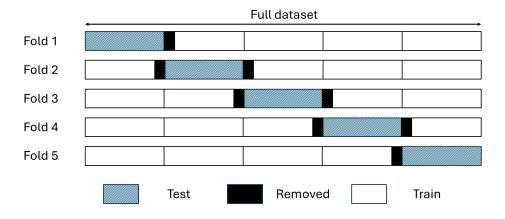


Figure 1: hv-Cross Validation

3 Monte Carlo

3.1 Setup

Theorem 1 demonstrated the connection between Fused LASSO and non-crossing constraints. This theorem also shows that GNCQR is equivalent to having quantile-specific hyperparameters. In this section we explore the implications of this using Monte Carlo evidence, and evaluate variable selection properties of the various estimators as well as examine their ability to recover the true quantiles. Specifically, we consider the following estimators: the non-crossing quantile regression of Bondell et al. (2010) (BRW), GNCQR, and the Fused LASSO (FLQR).¹³ We will also look at the two-step approach proposed in Chernozhukov et al. (2010), i.e. running regular quantile regression and sorting the fitted quantiles. This will allow us to gauge how much better model selection results lead to improved fit.

Our Monte Carlo setup takes as its starting point the design used in Bondell et al. (2010). Each Monte Carlo experiment is generated using the location scale heteroskedastic error model of the form:

$$y_t = \beta_0 + \beta^T x_t + (\theta_0 + \vartheta_t \odot \theta^T x_t) \varepsilon_t, \ x_{t,k} \sim U(0,1), \ \varepsilon_t \sim N(0,1)$$
 (7)

An intercept is included in each setup (i.e., $\beta_0 = \theta_0 = 1$). Note the k^{th} element of $\vartheta_t \in \{0, 1\}$ which regulates whether the given variable has quantile variation at the given quantile. This term is included to allow 'quantile varying sparsity', i.e. cases where a certain variable enters

¹³Note that we consider the Fused LASSO and not the Fused Adaptive LASSO in Jiang et al. (2013). We focus on the non-adaptive version as this allows us to examine the value added impact of quantile-specific hyperparameters.

only parts of the distribution (Kohns and Szendrei, 2021). A simple way to implement quantile-specific sparsity is by setting ϑ_t as an indicator function, where it takes the value of 1 only for cases when ε_t is below (or above) a specific quantile. For cases where there is full quantile variation, $\vartheta_t = \mathbf{1}_k$. The t subscript is needed the presence of quantile variation will be dependent on the magnitude of ε_t .

There are four data generating processes (DGPs), each with 500 generated datasets. The first three $(y_1, y_2, \text{ and } y_3)$ are identical to Examples 1-3 of Bondell et al. (2010). The fourth DGP (y_4) is an amendment of Example 2, with some variables only varying at the tails. Specifically:

- y_1 : k = 4, with the parameters $\beta = \mathbf{1}_k$, $\theta = 0.1 * \mathbf{1}_k$, and $\theta = \mathbf{1}_k$.
- y_2 : k = 10, with the parameters $\beta = (\mathbf{1}_4^T, \mathbf{0}_6^T)^T$, $\theta = (0.1 * \mathbf{1}_4^T, \mathbf{0}_6^T)^T$, and $\theta = \mathbf{1}_k$.
- y_3 : k = 7, with the parameters $\beta = \mathbf{1}_k$, $\theta = (\mathbf{1}_3^T, \mathbf{0}_4^T)^T$, and $\theta = \mathbf{1}_k$.
- y_4 : k = 10, with the parameters $\beta = (\mathbf{1}_4^T, \mathbf{0}_6^T)^T$, $\theta = (0.1 * \mathbf{1}_8^T, \mathbf{0}_2^T)^T$, and $\theta = (\mathbf{1}_4^T, \mathbf{1}_4^T \times [I(\epsilon_t > F_{\varepsilon}^{-1}(0.9)) + I(\epsilon_t \leq F_{\varepsilon}^{-1}(0.1))], \mathbf{0}_2^T)^T$.

Note that for y_4 , variable selection and fused shrinkage will both be necessary. The models considered will only allow for fused shrinkage, and as such this DGP is only included to judge the performance of the estimators in unideal situations.

To test the performance of the different estimators three sample sizes are considered for each DGP: $T \in \{50, 100, 200\}$. Sample sizes 100 and 200 were also considered in Bondell et al. (2010), but 50 was not. We include this small sample setting because in macroeconometric applications it is not uncommon to apply quantile regression for such small samples.¹⁴ Applying quantile regression to analyse GDP distribution for European economies has been particularly daunting on account of data availability, so we feel that these Monte Carlo results are particularly useful for policy makers.

We also consider variation in the number of quantiles to be estimated, by generating equidistant quantiles with varying distance between the quantiles: $\Delta_{\tau} \in \{0.2, 0.1\}^{15}$ Note that by increasing the number of quantiles, the number of estimated parameters varies. The choice to vary the estimated quantiles was driven by the fact that Theorem (1) shows how γ_0 acts as

¹⁴See Szendrei and Varga (2023) for an application with around 50 observations, Figueres and Jarociński (2020) for an application with around 100 observations, and Adrian et al. (2019) for an application with around 200 observations.

 $^{^{15}}$ for $\Delta_{\tau} = 0.2$, the first quantile is set to 0.1.

a quantile-specific hyperparameter, which is a claim we can verify by examining the variable selection properties of BRW as the number of quantiles increases.

Two measures are used to judge the variable selection performance of the estimators: True Positive Rate (TPR) and True Negative Rate (TNR). Both measures take a value between 0 (worst performance possible) and 1 (best performance possible). TPR measures the degree to which the estimator is able to capture the quantile varying coefficients, while the TNR measures the ability of the estimator to identify where no quantile variation occurs. Considering these measures together allows one to conclude whether a given estimator over-shrinks or undershrinks. When calculating these measures, we only consider the difference in β coefficients of the variables (without the intercept).

To measure fit, we follow Bondell et al. (2010) and report the average root mean integrated square error ($\times 100$):

$$RMISE = \frac{1}{n} \left[\sum_{iter=1}^{n} \left(\hat{g}_{\tau}(x_{iter}) - g_{\tau}(x_{iter}) \right)^{2} \right]^{1/2}$$

where *iter* indexes a given Monte Carlo experiment, n = 500 is the total number of evaluated Monte Carlo experiments for each DGP, \hat{g}_{τ} is the estimated quantile and g_{τ} is the true quantile given equation (7). We also report the standard error of the RMISE for all the estimators.

We create a grid of 100 equally-spaced points between [0,1), and use grid-search to obtain the optimal α parameter. To consider large α options, we append to this list of candidate solutions 200 points between [0,6] as exponents of base 10. Setting the hyperparameter to 10^6 yields solutions that are close to the composite quantile regression solution. For FLQR and GNCQR, we use 10-fold cross-validation.

Note that while Theorem (1) implies that changing the α parameter will have a similar impact on the β 's as the λ of a Fused LASSO estimator, the same α and λ values will not lead to the same β coefficients. This is because the quantile-specific difference in constants is an upper limit of quantile variation for GNCQR.

Table (2) presents the results for the average fit of the different estimators based on the Monte Carlo experiments, while Table 1 shows the variable selection performance. The RIMSE set of results for y_1 , y_2 , and y_3 when T=100 (and T=200 for y_3) and $\Delta \tau = 0.2$ is the closest setup to Bondell et al. (2010). The results for these setups for BRW and QR are almost identical and as such our new findings relate to the extensions to the Monte Carlo experiments of Bondell et al. (2010).

3.2 Variable Selection results

The results on variable selection presented in Table 1 are more revealing. Since the GNCQR can recover both the BRW (when setting $\alpha = 1$) and the QR (as $\alpha \to 0$), we can compare the performance of these estimators to see the impact α has on the coefficient profiles. Given the findings of Theorem 1, we expect the TPR to decrease and the TNR to increase as α increases. We find that the TPR of BRW is below 1 (and the TNR is above 0) for all DGPs, which corroborates Theorem 1, which states that non-crossing constraints are a special type of fused shrinkage. This is not simply a feature of the Monte Carlo design, as the QR yields a TPR of 1 and a TNR of 0 in all cases. Note that for y_1 , all variables are quantile-varying and as such TNR does not exist for this case.

Among the estimators that have some fused shrinkage, BRW yields the highest TPR for all DGPs for all cases considered. GNCQR always ranks second and FLQR has the worst performance for TPR. However, this superior performance in TPR for BRW is coupled with the worst performance when it comes to TNR. For TNR, FLQR produces the best results, with GNCQR a close second. From these results we can see that BRW undershrinks quantile variation, FLQR overshrinks, and GNCQR yields a middle-ground option.

As will be seen in the fit results of Table (2), we find that GNCQR not only provides robustness over FLQR for y_4 , but is also able to give fits close to FLQR while retaining better model selection properties. In particular, GNCQR is able to yield much better TPR than FLQR without substantially compromising its ability to identify the true negative differences.

Increasing the number of quantiles has a marked impact on variable selection: it lowers TPR and increases TNR for all estimators (except the QR). This further corroborates Theorem 1. Increasing sample size also influences the TPR and TNR of all estimators. For BRW and GNCQR, increasing the sample size increases TPR but lowers TNR for all DGPs and all Δ_{τ} . However, for FLQR we see similar tendency when $\Delta_{\tau} = 0.2$ but not when $\Delta_{\tau} = 0.1$. In particular, for FLQR for y_4 , the TPR decreases as the sample size increases when $\Delta_{\tau} = 0.1$. This is particularly troubling since FLQR has the worst TPR of all the estimators.

3.3 Fit results

Results on fit in Table (2) reveal that GNCQR and FLQR provide best fits, even beating BRW estimator, for y_1 , y_2 , and y_3 . However, for y_4 FLQR fails to yield improvements over BRW. While GNCQR also has difficulties in y_4 , it is much closer to the fits of BRW, with both

Table 1: True Positive and True Negative Rates for the different Monte Carlo experiments

	y_1 y_2		y_3		y_4		y_1	y_2		y_3		y_4		
	TPR	TPR	TNR	TPR	TNR	TPR	TNR	TPR	TPR	TNR	TPR	TNR	TPR	TNR
	$\Delta \tau = 0.2$				$\Delta \tau = 0.1$									
T-50														
$_{\mathrm{BRW}}$	0.806	0.360	0.639	0.535	0.525	0.305	0.691	0.618	0.246	0.752	0.383	0.669	0.224	0.776
GNCQR	0.372	0.232	0.769	0.345	0.715	0.243	0.753	0.304	0.195	0.812	0.271	0.788	0.173	0.825
QR	1.000	1.000	0.000	1.000	0.000	1.000	0.000	1.000	1.000	0.000	1.000	0.000	1.000	0.000
FLQR	0.215	0.136	0.868	0.246	0.790	0.194	0.819	0.127	0.107	0.901	0.154	0.870	0.114	0.888
T=100														
$_{\mathrm{BRW}}$	0.938	0.510	0.501	0.694	0.393	0.390	0.582	0.792	0.363	0.641	0.531	0.557	0.318	0.687
GNCQR	0.364	0.228	0.780	0.430	0.689	0.279	0.715	0.294	0.183	0.828	0.349	0.767	0.179	0.833
QR	1.000	1.000	0.000	1.000	0.000	1.000	0.000	1.000	1.000	0.000	1.000	0.000	1.000	0.000
FLQR	0.202	0.122	0.885	0.299	0.776	0.184	0.834	0.110	0.074	0.927	0.196	0.860	0.078	0.932
T=200														
$_{\mathrm{BRW}}$	0.988	0.659	0.340	0.846	0.262	0.481	0.459	0.923	0.488	0.517	0.690	0.445	0.409	0.575
GNCQR	0.381	0.225	0.787	0.564	0.672	0.280	0.700	0.325	0.166	0.846	0.465	0.765	0.203	0.797
QR	1.000	1.000	0.000	1.000	0.000	1.000	0.000	1.000	1.000	0.000	1.000	0.000	1.000	0.000
FLQR	0.215	0.130	0.883	0.413	0.746	0.209	0.821	0.139	0.061	0.946	0.298	0.832	0.098	0.917

yielding better fits than the traditional QR or FLQR. This is because GNCQR recovers BRW when $\alpha = 1$, so it will not do much worse than BRW. Hence, GNCQR is more robust to DGPs that have quantile-specific sparsity than the simple FLQR.

Unsurprisingly, y_4 produces the worst fit for all estimators, but as more data becomes available, the performance of all the estimators initially improves. The key difference lies with FLQR, where the fits do not improve as much as the other estimators when the sample size increases from T = 100 to T = 200. This also highlights that to yield improvements in the traditional LASSO setting for such DGPs one needs to explicitly penalise the level of β_{τ} coefficients too; see also Jiang et al. (2014). Overall, increasing sample size improves the fit for all estimators, while increasing the fineness of the grid of quantiles yields no significant differences. The key takeaway is that GNCQR either gives further improvements in fit over BRW, or (at worst) does as well as BRW.

Looking at the rearrangement method proposed by Chernozhukov et al. (2010) reveals that sorting the fitted quantiles for the quantile regression estimator yields modest improvements. These improvements increase as the number of estimated quantiles increases especially for smaller sample sizes. This is intuitive, as with more quantiles there is a higher chance of quantile crossing. However, the improvements in fit from rearrangement get smaller as the sample size increases.

Table 2: RIMSE of different models for the different Monte Carlo experiments

0.9 std. err	0.42 0.39 0.44 0.41	0.37 0.37 0.45 0.41	0.83 0.80 0.91 0.86	2.34 3.33 1.52 3.11	0.42 0.37 0.44 0.38	0.37 0.35 0.45 0.38	0.81 0.80 0.91 0.85	2.45 3.54 1.52 3.41
0.9 bias std	30.4 24.3 31.0 24.9	40.7 33.8 47.2 34.7	74.0 74.4 81.1 74.5	180.5 186.0 201.6 192.2	29.7 23.7 31.0 24.0	39.9 33.3 47.2 33.3	73.0 73.1 81.1 74.4	179.4 182.9 201.6 190.0
err	0.33 0.30 0.33 0.30	0.32 0.30 0.36 0.31	0.65 0.65 0.70 0.66	0.38 0.39 0.37 0.38	0.32 0.29 0.33 0.30	0.32 0.30 0.36 0.30	0.67 0.65 0.70 0.65	0.36 0.37 0.37 0.36
0.7 bias std.	23.2 20.5 23.5 23.5	33.2 30.8 36.7 30.9	57.6 56.9 61.8 58.6	37.4 36.0 36.8 35.1	22.9 20.3 23.5 20.3	33.0 30.5 36.7 30.4	57.3 56.2 61.8 57.8	35.4 33.4 36.8 33.0
0 err	0.32 0.29 0.33 0.30	0.30 0.29 0.33 0.30	0.62 0.62 0.65 0.65	0.36 0.37 0.35 0.35	0.33 0.29 0.33 0.29	0.30 0.29 0.33 0.29	0.62 0.61 0.65 0.62	0.34 0.35 0.35 0.35
T	22.1 20.0 22.4 20.1	31.7 29.9 34.6 30.1	54.8 52.9 58.1 53.8	33.6 34.1 34.2 34.0	21.8 19.9 22.4 20.0	31.6 29.7 34.6 29.6	54.1 52.7 58.1 53.4	32.5 32.3 34.2 32.1
0.3 std. err	0.33 0.30 0.33 0.29	0.32 0.31 0.35 0.35	0.65 0.63 0.71 0.64	0.40 0.40 0.41 0.40	0.33 0.29 0.33 0.29	0.32 0.30 0.35 0.31	0.64 0.63 0.71 0.64	0.41 0.39 0.41 0.37
bias	23.5 20.9 23.8 21.0	33.1 30.6 36.5 30.6	58.8 57.2 62.8 57.4	37.4 36.1 36.7 35.4	23.3 20.9 23.8 20.9	32.9 30.3 36.5 30.3	57.9 56.8 62.8 57.4	35.6 33.7 36.7 33.1
0.1 std. err	0.45 0.38 0.46 0.39	0.39 0.38 0.47 0.38	0.83 0.83 0.96 0.85	2.32 3.28 1.50 3.39	0.44 0.37 0.46 0.39	0.38 0.36 0.47 0.35	0.82 0.80 0.96 0.83	2.43 3.51 1.50 3.64
bias	30.4 23.9 30.8 24.1	40.8 34.3 47.6 33.8	75.7 75.2 83.1 77.9	176.9 181.6 199.5 195.3	29.7 24.0 30.8 24.2	40.0 33.6 47.6 33.1	73.9 74.4 83.1 78.2	175.4 179.8 199.5 190.5
0.9 std. err	0.61 0.57 0.66 0.59	0.52 0.53 0.66 0.55	1.19 1.16 1.37 1.13	2.22 3.06 1.44 3.06	0.59 0.54 0.66 0.56	0.52 0.54 0.66 0.55	1.19 1.14 1.37 1.14	2.30 3.33 1.44 3.36
bias	41.1 33.6 43.5 34.2	54.4 48.5 67.9 49.5	$100.0 \\ 98.6 \\ 115.2 \\ 100.1$	197.5 199.8 232.2 205.7	40.0 32.7 43.5 33.2	53.6 48.6 67.9 48.8	98.0 97.3 115.2 98.0	196.9 194.5 232.2 199.9
0.7 std. err	0.49 0.45 0.49 0.45	0.42 0.43 0.49 0.43	0.91 0.92 0.97 0.92	0.63 0.66 0.75 0.65	0.49 0.44 0.49 0.44	0.43 0.43 0.49 0.43	0.89 0.90 0.97 0.90	0.64 0.62 0.75 0.62
0 err bias s	32.0 29.3 33.0 29.3	46.4 44.0 52.2 44.0	80.8 80.0 89.3 81.7	52.9 51.9 54.7 51.2	31.6 28.9 33.0 29.1	46.1 43.9 52.2 43.9	80.5 78.2 89.3 79.4	51.1 49.0 54.7 48.6
T=100 0.5 s std. err	0.45 0.42 0.47 0.44	0.43 0.42 0.47 0.42	0.83 0.83 0.91 0.86	$\begin{array}{c} 0.55 \\ 0.59 \\ 0.51 \\ 0.61 \end{array}$	0.44 0.42 0.47 0.47	0.43 0.43 0.47 0.42	0.82 0.82 0.91 0.84	0.53 0.56 0.51 0.57
bia	30.3 27.8 31.2 27.9	44.6 43.3 49.3 43.3	76.2 74.7 82.7 75.8	48.4 48.6 48.7 49.2	29.8 27.7 31.2 27.8	44.6 43.2 49.3 43.1	76.2 74.1 82.7 74.6	47.0 47.1 48.7 47.3
0.3 std. err	0.47 0.44 0.50 0.45	0.44 0.42 0.47 0.43	0.94 0.92 1.01 0.92	0.66 0.68 0.81 0.67	0.46 0.45 0.50 0.44	0.43 0.42 0.47 0.43	0.91 0.90 1.01 0.96	0.64 0.63 0.81 0.60
err bias	32.9 29.2 33.9 29.5	45.9 43.8 51.9 43.6	79.7 78.0 87.8 78.3	53.1 52.2 54.5 51.9	32.5 29.2 33.9 29.0	45.9 43.8 51.9 43.9	79.6 77.8 87.8 78.6	50.9 49.1 54.5 48.7
0.1 std. er	0.58 0.54 0.63 0.55	0.51 0.53 0.65 0.54	1.19 1.22 1.37 1.18	2.28 3.18 1.42 3.55	0.57 0.53 0.63 0.53	0.50 0.53 0.65 0.53	1.18 1.20 1.37 1.19	2.37 3.53 1.42 3.75
err bias	41.7 33.6 43.9 33.2	55.0 48.7 67.1 47.9	101.3 98.6 116.8 99.9	194.4 196.5 231.0 200.9	40.6 33.5 43.9 32.5	54.3 48.8 67.1 48.0	100.1 97.5 116.8 100.6	193.5 190.6 231.0 194.0
0.9 std. err	0.80 0.78 0.93 0.77	0.72 0.76 0.90 0.78	1.70 1.69 1.95 1.67	2.29 2.97 1.78 2.91	0.78 0.76 0.93 0.81	0.71 0.78 0.90 0.79	1.67 1.65 1.95 1.64	2.36 3.22 1.78 3.27
	56.5 48.2 62.4 48.2	73.2 68.3 94.8 69.3	137.6 133.9 165.7 136.4	224.8 222.9 282.4 228.6	55.6 47.6 62.4 48.1	72.3 69.1 94.8 69.6	137.3 133.3 165.7 135.2	223.2 217.1 282.4 222.7
0.7 std. err bias	0.66 0.63 0.71 0.65	0.62 0.63 0.73 0.62	1.36 1.32 1.49 1.37	1.66 1.78 2.32 1.84	0.66 0.64 0.71 0.64	0.61 0.63 0.73 0.63	1.36 1.32 1.49 1.34	1.67 1.78 2.32 1.80
-	44.8 41.9 48.0 42.1	63.1 62.0 72.3 61.7	114.2 112.5 127.8 113.5	85.5 85.4 98.9 85.8	44.8 41.7 48.0 42.0	63.2 62.5 72.3 62.4	113.6 112.1 127.8 113.5	84.1 82.9 98.9 82.7
T=50 0.5 std. err bias	0.62 0.61 0.65 0.65	0.57 0.58 0.67 0.59	1.28 1.27 1.39 1.29	1.27 1.36 1.09 1.38	0.62 0.61 0.65 0.60	0.58 0.59 0.67 0.60	1.25 1.27 1.39 1.30	1.26 1.29 1.09 1.30
T=50 0.5 bias std.	42.8 40.8 45.3 40.6	61.2 60.1 67.6 59.9	108.3 106.9 118.2 106.9	77.3 77.8 72.5 79.0	42.8 40.7 45.3	61.4 60.6 67.6 60.3	108.0 106.5 118.2 106.7	75.5 75.4 72.5 75.5
0.3 std. err bias	0.65 0.61 0.68 0.61	0.58 0.60 0.70 0.61	1.28 1.30 1.38 1.30	1.61 1.66 2.34 1.66	0.64 0.61 0.68 0.61	0.59 0.61 0.70 0.63	1.26 1.25 1.38 1.29	1.63 1.64 2.34 1.84
	44.3 41.7 46.8 41.8	63.0 61.9 72.2 61.8	110.4 109.4 124.1 109.7	85.9 85.5 100.0 85.7	44.0 41.6 46.8 41.5	63.1 62.4 72.2 62.3	110.6 109.7 124.1 110.9	84.2 82.6 100.0 84.5
0.1 std. err bias	0.76 0.72 0.84 0.72	0.69 (0.75 (0.84 70.74 (1.62 1 1.71 1 2.00 1 1.74 1	3.06 8 3.06 8 1.74 1 3.40 8	0.76 0.72 0.84 0.72	0.69 0.74 0.84	1.62 1 1.67 1 2.00 1 1.73 1	2.52 8 3.33 8 1.74 1 3.60 8
0.1 bias ste	55.3 (46.5 (60.7 (46.6 (73.4 (69.4 (94.5 (68.5 (54.3 (46.2 (60.7 (46.4 (73.2 (69.9 (94.5 (
τ Di	$\begin{array}{c c} \Delta \tau = 0.2 \\ & \text{BRW} \\ \text{GNCQR} \\ & \text{QR} \\ \text{GR} \\ & \text{FLQR} \\ \end{array}$	BRW 7 GNCQR 6 QR 9 FLQR 6	^{l3} BRW 136.8 GNCQR 132.5 QR 167.7 FLQR 133.7	- ~ ~ ~ <u>-</u>	_ ~ ~ ~ ~	BRW 7 GNCQR 6 QR 9 FLQR 6	³ BRW 135.2 GNCQR 130.9 QR 167.7 FLQR 132.6	H BRW 224.3 GNCQR 222.0 QR 279.7 FLQR 226.2
	$S_1 = S_1$ S_1 S_2 S_3 S_4 S_1 S_2 S_3 S_4	£ _	y ₃ GNC	gNG GNG	L 5 H	5 7	Z H	garanta da

Comparing the rearrangement method with the alternative estimators shows that although rearrangement helps improve fit, the alternative estimators still perform better than the sorted QR. This highlights how better interquantile variable selection translates to superior fit beyond simple quantile sorting.

Note that in all the Monte Carlo runs, we have estimated a correctly specified model. In such situations the BRW estimator will (almost) always yield improvements over the QR. When the estimated model is misspecified, imposing strict non-crossing constraints can lead to worse coefficient bias than the QR. The fact that the GNCQR allows for some quantile crossing when $\alpha < 1$, means that the estimator is more robust to misspecification than the BRW. In essence, when $0 < \alpha_{opt} < 1$ the GNCQR provides the best linear approximation of the quantiles while regularising some of the quantile variation in the coefficients.

In summary, GNCQR and FLQR provides better fit than BRW for y_1 , y_2 , and y_3 , while for y_4 BRW performs best but GNCQR is nearly as good and both are preferred over the FLQR. This robustness of GNCQR is attractive in quantile applications as it is difficult to know ex ante whether quantile specific sparsity is present in the DGP. The variable selection results demonstrate that BRW undershrinks while FLQR overshrinks. Taking the fit and variable selection results together, we can see that GNCQR provides good fit while retaining better variable selection properties than FLQR.

4 Growth-at-Risk results

The canonical GaR uses US quarterly GDP growth in conjunction with the NFCI to obtain estimates of downside risk. Using quantile regression, GaR estimates can be obtained by estimating the following model:

$$y_{t+h} = x_t' \beta_{\tau_a} + \varepsilon_{t+h} \tag{8}$$

for $t = 1, \dots, T-h$, where h refers to the forecast horizon. x_t includes a constant (intercept), current quarterly GDP growth (annualised), and the average of the NFCI for the given quarter. The quarterly data cover 1973Q1 to 2023Q1. This sample includes the original sample of Adrian et al. (2019), augmented by the COVID-19 crisis. We consider one- and four-quarter ahead forecast horizons (h = 1, 4). We follow Adrian et al. (2019), and forecast average growth rates over the specific horizons.

To evaluate the performance of the different estimators, we will first conduct a small psuedo forecasting exercise similar to Carriero et al. (2025). This is followed by looking at the coefficient

profiles of the different models to see whether we corroborate the finding of Adrian et al. (2019), i.e. that lagged GDP growth acts as a location shifter, and NFCI is the key driver of nonlinearities in GDP growth. We will briefly look at the in-sample and out-of-sample weighted residuals for several values of α to gauge the bias-variance trade-off. We then turn to the policy implications of selecting models that incorporate interquantile shrinkage.

4.1 Out-of-Sample Performance

To evaluate out-of-sample performance of the estimators we will do a pseudo forecasting exercise similar to the one in Carriero et al. (2025). For the one- and four-quarter ahead forecast horizon we will compute the out-of-sample GDP densities on an expanding window, where the initial in-sample period uses the first 50 observations of the sample. This means that there are in total 150 - h forecast windows to evaluate.

4.1.1 Crossing incidence

Before evaluating the out-of-sample performance, we note that having $\alpha=1$ only ensures non-crossing quantiles in-sample but not out-of-sample. However, when setting $\alpha>1$, the non-crossing constraints are tighter which leads to shrinking the fitted quantiles towards parallel lines. This in turn leads to less out-of-sample crossing as well. Since the FLQR also shrinks towards parallel fitted quantiles, one question is whether the FLQR also leads to less out-of-sample crossing. To this end we will compare the out-of-sample crossing incidence for all the estimators (QR, BRQ, GNCQR, FLQR). The crossing incidence is calculated by comparing the fitted quantiles with the sorted quantiles following the procedure of Chernozhukov et al. (2010):

$$CrossI = \frac{1}{Q(T-50)} \sum_{t=51}^{T} \sum_{q=1}^{Q} \mathbf{I}[\hat{\mathcal{Q}}(\tau_q, t+h|\mathcal{F}_t) = \hat{\mathcal{Q}}_{sort}(\tau_q, t+h|\mathcal{F}_t)]$$
(9)

where $\hat{Q}(\tau_q, t + h|\mathcal{F}_t)$ is the forecasted quantile at time t and forecast horizon h and $\hat{Q}_{sort}(\tau_q, t + h|\mathcal{F}_t)$ is the sorted forecast quantile. The crossing incidence measures what proportion of quantiles need sorting after estimation to obtain a valid CDF. The lower the CrossI value the less quantiles need to be rearranged after estimation. The out-of-sample crossing incidence table is presented in table (3)

The first thing to note is that more crossing occurs at the shorter forecast horizon than the

Table 3: Out-of-Sample Crossing Incidence for the different estimators

	h=1	h=4
\overline{QR}	6.32%	5.62%
BRW	1.37%	1.47%
FLQR	0.91%	0.00%
GNCQR	3.19%	1.79%

longer one for all estimators. Second, the out-of-sample crossing incidence is the highest for the traditional quantile regression method. This indicates that the rearrangement algorithm of Chernozhukov et al. (2010) would have the most gains for the traditional quantile estimator. Note, that as mentioned by Bondell et al. (2010), post-processing methods like the rearrangement method have the potential to improve fitted quantiles without changing the estimated coefficients. Third, GNCQR yields the lowest out-of-sample crossing incidence, even yielding proper densities for h=4. Finally, the FLQR estimator leads to worse out-of-sample crossing than the BRW estimator. Taken together these results indicate that the GNCQR is most likely to yield proper forecasted densities.

4.1.2 Forecast Performance

Given that the sorting procedure is widely used in forecasting settings, we will evaluate forecast performance of sorted and unsorted quantiles. We opt to look at both sorted and unsorted quantiles because of the results in table (3) showing that sorting is likely to improve the performance of the QR and FLQR more than the BRW and GNCQR. To evaluate forecast performance, several measures are used. First, the quantile score (QS) is used with is the tick-loss weighted residual for a given forecast observation (Giacomini and Komunjer, 2005):

$$QS_{\tau_q,t+h} = (y_{t+h} - \hat{\mathcal{Q}}(\tau_q, t+h|\mathcal{F}_t))(\tau_q - \mathbf{I}[|y_{t+h} \le \hat{\mathcal{Q}}(\tau_q, t+h|\mathcal{F}_t)|])$$
(10)

where $\hat{\mathcal{Q}}(\tau_q, t + h|\mathcal{F}_t)$ is the forecasted quantile and y_{t+h} is the unobserved value of GDP at time t.

To get an overall picture of density forecast performance we will follow Knotek and Zaman (2019) and Carriero et al. (2025) in using diffusion indices to gauge relative forecast performance of the different estimators. For our model comparison we will use the traditional quantile estimator (QR) to be the baseline model. Let $rQS_{i,\tau_q,h}$ be the ratio of the Quantile Score for estimator i, at quantile τ_q , for forecast horizon h = 1, 4. Given the ratios of QS we calculate

the case specific index as follow:

$$f_{i,\tau_q,h} = \begin{cases} 1 & \text{if } rQS \ge 1+s \\ -1 & \text{if } rQS \le 1-s \\ 0 & \text{otherwise} \end{cases}$$
 (11)

where s is the chosen sensitivity in relative performance. To obtain the diffusion index we take the average of $f_{i,\tau_q,h}$ over τ_q , and h, i.e. $DI_i = N^{-1} \sum_{\forall \tau_q,h} f_{i,\tau_q,h}$. By construction DI_i is between 1 and -1, with lower values indicating better performance relative to the QR forecasts. Values closer to 0 indicate no difference between the estimator and traditional quantile regression. Note, that the diffusion index is sensitive to the value of s chosen. Setting s to smaller values makes the performance index more sensitive to changes relative to the QR, while setting s to larger values will only consider large improvements in forecasting performance. In Carriero et al. (2025) s = 0.05 while in Knotek and Zaman (2019) s = 0.1. Rather than pick one specific value for s we will compare various sensitivity values: s = 0.005, 0.01, 0.05. The results for the diffusion indices for the various s values for the unsorted and sorted. The results for the diffusion index are shown in table (4). Note that in this table the reference model is always the respective QR model, i.e. for unsorted quantiles the reference is QS derived from the unsorted QR fits while for the sorted quantiles the reference is the QS derived from the sorted QR fits.

The table reveals that the GNCQR and BRW has gains over the QR estimator regardless if the sorted or unsorted QS are used. The table also shows that much of the improvements are smaller given that the DI becomes less negative the larger s is. For the unsorted quantiles, the GNCQR and BRW yield improvements over the QR even when setting s = 0.05. The same cannot be said for the sorted quantiles, where s has to be set to 0.01 to see any difference in performance between the sorted QR quantiles and the sorted GNCQR quantiles. Furthermore, the sorted BRW quantiles perform on par with the sorted QR. Note that from table (3) we know that the GNCQR improves the least from sorting given it has the lowest crossing incidence.

Table 4: Diffusion Index for the estimators at different sensitivities

	Unse	orted quant	tiles	Sorted quantiles				
	BRW	GNCQR	FLQR	BRW	GNCQR	FLQR		
s = 0.005	-28.95%	-36.84%	15.79%	-15.79%	-10.53%	31.58%		
s = 0.01	-13.16%	-31.58%	5.26%	0.00%	-7.89%	21.05%		
s = 0.05	-7.89%	-5.26%	0.00%	0.00%	0.00%	2.63%		

As such, it is reassuring that even in the sorted quantiles, the GNCQR provides gains over sorted quantiles corroborating the fit results from our Monte Carlo exercise. The table also shows that the FLQR yields no improvements in quantiles score compared to QR regardless of what s is set and whether we sort the quantiles. In essence, these findings add to Carriero et al. (2025), namely that fused shrinkage is also important for growth-at-risk. Importantly, our results indicate that perhaps just as important is how this interquantile shrinkage is induced: adaptive non-crossing constraints yield improvements, while the traditional fused LASSO seem to lead to worse forecasting performance.

While the diffusion index of the QS gives a broad view of how the estimator performs, it doesn't inform us about what part of the distribution the estimator does better (or worse). To this end we also construct quantile weighted CRPS (qwCRPS) scores of Gneiting and Ranjan (2011) for all the estimators for the different forecast horizons. To calculate these measures, we take the QS and apply quantile specific weights:

$$qwCRPS_{t+h} = \int_0^1 w_{\tau_q} QS_{t+h,\tau_q} d\tau_q, \qquad (12)$$

where w_{τ_q} denotes a weighting scheme to evaluate specific parts of the forecast density. We choose this measure as the scoring rule because through different weighting schemes we can evaluate differences at different parts of the distribution. We consider 3 different weighting schemes: $w_{\tau_q}^1 = \frac{1}{Q}$ places equal weight on all quantiles (denoted as CRPS), which is equivalent to taking the average of the weighted residuals; $w_{\tau_q}^2 = \tau_q(1 - \tau_q)$ places more weight on central quantiles; $w_{\tau_q}^3 = (1 - \tau_q)^2$ places more weight on the left tail; and $w_{\tau_q}^4 = \tau_q^2$ places more weight on the right tail. The results using the different weighting schemes are presented in Table (5).

Looking at the unsorted results of Table 5, we can see that GNCQR yields the lowest weighted residual for all forecast horizons, for all weighting schemes except the right tail. These results indicate that the gains in forecast performance in GNCQR are largely from better left tail and centre forecasts. Given that the growth-at-risk framework was designed to capture downside (left tail) risk, the inability to yield better right side forecasts for the short forecast horizon is less of a concern. These results also show how BRW is a strong candidate to obtain density estimates. In particular, it is often only beaten by GNCQR. This highlights the usefulness of BRW in density estimation and for most applications should suffice as a first candidate to estimate.

The improvement in left tail forecasts is particularly important as it means that GNCQR is

Table 5: Forecast results of the different estimators

		Unson	rted		Sorted				
	CRPS	Centre	Left	Right	CRPS	Centre	Left	Right	
h=1									
QR	0.916	0.173	0.280	0.463	0.909	0.172	0.277	0.459	
BRW	0.914	0.172	0.279	0.462	0.911	0.173	0.278	0.461	
GNCQR	0.910	0.172	0.274	0.464	0.909	0.172	0.274	0.463	
FLQR	0.918	0.173	0.279	0.466	0.914	0.173	0.277	0.463	
h=4									
QR	0.559	0.108	0.168	0.284	0.554	0.107	0.167	0.280	
BRW	0.554	0.107	0.166	0.281	0.553	0.107	0.166	0.280	
GNCQR	0.552	0.107	0.165	0.280	0.552	0.107	0.165	0.280	
FLQR	0.560	0.108	0.167	0.285	0.557	0.108	0.166	0.284	

better at capturing low-probability, high-impact negative events. This is particularly useful for policymakers and financial institutions. For example, it allows for more precise calibration of macroprudential policies, such as such as the Countercyclical Capital Buffer, that aim to safeguard financial stability during periods of economic stress. With more accurate GaR measures, one can better tune the regulatory tools available to policymakers.

Overall proposed method provides gains in forecasting. This highlights how even in small (covariate) dimensional settings, one can have better forecast performance through correctly identifying which variables are quantile varying.

4.2 Coefficients

While quantile regression is robust to outliers, extreme values in the covariates can still cause issues. Since the GaR is essentially a quantile autoregressive model of Koenker and Xiao (2006), the extreme GDP observations of the COVID period enter the independent variables through the lag. This could potentially have undue influence on the coefficients and as such we will run the model on two sample periods: Pre-Covid and Full sample.

The quantile profiles of the estimated β coefficients are presented in Figures 2 and 3, for pre-COVID and full samples. β_1 is the coefficient for GDP growth and β_2 corresponds to the NFCI. A key observation is that the coefficient for the NFCI has a distinct quantile-varying profile for both samples and forecast horizons. Almost all estimators showcase a general increasing tendency across the quantiles. Nevertheless, there are some difference between the different estimators. First, FLQR is likely to shrink quantile variation at the tails, particularly the lower

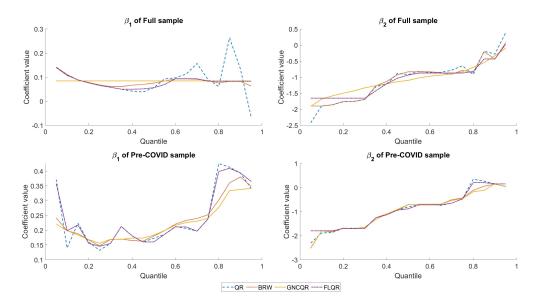


Figure 2: β coefficients for the different estimators at h=1

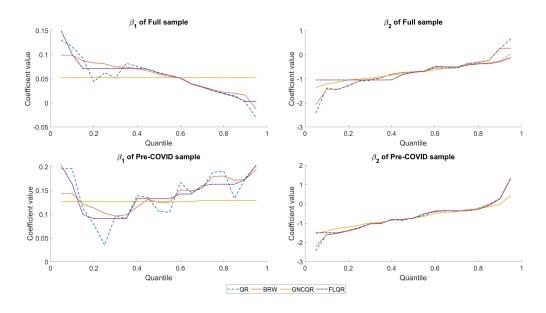


Figure 3: β coefficients for the different estimators at h=4

tail. This tendency sees FLQR have a less pronounced impact of NFCI on the lowest quantiles than the other estimators. Second, the traditional QR estimator has a positive coefficient for the upper quantiles of NFCI, while GNCQR is more likely to give a zero coefficient. Given the argument of Adrian et al. (2019), a zero coefficient for the upper quantiles of NFCI makes more sense than positive coefficients. Note that none of the estimators have shrinkage imposed on the β , i.e. there is no explicit variable selection, yet GNCQR obtains $\hat{\beta}$'s much closer to 0 at the upper quantile of NFCI. Third, GNCQR has less 'jagged' quantile profile. In fact, looking at the NFCI coefficients quantile profile, GNCQR only has negative slope for the pre-covid sample

for the h=1 forecast horizon. For all the other cases, GNCQR portrays a gentle upwards sloping profile.

While there is significant quantile variation for the coefficient of NFCI, the same cannot be said for the coefficient of GDP growth. The GNCQR yields virtually no quantile variation in almost all cases, with only pre-COVID of h = 1 showing some variation. The other estimators also show less quantile variation for GDP growth than the NFCI, but they do not fully smooth out spurious quantile variation. Interestingly, FLQR also does not shrink away the quantile variation in GDP growth.

The coefficient profile figures demonstrate that GNCQR is capable of identifying quantile variation better than simple FLQR. Furthermore, for the variables that have quantile variation, GNCQR shrinks away the "jagged" edges resulting in smoother quantile profiles. This is likely the reason for the GNCQR's superior forecast performance: the GNCQR is able to identify quantile varying variables and shrink away any unnecessary variation in these variables profiles.

These results verify the findings of Adrian et al. (2019), namely that macro-financial linkages are important for downside risk of GDP growth. Notably, the GNCQR completely shrinks the quantile variation in the GDP coefficient, which signifies that, while past values of GDP are important in determining the location of the GDP distribution, most variation in the shape of the distribution is on account of changes in the financial conditions. Furthermore, we add to the finding of Carriero et al. (2025): beyond shrinkage, interquantile shrinkage has benefits for empirical macroeconomics. This is particularly important if the goal is to obtain the coefficient profiles, since post-processing sorting procedures are applicable to the fitted quantiles and not the coefficient profiles.

4.3 Bias-variance trade-off

Figures (4) and (5) show the in-sample and out-of-sample average quantile score for different α values and compares the profiles with the fused LASSO estimator with the same hyperparameter value.

The figures show that the equivalent hyperparameters of the different estimators do not lead to the same values. This in turn leads to different profiles. In particular, GNCQR shows a more gradual change in in-sample fit for all horizons and subsamples. Furthermore, the out-of-sample fits of the GNCQR have a more pronounced minimum. Another feature of the GNCQR is that its in-sample and out-of-sample profiles are less "jagged" than that of the FLQR. This

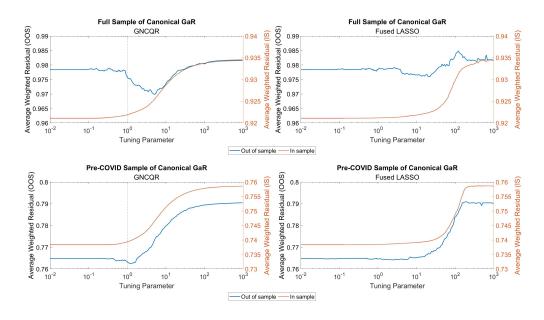


Figure 4: In-sample vs. Out-of-sample fit for various hyperparameter values (h = 1)

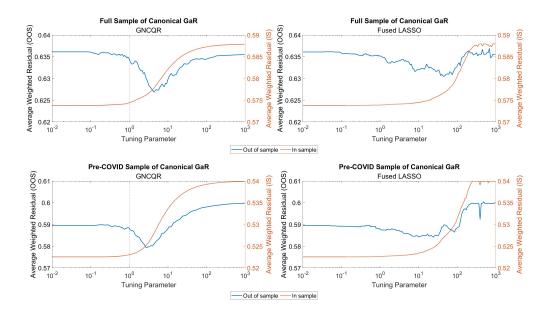


Figure 5: In-sample vs. Out-of-sample fit for various hyperparameter values (h = 4)

is true for all forecast horizons, as well as pre- and post-COVID. These are important for model selection: it is easier to optimise the tuning parameter selection when the minimum is obtain gradually. Furthermore, these properties indicate that the GNCQR shrinks quantile variation of the correct variable.

4.4 Policy implication

The previous section highlighted the importance of fused shrinkage for forecasting and getting coefficient profiles that have less spurious quantile variation. In this section we will highlight how these differences in the estimators could impact policy.

4.4.1 Expected Shortfall

The first policy tool we will investigate is expected shortfall as outlined in Adrian et al. (2019), which measures the total probability mass that the conditional distribution assigns to the left tail:

$$SF_{t+h} = \frac{1}{\pi} \int_0^{\pi} \hat{F}_{y_{t+h|\mathcal{F}_t}}^{-1}(\tau|\mathcal{F}_t) d\tau$$
 (13)

To construct expected shortfall from the estimated conditional quantiles we will utilise the methodology of Mitchell et al. (2024), which uses a nonparametric method to construct densities from estimated quantiles. We will construct the expected shortfall for the 5^{th} quantile using the nonparametric methodology. The constructed expected shortfall are in figure (6). While the same procedure can be used to estimate Expected Longrise, we opt to only focus on Expected Shortfall, as the variables used in the GaR model are more likely to capture downside risk.

The figure reveals that there is a stark difference between the estimators for ES, especially at the one year horizon (h=4). These ES estimates are critical for policy-makers as they give an overview for the evolution of downside risk.

Looking at the difference between the estimators, we can see that the QR estimator produces the most volatile ES measure compared to the other estimators, especially at the start of the estimation period. Compared to QR, the BRW estimator produces smoother ES over time, while still dipping during extreme financial stress: namely the global financial crisis and the 1970s.

FLQR deviates the most from the QR and BRW, with producing more optimistic ES during financial stress episodes, but more pessimistic ES during more tranquil periods. This tendency is particularly clear for the one year horizon. Notably, FLQR shows consistently more negative ES estimates in this horizon during relatively stable market periods (e.g., 1990-2005), suggesting a potential pessimistic bias for the FLQR.

Just like with the variable selection results, the ES made from GNCQR coefficients yields a middle-ground: between the QR's ES and the FLQR's ES. In particular, for the one-year

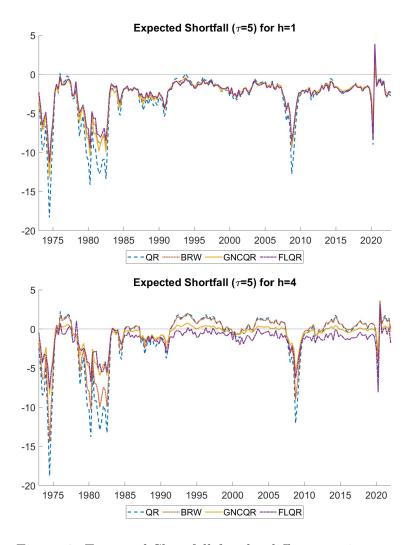


Figure 6: Expected Shortfall for the different estimators

horizon the GNCQR ES has less negative values than QR during financial stress episodes and during tranquil periods the method yields close to 0 or positive ES values, which are above the FLQR. This in turn leads to smoother ES dynamics than QR. Smooth measures of tail-risk that still capture episodes of stress are important for policy makers as abrupt and potentially unnecessary changes in regulatory tools imposes costs on the financial sector. These advantages of GNCQR are likely on account of its ability to shrink away spurious quantile variation, which helps in identifying variables that drive downside risk.

The difference between QR and the other estimators, particularly during the early sample period, highlights the value of identifying variables that drive quantile variation. GNCQR can effectively temper the extreme volatility of traditional QR while maintaining sensitivity to genuine downside risks, potentially leading to more reliable and practical risk assessments for financial stability purposes.

4.4.2 Tail local projection

Another important policy tool for GaR is local projection. Ruzicka (2021) notes that there is an identification assumption embedded in Adrian et al. (2019), namely that NFCI has no contemporaneous effects on GDP growth distribution. This assumption has been used by Ruzicka (2021), Wojciechowski (2024), and Chavleishvili and Manganelli (2024) to create quantile IRFs and quantile local projections. Importantly, these methods rely on the estimated coefficients to construct the quantile projections through time, which cannot be sorted. We will produce quantile local projections for the 4 estimators for the 5^{th} , 10^{th} , and 25^{th} quantile. We will follow Ruzicka (2021) and deviate from Adrian et al. (2019) on two fronts: a) we will estimate future growth directly, rather than averaged over the forecast horizon; and b) we will include additional lags of GDP and NFCI. The quantile LPs of the response of GDP to a unit shock in NFCI are shown in figure (7).

Across all three quantiles, there's a general pattern of initial decline followed by recovery, but the timing and magnitude of these movements vary considerably by estimator. The QR consistently shows the deepest and most volatile projections, while FLQR shows the quickest recovery. BRW and GNCQR typically fall between these extremes, with GNCQR often showing slightly less negative projections than BRW, particularly in the medium run (h = 4 - 8).

In the extreme lower tail (5th quantile), we observe substantial heterogeneity across estimators. The QR estimator exhibits the highest volatility, with deeper initial drops reaching below -4 around h = 1 - 2 and again near h = 6 - 7. Indeed this volatility was the primary motivation for Ruzicka (2021) to propose smoothing the projections dynamically over the horizons. In contrast, the FLQR estimator shows a remarkably different pattern, returning to near-zero values much more quickly after horizon 4, indicating it might be producing more conservative estimates of tail risk. When looking at the coefficient profiles we saw that the FLQR was prone to shrinking quantile variation at the tails. As such it is likely that the quick recovery suggested by the FLQR is a consequence of this overshrinking behaviour. The GNCQR and BRW produce smoother projections than the QR but do not return to 0 as quickly as the FLQR. This finding is particularly interesting and suggests that potentially the smoothed quantile LP framework of Ruzicka (2021) induced fused shrinkage at each horizon. We leave for future research to explore the link between smoothing quantile variation and smoothing projections over time.

The differences between estimator choices diminish as we move up the quantiles, with the 25th quantile showing limited difference between the estimated profiles especially in the short

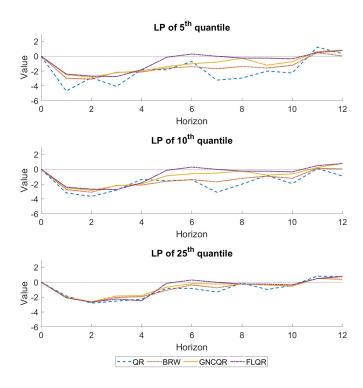


Figure 7: Local projections for the different estimators at different quantiles

run. After h = 4, the estimators begin to diverge again, though less dramatically than in the lower quantiles.

These differences highlight the importance of estimator choice in growth-at-risk analyses, especially when focusing on extreme quantiles that are of particular interest to policymakers concerned with tail risks to economic growth. The varying persistence and magnitude of negative projections across estimators could lead to significantly different assessments of downside risks and potentially different policy recommendations.

5 Conclusion

This paper develops an adaptive non-crossing constraint, encompassing the traditional quantile regression estimator, the Bondell et al. (2010) non-crossing estimator, as well as the composite quantile regression, as special cases by varying tightness of the constraint. By developing non-crossing constraints that can be tightened, we study the properties of these constraints on the estimated β parameters. Doing so reveals that non-crossing constraints are simply a type of Fused LASSO with quantile specific shrinkage parameters.

We verify that imposing non-crossing constraints is equivalent to introducing fused shrinkage

with a Monte Carlo experiment. We also show how our proposed Generalised Non-Crossing Quantile Regression (GNCQR) estimator provides fits that are either better or nearly as good as those of the BRW estimator (Bondell et al., 2010). When looking at the variable selection properties of the different estimators we found that traditional Fused LASSO (FLQR) estimator overshrinks while BRW estimator undershrinks compared to GNCQR.

Taking an interquantile shrinkage lens to US Growth-at-Risk, we found that our proposed GNCQR estimator outperforms both traditional quantile regression and standard fused LASSO approaches, particularly in forecasting left-tail risks. The GNCQR effectively identifies variables that drive quantile variation while shrinking away spurious variation, resulting in smoother coefficient profiles and improved forecast performance. Notably, our results confirm the findings of Adrian et al. (2019) that financial conditions are the drivers of nonlinearities in GDP distribution, while past GDP growth acts primarily as a location shifter.

The policy implications of better GaR coefficients is also explored. Expected Shortfall estimates derived from GNCQR exhibit less volatility while remaining sensitive to genuine financial stress episodes, offering policymakers a more reliable assessment of downside risks. Similarly, quantile local projections based on GNCQR provide balanced medium-term forecasts that avoid both the excessive volatility of traditional quantile regression and the potentially over-optimistic recovery paths suggested by standard fused LASSO. These improvements in policy-relevant metrics underscore the importance of appropriate interquantile shrinkage in macroeconomic risk assessment.

In sum, the non-crossing constraints of Equation (5) bestow upon the quantile estimator additional attractive properties: (1) it can distinguish variables with quantile variation from those without, and shrinks the correct variable; and (2) it renders the estimated quantile profiles less 'jagged', i.e., it removes sudden reversions in the difference in the β coefficient.

The findings of this paper reach beyond the estimator proposed here. In particular, because of the equivalence between non-crossing and fused-shrinkage, one can extend the methodology to obtain non-crossing Bayesian quantile regression estimators like in Lancaster and Jae Jun (2010). To our knowledge, currently the most popular way to obtain non-crossing Bayesian quantiles involve post-processing methods such as Rodrigues and Fan (2017). Implementing theorem (1) to the Bayesian realm has clear advantages.

The formulation of the GNCQR is very simple: the hyperparameter is simply a scalar. It is probable that one can gain further improvements in estimation by making the hyperparameter have more than just one dimension: making the hyperparameter variable specific has been

shown to offer better performance in other contexts (Zou, 2006). Naturally, doing so will require a better hyperparameter tuning procedure, since grid-search is a computation-intensive choice even with just a single parameter. We leave this extension to future research.

Another avenue of research is shrinking both the level and difference coefficients, similar to Jiang et al. (2014). However, following Szendrei and Varga (2023), shrinkage in levels would require a separate dedicated hyperparameter. This way the two types of shrinkage could be incorporated yielding the possibility to identify quantile varying sparsity like in Kohns and Szendrei (2021) and Szendrei and Varga (2023), while being principled about the degree of fused shrinkage one needs to impose. We leave this work for future research.

Finally, since the bias introduced by non-crossing constraints is similar to biased bootstrap methods proposed for regression under monotone or convexity constraints, one can use degree of bias to test model adequacy. Developing tests of model adequacy using bias induced by the quantile non-crossing constraints is a promising avenue for future research.

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A Estimators

• Quantile regression (Koenker and Bassett, 1978)

$$\hat{\beta}_{QR} = \underset{\beta}{argmin} \sum_{q=1}^{Q} \sum_{t=1}^{T} \rho_q (y_t - x_t^T \beta_{\tau_q})$$

• Composite quantile regression (Koenker, 1984; Zou and Yuan, 2008)

$$\hat{\beta}_{CQR} = \underset{\beta}{argmin} \sum_{q=1}^{Q} \sum_{t=1}^{T} \rho_q(y_t - x_t^T \beta)$$

• Non-crossing quantile regression (Bondell et al., 2010). Here, $z \in [0, 1]$, i.e. it is the x rescaled.

$$\hat{\beta}_{BRW} = \underset{\beta}{argmin} \sum_{q=1}^{Q} \sum_{t=1}^{T} \rho_q (y_t - z_t^T \beta_{\tau_q})$$

$$s.t. \ \gamma_{0,\tau_p} \ge \sum_{k=1}^{K} \gamma_{k,\tau_q}^-$$

• Fused shrinkage quantile regression (Jiang et al., 2013)

$$\hat{\beta}_{JWB} = \underset{\beta}{argmin} \sum_{q=1}^{Q} \sum_{t=1}^{T} \rho_{q} (y_{t} - x_{t}^{T} \beta_{\tau_{q}})$$

$$s.t. \ k^{*} \geq \sum_{q=2}^{Q} \sum_{k=1}^{K} (\gamma_{k,\tau_{q}}^{+} + \gamma_{k,\tau_{q}}^{-})$$

• Generalised Non-Crossing Quantile Regression

$$\hat{\beta}_{GNCQR} = argmin \sum_{q=1}^{Q} \sum_{t=1}^{T} \rho_{q} (y_{t} - x_{t}^{T} \beta_{\tau_{q}})$$

$$s.t. \ \gamma_{0,\tau_{p}} + \sum_{k=1}^{K} \left[\bar{X}_{k} - \alpha (\bar{X}_{k} - min(X_{k})) \right] \gamma_{k,\tau_{q}}^{+} \ge \sum_{k=1}^{K} \left[\bar{X}_{k} + \alpha (max(X_{k}) - \bar{X}_{k}) \right] \gamma_{k,\tau_{q}}^{-}$$

B Bias-variance trade-off discussion

GNCQR allows us to gradually enforce non-crossing constraints by inducing fused shrinkage. This naturally entails that there is some type of bias-variance trade-off. In essence, as we increase α we increase the amount of non-crossing we want to impose. By introducing non-crossing constraints we impose a bias on the quantile property (i.e. we no longer obtain the minimum of the tick-loss weighted ℓ_1 residuals) and instead enforce monotonically increasing quantiles. This implicitly enforces that the data below τ_q must be a subset of the data below τ_{q+1} . As we move from $\alpha = 0$ to $\alpha = 1$, we gradually move from the quantile property to the 'quantile subset' property. Once $\alpha > 1$, GNCQR starts to become more restrictive in how the quantile subset property is satisfied. In particular, it will start to penalise quantile closeness as well as quantile crossing. Formally this can be shown as follows:

$$\{I(\varepsilon_{\tau_{q-1}} - \xi \le 0)\} \subseteq \{I(\varepsilon_{\tau_q} \le 0)\} \tag{14}$$

Here $\{I(\varepsilon_{\tau_q} \leq 0)\}$ is the elements of the quantile residual that are negative for the fitted τ_q^{th} quantile, i.e. the observations below the fitted quantile. The quantile subset property necessitates that the observations with negative residuals of the τ_{q-1} quantile must be a subset of the observations with negative residuals of the τ_q quantile. This is achieved in equation (14) when $\xi = 0$.

As α of equation (6) goes above 1, the ξ term in the quantile subset property (equation (14)) increases as well. Increasing the ξ parameter means that even some observations that are above the τ_{q-1} fitted quantile form the set that has to be below τ_q quantile. In this way, quantile that get too close to each other are treated as if they cross. As such, when $\alpha > 1$, the ξ term in equation (14) becomes positive, which in turn leads to penalising quantile closeness in addition to quantile crossing. Conversely, when $\alpha < 1$, the ξ term is negative and the quantile subset property allows for some degree of quantile crossing. When $\alpha = 0$, the ξ term is some large number, making equation (14) trivial to satisfy.

¹⁶The bias thus introduced by non-crossing constraints is similar to biased bootstrap methods (data tilting, data sharpening, etc) proposed for regression under monotone or convexity constraints. The degree of bias can be used to test model adequacy, which is a topic retained for future research.

C Hyperparameter selection methods

The choice of cross-validation technique can lead to differences in results. In particular, there is a documented trade-off between model identification and minimising predictive risk (Yang, 2005). Broadly speaking, leave-one-out cross-validation is asymptotically equivalent to AIC (Stone, 1977), while the various block cross-validation methods are closer to the BIC (Shao, 1997), if the size of the training sample (relative to the validation sample) goes to zero as $n \to \infty$. This underlies the key point of Yang (2005) that one cannot simultaneously achieve model consistency (in terms of model selection) and efficiency (in terms of achieving lowest error variance). Note also that by choosing the hv-block CV of Racine (2000), we are implicitly expressing a preference in favour of model selection. If the interest is purely data fit, it might be beneficial to opt for a leave-one-out CV. We retain consideration of the GNCQR using other CV methods to future research.

Given the results of Stone (1977) and Shao (1997), one can also rely on information criteria for hyperparameter selection. The equation for AIC and BIC for quantile regression can be found in Jiang et al. (2014). Utilising the information criteria has the advantage of only needing one estimation (per hyperparameter), rather than one per block. This can speed up computation especially for larger sample sizes.

Cross-validation has been shown to have some drawbacks for some applications (Bates et al., 2024). Specifically, it has been shown that cross-validation estimates the average prediction error of models fit on other unseen training sets drawn from the same population, rather than the prediction error of the model fit on the specific training set. On account of this, confidence intervals for prediction error may have lower coverage. While these results are undoubtedly important, they are less of a concern for model selection. In essence, all candidate hyperparameters are biased in the same way which has little influence on their relative rank. This entails that the optimal hyperparameter choice remains valid.^{17, 18}

¹⁷For a more rigorous mathematical treatment of the consistency of cross-validation for model selection, see Theorem 2 of Yang (2007) and Proposition 2 of Wager (2020).

¹⁸While our analysis assumes that the relative ranking of hyperparameters remains valid, further investigation might reveal contexts in which Bates' nested cross-validation method improves hyperparameter tuning.