Forecasting in Data-Rich Environments: An Application to Inflation Rates^{*}

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Abstract

Stock and Watson (1998 and 1999) developed a factor-model approach which allows for large data sets to be systematically summarized by to a few explanatory factors. In this paper two other methods are proposed. The first one, Partial Least Squares is imported from the Chemometrics literature. The second one, which is based on the Combination of Forecasts literature is a modification of Stock and Watson's method. We call it Principal Components Combination. These methods are compared in an empirical application to inflation. It is found that overall the Principal Components Combination performs the best.

Keywords: combination of forecasts; factor analysis; forecasting; inflation; partial least squares; principal components

1 Introduction

With enormous amounts of new information, on several economic indicators, arriving in real time, applied Macroeconomists have the problem of dealing with huge data sets, with hundreds of explanatory variables that can be useful for forecasting purposes. Usually, we have, at most, a few hundred observations, making the use of so many variables impossible in a single regression model. Even with financial data, where much longer time series may

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easily be found, it is of dubious interest to consider hundreds of regressors. Nevertheless, it is inefficient not to use all available information. More information should be helpful, not a problem.

One popular method to deal with this problem of many explanatory variables is the Principal Components Regression (PCR), which was applied by Sargent and Sims (1977) and Geweke (1977). More recently, this method has been successfully applied to US Macroeconomic data (Stock and Watson (1998, 1999, and 2002)), Bernanke and Boivin (2003). Marcellino, Stock and Watson (2003) also applied this method to European data, but there the Principal Components Regression could not consistently improve upon a simple Auto Regression model.

This literature is growing, and some nice asymptotic results have already been derived — see Stock and Watson (1998), Forni, Hallin, Lippi, and Reichlin (2000) and Bai and Ng (2002). Still, some criticisms to this approach remain:

- the results are very sensitive to the scale measurement of the variables, and
- the principal components are constructed without taking into consideration any relationship between the regressors and the dependent variable.

One method, which tries to overcome the second problem is the Partial Least Squares (PLS). This method, popular in the Chemometrics literature, was proposed by Wold (1975). PLS became popular during the 80's and, a decade later, several papers appeared in the Statistics literature, analyzing the properties of this method. Although popular among chemometricians, this method has never become popular among econometricians and economists. An exception is Gibson and Prisker (2000) who applied this method to economic data.

A different branch of literature is the Combination of Forecasts proposed by Bates and Granger (1969) — see also Granger (1989) and Deutsch, Granger and Terävirsta (1994). This literature deals with the problem of having multiple forecasts for the same variable. These authors, and others, argue that combining the different forecasts in a suitable manner leads to better predictions than the individual ones. Bates and Granger (1969) argued that a simple way to combine the different forecasts is to run a simple regression (OLS) to find the best combination. Note that if one has a large number of forecasts then simple OLS will not be appropriate. Chan, Stock and Watson (1999) make the argument that a suitable way to combine a large number of different forecasts is by PCR.

As an alternative to PCR and PLS, we will combine PCR with the forecast combination approach. More specifically, we will use each explanatory variable to obtain a forecast for the dependent variable, and then combine a large number of forecasts using the PCR method. The proposed method has two advantages: it is scale invariant, thereby dealing with the first criticism, and it takes into consideration the explanatory power of the independent variables on the dependent variable. There is also a third advantadge, which is left for future research, we don't need to restrict ourselves to linear models when we produce the forecasts, so this method has more modelling flexibility than PCR or PLS.

The rest of the paper is organized as follows: section 2 sets up the basic model, and describes and relates two well-known estimation methods: PCR and PLS. In section 3, a new method is proposed: PCC. In section 4 the different methods are applied to inflation forecasting and compared. Section 5 concludes.

2 The Model

Let the basic data be given by $X = (x_1, ..., x_N)$, a matrix of T observations of N independent variables, and y, a vector with T observations of the dependent variable. To facilitate interpretation, we assume that all the variables have already been demeaned.

Consider a factor model of the form:

$$\begin{cases} x_n = \lambda_{n,1} F_1 + \dots + \lambda_{n,K} F_K + e_n & n = 1, \dots, N \\ (T \times 1) & (1 \times 1)(T \times 1) & (1 \times 1)(T \times 1) & T \times 1 \\ y = \beta_1 F_1 + \dots + \beta_K F_K + \varepsilon \\ (T \times 1) & (1 \times 1)(T \times 1) & (1 \times 1)(T \times 1) & T \times 1 \end{cases}$$

or, stacking the vectors together:

$$\begin{cases} X = F \lambda + e \\ {}_{(T \times N)} = F \lambda + e \\ y = F \beta + \varepsilon \\ {}_{(T \times 1)} = F \lambda + \varepsilon \\ {}_{(T \times K)(K \times 1)} + F \lambda + \varepsilon \end{cases}$$
(1)

The crucial assumption of this model is that y depends on X by only a few unobserved factors F. A factor model of this type is useful when the number of predictor variables is large (possibly larger than T), making more common forecasting techniques unattractive or not feasible. Since F may contain lagged values of the underlying factors, this model is also called a dynamic factor model.

A natural way to estimate the parameters of the second equation of system 1 is to replace the unobservable factors by estimated factors, and then estimate β by Ordinary Least Squares (OLS).

In the next subsections of the paper we consider two different methods to estimate the unobserved factors:

• Principal Components Regression (PCR), and

• Partial Least Squares (PLS).

The first is becoming increasingly popular among econometricians, while the latter is most popular in the Chemometrics literature. We will then propose a modification of the PCR based on the forecast combination literature. This modification follows the spirit of PLS (by taking into consideration the effect of each predictor for the dependent variable) but essentially uses the analytical tools of PCR, with the advantage of being scale invariant.

2.1 Principal Components Regression

If the model described above is correct, a possible procedure is to use the principal components of X as an estimate of the factors, and use them to estimate the second equation of system 1.

As Stone and Brooks (1990) showed, the idea of this method is to find the linear combinations of the X variables, such that a vector of weights, p_1 , maximizes p'X'Xp, and then p_2 is chosen to maximize p'X'Xp subject to the constraint that $p'p_1 = 0$. The vectors of weights are normalized to have unit distance. Thus p_1 is the normalized eigenvector of X'Xassociated with the largest eigenvalue, p_2 is the normalized eigenvector associated with the second largest eigenvalue, and so on.

By choosing the components associated with the largest eigenvalues, one obtains the linear combinations of X that are orthogonal to each other and simultaneously have the largest variance. Intuitively, by choosing linear combinations with the possible largest variance, one is, in a certain sense, maximizing the information contained in those linear combinations. When carrying out the empirical applications, we discuss how to estimate the numer of factors.

Stock and Watson (1998), Forni *et al.* (2000) and Bai and Ng (2002) provide consistency results for this method. The asymptotic theory of this method assumes not only $T \to \infty$ but also $N \to \infty$. For example, Bai and Ng assume that $E ||F_t||^4 < \infty$ and $\frac{1}{T} \sum_{t=1}^T F_t' F_t \to \Sigma_F$ as $T \to \infty$, with Σ_F being some positive definite matrix. They also assume that each factor has a nontrivial contribution to the variance of X: $\left\|\frac{\lambda'\lambda}{N} - D\right\| \to 0$ as $N \to \infty$, with D being some positive definite matrix, and $\|\lambda_n\| \leq \bar{\lambda} < \infty$. They further impose some conditions on the error terms of the X variables, allowing for heteroskedasticity in both time and cross section dimensions and some dependence between factors and the errors. Bai and Ng (2002), and Stock and Watson (1998) with a different set of assumptions, show that, asymptotically, the estimated factors and the true factors span the same space.

2.2 Partial Least Squares

Using PCR only the information contained in the X-data is used to estimate the factors. Not all the information is used, as the relationship to the dependent variable is not considered.

PLS first appeared in the form of an algorithm (which is described bellow). Stone and Brooks (1990) showed that with PLS a vector of weights p_1 is chosen to maximize p'X'yy'Xp, then p_2 is chosen to maximize p'X'yy'Xp subject to the constraint that $p'(X'X)p_1 = 0$. So one is finding the linear combination of the X variables which maximizes the squared sample covariance between X and y. Although PLS deals with the second criticism to PCR, it fails to address the first, as it is scale dependent as well. The usual procedure is to normalize all the variables to have unit variance. By doing this, maximizing the squared sample covariance amounts to maximizing the squared sample correlation.

There are at least two algorithms (one proposed by Wold (1975) and the other proposed by Martens (1985)). Helland (1988) proved the equivalence between them and also proposed a third method, which is computationally more convenient. Next we will describe the algorithm that Wold (1975) proposed and, after that, the alternative basis that Helland (1988) proposed. For a description of both algorithms and the proof of their equivalence and also the equivalence of the alternative basis, the reader is referred to Helland (1988). For some consistency results of PLS the reader can consult Naik and Tsai (2000)¹.

2.2.1 The original PLS algorithm

Define $E_0 = X$ and $f_0 = y$. Define E_a and f_a recursively as:

$$E_a = E_{a-1} - \hat{F}_a \hat{\lambda}'_a$$

$$f_a = f_{a-1} - \hat{F}_a \hat{\beta}_a$$
(2)

where \hat{F} stands for the factor estimate.

We will need to determine \hat{F}_a , $\hat{\lambda}_a$ and $\hat{\beta}_a$ in these equations. As with the Principal Components approach, each estimated factor \hat{F}_a will be a linear combination of the X variables. For example, for a = 1 we have:

$$\hat{F}_{1}_{T\times 1} = \sum_{n=1}^{N} x_n p_{n1}_{N\times 11\times 1} = \sum_{T\times N}^{N} p_{1}_{N\times 11}$$
(3)

Since we would like to use the information contained in y to estimate the factors the

¹Assuming that the explanatory variables are i.i.d., these authors prove consistency of the PLS for $T \to \infty$. Extension to stationary variables is immediate.

weights will be chosen as:

$$p_1 = X'y \tag{4}$$

With this method, explanatory variables with a higher covariance with Y will receive a higher weight.

In general we have:

$$\hat{F}_a = E_{a-1}p_a \tag{5}$$

$$p_a = E'_{a-1} f_{a-1} (6)$$

We still need to determine $\hat{\lambda}_a$ and $\hat{\beta}_a$. To have the best fit in equation (2) we use the regression coefficients. For a = 1 we have $y = \hat{F}_1 \hat{\beta}_1 + f_1$ and $X = \hat{F}_1 \hat{\lambda}'_1 + E_1$, so the regression coefficients are given by $\hat{\beta}_1 = (\hat{F}'_1 \hat{F}_1)^{-1} y'_1 \hat{F}_1$, and $\hat{\lambda}'_1 = (\hat{F}'_1 \hat{F}_1)^{-1} \hat{F}'_1 X$. In general we have:

$$\hat{\lambda}_a = \left(\hat{F}'_a \hat{F}_a\right)^{-1} E'_{a-1} \hat{F}_a \tag{7}$$

$$\hat{\beta}_a = \left(\hat{F}'_a \hat{F}_a\right)^{-1} f'_{a-1} \hat{F}_a \tag{8}$$

Note that since the \hat{F}_a 's are orthogonal to each other (again see Helland (1988)), instead of equations (7) and (8) we can use:

$$\hat{\lambda}_{a} = \left(\hat{F}'_{a}\hat{F}_{a}\right)^{-1}X'\hat{F}_{a}$$
$$\hat{\beta}_{a} = \left(\hat{F}'_{a}\hat{F}_{a}\right)^{-1}y'\hat{F}_{a}$$

With this method, the first factor to be estimated is $\hat{F}_1 = (X)(X'y)$. So instead of finding $T \times N \xrightarrow{N \times 1} N \times 1$ the linear combination of the X variables that maximizes the variance, one is using the covariance between each predictor and the dependent variable as the weight of that variable. Then the second factor will be estimated using the covariance between $(X - \hat{F}_1 \hat{\lambda}'_1)$ and $(y - \hat{F}_1 q'_1)$, and so on.

2.2.2 An alternative basis

The next proposition allows us to use a computationally more convenient method.

Proposition 1 Let S_A be the space spanned by $p_1, \ldots p_A$. As long as p_A is nonzero, an alternative basis for S_A is given by the vectors $(X'y), (X'X)(X'y), \ldots, (X'X)^{A-1}(X'y)$.

Proof. See Helland (1988) or Stone and Brooks (1990). ■

This algorithm is computationally easier to implement than the original one proposed by Wold (1975), without requiring any iterative procedure.

2.3 Prediction, spectral representation and relation between PLS and PCR

For a moment, let us consider a population version of the model described in system 1, where there is no noise.

Consider the spectral decomposition of $S = X'X = \sum_{k=1}^{K} \varphi_k p_k p'_k$, where p_k is the eigenvector associated with the strictly positive eigenvalue φ_k (assuming that X'X has rank K).

Using the principal components regression, the predicted value for y is given by:

$$\hat{y} = F(F'F)^{-1}F'y = \sum_{k=1}^{K} Xp_k (p'_k X' X p_k)^{-1} p'_k (X'y)$$

For prediction purposes all the non-relevant eigenvectors of X'X can be deleted. Also, if an eigenvalue has multiple eigenvectors associated with it, the corresponding terms can be substituted by only one term by rotating in eigenspaces with equal eigenvalue, such that we get only one eigenvector. For example, suppose that $\lambda_1 = \lambda_2$, then we can replace p_1 and p_2 by $p_1^* = \left(\frac{p_1 p'_1 + p_2 p'_2}{\left(\left(p'_1 s\right)^2 + \left(p'_2 s\right)^2\right)^{\frac{1}{2}}}\right)(X'y)$. Note that $p_1^* p_1 = 1$, and $p_1 p'_2(X'y) + p_2 p'_2(X'y) = p_1^* p_1^{*'}(X'y)$.

Definition 2 The relevant eigenvectors of X'X to predict y are the ones associated with different eigenvalues which satisfy $p'_k(X'y) \neq 0$. The corresponding factors $F_k = Xp_k$ are the relevant factors in X for prediction of y. Let A be the total number of relevant eigenvectors.

Proposition 3 The population PLS space has dimension A, and when this minimal number of terms is used, the population PLS regression vector and the population PCR regression vector are equivalent.

Proof. See Helland 1990. ■

This proposition tells us that the PLS and PCR regression vectors are equivalent when all the relevant components are included. Some stopping rule must be defined when applying the algorithm and hence the previous results will only be approximate: with real and noisy data it is highly unlikely that we find exact repeated values for the eigenvalues or that $p'_k(X'y) = 0$ (the sample relevant components will be very close to min (N, T - 1)). Perhaps the largest advantage of PLS over PCR is that the possible nonsense of giving a large weight to an irrelevant explanatory variable is avoided. For example, suppose that the variable Xg is completely uncorrelated with y. Using the PCR algorithm there is nothing to prevent this variable from receiving a possibly large weight, while with the PLS approach, this variable receives approximately zero weight.

3 Forecast Combination and Principal Components

Bates and Granger (1969) — see also Granger (1989) and Deutsch, Granger, and Terävirsta. (1994) — suggest that when there are several forecasts for the same variable, one sensible approach is to combine these several forecasts. Several combination methods have already been by proposed. Chan, Stock and Watson (1999) argue that a suitable way to combine the different forecasts is to model them as an approximate factor model.

If one has N explanatory variables, then, using univariate regressions it is possible to produce N forecasts that can be combined using the PCR approach. We will call this procedure Principal Components Combination (PCC).

Let us consider in detail how to implement the PCC method. Step 1, project y onto the space spanned by each of the N explanatory variables: $z_n = x_n (x'_n x_n)^{-1} x'_n y$, for n =1,2,..., N. Step 2, create a new matrix of explanatory variables: $Z = (z_1, ..., z_N)$. Step 3, find the eigenvectors u_i of Z'Z associated with positive eigenvalues. Let u_1 be the eigenvector associated with the largest eigenvalue, u_2 with the second largest, and so on. Step 4, use as new regressors the variables Zu_A associated with the A highest eigenvalues.

By choosing the principal components one is choosing a linear combination of the explanatory variables (Z) that maximizes the variance. In this case the variance of each individual predictor has a natural interpretation: it is the explained variance of y by the corresponding original explanatory variable. One is no longer finding the principal components without taking into account the information contained in y. The weight that each variable receives is not independent from the relationship between the regressors and the dependent variable. Variables with higher explanatory power are also the variables with the highest variance, and hence they will tend to receive a higher weight. On the other extreme, if some variable x_n has no explanatory power over y, then the estimated y's will be constant (since all variables are in deviations from the mean, z_n will be a column of zeros), and this variable will receive zero weight when constructing the principal components.

If we choose A components the estimated value for y is

$$\hat{y} = Z(u_1, ..., u_A) \left[(Z(u_1, ..., u_A))' Z(u_1, ..., u_A) \right]^{-1} (Z(u_1, ..., u_A))' y$$

The final forecasts will be independent of the scale of the original variables X, because the matrix Z will not be changed with the scale of the original variables, so practitionersare free of choosing appropriate scaling of data.

Proposition 4 Let K be the number of eigenvectors (p_k) of X'X associated with nonzero eigenvalues and assume that $cov(y, x_n) \neq 0$, n = 1, ..., N. Then $(Zu_1, ..., Zu_K)$ and $(Xp_1, ..., Xp_K)$ span the same space.

Proof. Note that $a_n = (x'_n x_n)^{-1} X'_n y$ is a scalar different from zero asymptotically as long as $cov(y, x_n) \neq 0$. So $z_n = a_n x_n$ and hence X and Z span the same space and the number of eigenvectors associated with nonzero eigenvalues of X'X and Z'Z are the same (i.e., K). Since $(Xp_1, ..., Xp_K)$ span the same space as X, and $(Zu_1, ..., Zu_K)$ span the same space as Z, we must have that $(Xp_1, ..., Xp_K)$ and $(Zu_1, ..., Zu_K)$ span the same space.

This proposition tells us that, when considering the population version of the model, PLS and PCC are equivalent, as long as all the components associated with strictly positive eigenvalues are used. In a sample regression this result will have some noise because the number of positive eigenvalues will be min (N, T - 1), and obviously it is unfeasible to use so many components. In small samples, one would expect that when only a few components are considered then the components estimated by PCC will produce better forecasts (we will be able to confirm this later) but asymptotically, with N and T approaching infinity, the results should converge.

We are restricting ourselves to produce the forecasts using a linear model. Although we do not pursue this route here, there is nothing fundamental about that restriction. If we believe that a nonlinear model is better to capture the relationship between, say, x_i and y then we can use for that variable a nonlinear model. This is a possible extension of the method we are proposing.

4 Empirical Application

Inflation forecasts are at the center of policy deliberations at inflation-targeting central banks. They also play an important role in non-inflation-targeting central banks such as the Federal Reserve and the European Central Bank. In countries with centralized wages barganing, inflation forecasts also play a crucial role because unions and firms are mainly concerned in negociating real wages. Rational economic agents base their investment decisions on expected real interest rates.

In this section we will apply the methods described in sections 2 and 3 to forecast inflation. The data was taken from the DRI-Mcgraw Hill Basic Economics database spanning a time horizon from October 1968 to March 2003. This amounts to 413 monthly observations of 140 variables.

All these variables are economic indicators measuring different aspects of the economy activity, such as real output and income, employment, sales, consumption, housing starts inventories, stock prices, exchange rates, interest rates, monetary aggregates, wages and, obviously, inflation.

Most variables were logarithmized (namely all the strictly positive variables that were not in the form of rates or ratios). Using the ADF and Phillips Perron tests, we test each series to check if it was stationary or not. In the cases in which the series were not stationary we took first differences.

We will produce h-month-ahead inflation forecasts using different specifications. We will estimate the model using T observations and use the estimated model to produce an out of sample inflation forecast and compare this forecast with the realized inflation rate. This will be done recursively for the complete sample. Then the Mean Square Prediction Error (MSE) and the Mean Absolute Prediction Error (MAE) of the out of sample forecasts are obtained to compare the accuracy of the different methods proposed. Rolling window estimation is used. For example, if we consider a sample size of 100 observations, we use the first 100 observations to predict the inflation of period 101. Then we will reestimate the model using observations 2-101 to produce a forecast of the inflation in period 102, and so on

As in Stock and Watson (1999) we will consider two different measures of inflation. One is the Consumer Price Index (with the mnemonic PUNEW) — a Laspeyres index — and the other is the Personal Consumer Expenditure deflator (with the mnemonic GMDC) — a chain weighting.

In the more general form, the model to be estimated is:

$$\pi_{t+h}^{h} = \alpha + \beta (L) \mathbf{x}_{t} + \gamma (L) \pi_{t} + e_{t+h}^{h}, \ t = 1, 2, \dots$$
(9)

The dependent variable is π_{t+h}^h is given by $\pi_{t+h}^h = \left(\left[\prod_{i=1}^h (1+\pi_{t+i}) \right]^{\frac{1}{h}} - 1 \right)$. This specification can be thought of as predicting inflation over the next h months.

The regressor(s) \mathbf{x}_t is (are) some explanatory variable(s) available at time t. $\boldsymbol{\beta}(L)$ is a polynomial vector in the lag operator L, and $\gamma(L)$ is a polynomial in the lag operator L.

We will consider several competing methods for the choice of \mathbf{x}_t :

- the Phillips curve: \mathbf{x}_t is just the unemployment rate between all workers of 16 years or older of period t,
- the pure time series AR model: \mathbf{x}_t is omitted,

• PCR, PLS and PCC: \mathbf{x}_t is recursively chosen in each regression according to the methods described below.

For PCR we compute the principal components, using the procedure described in section 3, and choose the one associated with the largest eigenvalue. Then to determine whether we should include the component associated with the second highest eigenvalue we use a modified version of the Bayes Information Criterion (BIC), proposed by Bai and Ng (2002)². If the inclusion of the second component is rejected, the process stops; otherwise the same criterion is used again to evaluate the score associated with the third largest eigenvalue, and so on. A maximum of 10 components is allowed. With the PCC the procedure is the same as the PCR method. The only difference is that instead of considering the original variables, these are pre-transformed (as described in section 3).

For example, if the original variable is a vector X_i , we will work with $z_i = X_i (X'_i X_i)^{-1'} y$, where y is the dependent variable, the *h*-period ahead inflation rate. Finally to estimate the components using the PLS method, we use the alternative basis described in proposition 2. The first component to be included is X(X'y). Then one checks if X[(X'X)X'y] should be included. If the inclusion is rejected, the process stops; otherwise we check if $X[(X'X)^2 X'y]$ should also be included, and so on. Again a maximum of 10 components is allowed.

Two aditional matters should be mentioned. First since PLS and PCR are scale sensitive we followed the suggestion in the literature and, in each regression, we normalized all the variables to have unit variance. Although not reported, we also considered the case with no normalization. The performance of these two methods is severely worse without the normalization. We should also note that since we have 140 explanatory variables and when constructing the X matrix, we include two more lags of each explanatory variable, the matrix of explanatory variables has 420 columns.

To choose the order of the polynomials $\beta(L)$ and $\gamma(L)$, we use the typical BIC.

4.1 Results

In tables 1 to 5 we can check the performance of the various methods. On the top part of each table we have the relative (to PCR) mean square forecast errors and on the lower part the relative mean absolute forecast error. We considered several sample sizes, so that one can evaluate the performance on small and on larger samples. Naturally, the larger the sample size is, the lesser the number of feasible estimations is.

²Bai and Ng showed that the standard BIC can only consistently estimate the correct number of factors if the factors are known. If one has to estimate the factors then the BIC may not consistently estimate the correct number of factors. The same criterion was used by Marcellino et al. (2003).

By a simple counting procedure it is apparent that the PCC method gives the most accurate forecasts: in 76 times, out of 120, the PCC had the smallest out of sample relative forecast errors. PLS also performed reasonably well, being able to produce the smallest mean forecast errors 32 times, followed by PCR (8 times) and the AR model (4 times).

Taking the PCR model as the benchmark, we conclude that PCC was able to beat PCR 101 times (out of 120), while PLS produced more accurate forecasts than PCR (according to the two different criteria) 70 times. Comparing the PCC method with PLS we can see the PCC produces more accurate forecasts 84 times (out of 120).

To compare the performance of these methods in a more formal way we consider two tests. One is a sign test (see Diebold and Mariano (1995) for details), the other is the Diebold and Mariano Statistic (again see Diebold and Mariano (1995) for details) to test if the MSE and MAE of two different methods are statistically significantly different (the null being that the forecast performances are similar) — negative values of the test statistics mean that PCC performed better according to the criterion of the test. In tables 6 to 10 we have the results of the tests comparing PCR with PCC (bellow the value of each statistic is the p-value).

Of all the tests applied to each series of forecasts, only once it was concluded that the PCR had a significantly better performance (considering 10% significance level) than PCC – namely when predicting the 6 months inflation, using the GMDC price index, and the MAE criterion to evaluate the performance.

On the other hand we can see that PCC performs significantly better than PCR several times and according to the several tests. For example, when predicting the two years inflation, the PCC performance is always significantly better than PCR, according to the three different statistics (except when we have the sample size of 300). For shorter horizons, like one month or three month inflation forecasts although PCC systematically performs better, only sporadically the better performance is statistically significant. Looking at intermediate horizon forecasts (6 and 12 months), we conclude that about half of the times the difference between the performance of the two methods is statistically significant.

In tables 11 to 15, we can see the results of the same tests comparing PCC with PLS — as before, negative values for the test statistics mean that PCC performed better. PCC was significantly more accurate (considering a significance level of 10%) 81 times while PLS was significantly more precise 19 times. Given these results, it is fair to consider PCC as being the method with the overall best performance.

5 Conclusions

Stock and Watson (1999) considered several forecasting models to predict inflation in the US. Of the several models they considered, PCR was the one with the best performance. In this paper we used this model as a benchmark.

To overcome some of the criticisms to the PCR method, two other methods, which can be applied in similar situations, were proposed:

- the Partial Least Squares, which is very well-known in the Chemometrics literature, and its relation with PCR has already been widely studied, and
- the Principal Components Combination, which tries to overcome the shortcomings of the PCR method by combining this method with the literature on combinations of forecasts. This method is scale invariant with respect to the original explanatory variables, and takes into consideration the explanatory power of each of the explanatory variables when choosing the weights to give to each variable.

The main results of Stock and Watson was reproduced in this paper: PCR leads to significant improvements over the typical AR model, or over the traditional Phillips curve. However the new method PCC outperforms PCR in many cases.

PLS seems to produce better forecasts than PCR for longer horizons (one or two years inflation forecasts), but these results do not carry over to smaller horizons.

PCC outperforms PCR and PLS.

References

- Bai, J. and Ng, S. (2002), Determining the Number of Factors in Approximate Factor Models, Econometrica, 70, 191—221.
- Bates, J. and C.W. J. Granger (1969), The Combination of Forecasts, Operational Research Quarterly, 20, 451—468.
- [3] Bernanke, B and Boivin, J. (2003), Monetary Policy in a Data-Rich Environment, Journal of Monetary Economics, 50, 525—546.
- [4] Chan, Y., Stock, J. and Watson, M. (1999), A Dynamic Factor Model Framework for Forecast Combination, Spanish Economic Review, 1, 91—121.
- [5] Deutsch, M., C. W. J. Granger, and T. Terävirsta (1994), The Combination of Forecasts Using Changing Weights, International Journal of Forecasting, 10, 47—57.

- [6] Diebold, F. and Mariano, R. (1995), Comparing Predictive Accuracy, Journal of Business & Economic Statistics, 13, 134 — 144.
- [7] Forni, M., Hallin, M., Lippi, M. and Reichlin, L. (2000), The Generalized Dynamic Factor Model: Identification and Estimation, Review of Economics and Statistics, 82, 540—554.
- [8] Geweke, J. (1977), The Dynamic Factor Analysis of Economic Time Series, in Aigner and Goldberger, Latent Variables in Socio-Economic Models, Amsterdam: North Holland.
- [9] Granger, C. (1989), Combining Forecasts Twenty Years Later, Journal of Forecasting, 8, 167—173.
- [10] Helland, I. (1988), On the Structure of Partial Least Squares, Communications in Statistics – Simulation and Computation, 17, 581—607.
- [11] Helland, I. (1990), Partial Least Squares Regression and Statistical Models, Scandinavian Journal of Statistics, 17, 97-114.
- [12] Marcellino, M., Stock, J. andWatson, M. (2003), Macroeconomic forecasting in the Euro area: Country specific versus area-wide information, European Economic Review, 47, 1—18.
- [13] Martens, H. (1985), Multivariate Calibration, Doctoral Thesis, Technical University of Norway, Trondheim.
- [14] Gibson. M. and Pritsker, M. (2000), Improving grid-based methods for estimating value at risk of fixed-income portfolios, Finance and Economics Discussion Series 2000-25, Board of Governors of the Federal Reserve System.
- [15] Naik, P. and Tsai, C. (2000), Partial Least Squares for Single-Index Models, Journal of the Royal Statistics Association B, 62, 163—771
- [16] Sargent, T. and Sims, C. (1977), Business Cycle Modeling Without Pretending to Have Too Much A Priori Economic Theory, in C. Sims, New Methods in Business Cycle Research, Minneapolis: Federal Reserve Bank of Minneapolis, 45—109.
- [17] Stock, J., and Watson, M. (1998), Diffusion Indexes, NBER working paper.
- [18] Stock, J., and Watson, M. (1999), Forecasting Inflation, Journal of Monetary Economics, 44, 293—335.

- [19] Stock, J., and Watson, M. (2002), Macroeconomic Forecasting Using Diffusion Indexes, Journal of Business & Economic Statistics, 20, 147 — 162.
- [20] Stone, M. and Brooks, R. (1990), Continuum Regression: Cross-validated Sequentially Constructed Prediction Embracing Ordinary Least Squares, Partials Least Squares and Principal Components Regression, Journal of the Royal Statistics Association B, 52, 237—269
- [21] Wold, H. (1975), Soft Modelling by Latent Variables: the Nonlinear Partial Least Squares Approach, in Perspectives in Probability and Statistics, Papers in Honour of M. S. Bartlett, ed. J. Gani, London: Academic Press.

6 Tables

		GMDC								
		Relati	ve Mear	i Square	Error					
sample size	50	100	150	200	250	300				
PCR	1	1	1	1	1	1				
Phillips	1.007	1.054	1.246	1.242	1.308	1.396				
PLS	1.129	1.153	1.155	1.052	1.025	1.026				
AR	0.947	1.018	1.195	1.244	1.337	1.396				
PCRC	1.003	0.975	0.946	0.886	0.913	0.909				
		Relativ	/e Mean	Absolute	e Error					
PCR	1	1	1	1	1	1				
Phillips	1.021	1.078	1.157	1.123	1.137	1.137				
PLS	1.075	1.113	1.094	1.031	1.035	1.042				
AR	0.989	1.048	1.128	1.127	1.148	1.130				
PCRC	0.987	0.985	0.965	0.941	0.924	0.918				
			PUN	IEW						
		Relati	<u>ve Mear</u>	i Square						
sample size	50	100	150	200	250	300				
PCR	1	1	1	1	1	1				
Phillips	1.071	1.034	1.154	1.181	1.252	1.191				
PLS	0.946	1.013	1.137	1.370	1.342	1.179				
AR	0.883	0.996	1.081	1.117	1.229	1.134				
PCRC	0.948	1.085	0.931	0.929	0.936	0.882				
		Relativ	<u>re Mean</u>	Absolute	Error					
PCR	1	1	1	1	1	1				
Phillips	1.029	1.063	1.137	1.100	1.129	1.106				
PLS	0.996	1.008	1.089	1.173	1.171	1.127				
AR	0.964	1.043	1.092	1.062	1.109	1.059				
PCRC	0.989	1.018	0.960	0.963	0.956	0.965				

Table 1: one month inflation

	GMDC								
		Relati	ve Mear	i Square	Error				
sample size	50	100	150	200	250	300			
PCR	1	1	1	1	1	1			
Phillips	1.311	1.227	1.572	1.544	1.827	1.614			
PLS	0.896	0.936	1.011	0.916	1.207	0.800			
AR	1.165	1.187	1.433	1.512	1.896	1.569			
PCRC	0.993	0.887	0.900	0.899	0.894	0.830			
		Relativ	<u>e Mean</u>	Absolute	e Error				
PCR	1	1	1	1	1	1			
Phillips	1.061	1.224	1.364	1.243	1.393	1.298			
PLS	0.911	1.043	1.046	0.936	1.113	0.952			
AR	1.078	1.204	1.299	1.232	1.402	1.249			
PCRC	0.963	0.977	0.975	0.957	0.926	0.926			
			PUN						
		Relati	<u>ve Mear</u>	<u>i Square</u>					
sample size	50	100	150	200	250	300			
PCR	1	1	1	1	1	1			
Phillips	1.299	1.256	1.521	1.314	1.718	1.770			
PLS	0.849	0.910	1.017	1.076	1.484	1.445			
AR	1.097	1.211	1.335	1.137	1.633	1.560			
PCRC	0.786	0.951	0.992	0.929	0.931	0.797			
		Relativ	<u>re Mean</u>	Absolute	e Error				
PCR	1	1	1	1	1	1			
Phillips	1.103	1.203	1.304	1.161	1.323	1.390			
PLS	0.935	1.012	1.054	1.043	1.193	1.180			
AR	1.058	1.144	1.186	1.087	1.280	1.262			
PCRC	0.889	0.987	1.000	0.984	0.944	0.892			

Table 2: three months inflation

		GMDC								
		Relati	ve Mear	i Square	Error					
sample size	50	100	150	200	250	300				
PCR	1	1	1	1	1	1				
Phillips	1.376	1.475	1.729	1.889	2.160	1.892				
PLS	0.808	1.114	0.896	0.931	1.284	0.873				
AR	1.350	1.483	1.537	1.782	2.224	1.760				
PCRC	0.775	0.882	0.851	0.912	0.914	1.052				
		Relativ	'e Mean	Absolute	Error					
PCR	1	1	1	1	1	1				
Phillips	1.123	1.322	1.366	1.367	1.608	1.436				
PLS	0.889	1.086	0.916	0.931	1.171	0.942				
AR	1.193	1.304	1.291	1.318	1.607	1.358				
PCRC	0.827	0.927	0.927	0.962	0.918	1.006				
			PUN	IEW						
		Relati		<u>i Square</u>						
sample size	50	100	150	200	250	300				
PCR	1	1	1	1	1	1				
Phillips	1.531	1.541	2.159	1.828	2.051	2.180				
PLS	0.814	0.910	1.071	1.009	1.235	1.121				
AR	1.425	1.467	1.778	1.487	1.915	1.827				
PCRC	0.674	0.816	1.120	1.001	0.973	1.003				
		Relativ	<u>'e Mean</u>	Absolute	Error					
PCR	1	1	1	1	1	1				
Phillips	1.184	1.327	1.530	1.426	1.518	1.592				
PLS	0.868	0.936	1.025	1.040	1.144	1.088				
AR	1.156	1.192	1.315	1.263	1.442	1.398				
PCRC	0.813	0.906	1.089	1.085	0.972	1.080				

Table 3: six months inflation

		GMDC								
		Relati	ve Mear	i Square	Error					
sample size	50	100	150	200	250	300				
PCR	1	1	1	1	1	1				
Phillips	2.020	1.583	1.942	2.234	2.369	2.365				
PLS	0.814	0.895	0.746	0.763	0.848	0.704				
AR	1.712	1.614	1.660	1.878	2.353	2.009				
PCRC	0.711	0.569	0.795	0.772	0.685	1.063				
		Relativ	/e Mean	Absolute	Error					
PCR	1	1	1	1	1	1				
Phillips	1.293	1.375	1.465	1.512	1.610	1.552				
PLS	0.921	1.000	0.823	0.863	0.898	0.772				
AR	1.412	1.366	1.340	1.363	1.583	1.385				
PCRC	0.821	0.771	0.914	0.902	0.746	1.005				
			PUN	IEW						
		Relati	<u>ve Mear</u>	i Square	Error					
sample size	50	100	150	200	250	300				
PCR	1	1	1	1	1	1				
Phillips	1.936	1.792	2.373	2.189	2.404	2.234				
PLS	0.788	0.811	0.962	0.879	0.935	0.639				
AR	1.982	1.689	1.829	1.580	2.191	1.719				
PCRC	0.583	0.652	0.936	0.880	0.903	0.805				
		Relativ	<u>re Mean</u>	Absolute	Error					
PCR	1	1	1	1	1	1				
Phillips	1.285	1.349	1.588	1.556	1.707	1.654				
PLS	0.873	0.951	0.985	0.951	1.020	0.793				
AR	1.413	1.280	1.327	1.291	1.599	1.398				
PCRC	0.753	0.838	1.032	1.016	0.906	0.900				

Table 4: twelve months inflation

		GMDC								
		Relati	ve Mean	Square	Error					
sample size	50	100	150	200	250	300				
PCR	1	1	1	1	1	1				
Phillips	1.619	1.828	1.979	2.670	2.802	3.076				
PLS	0.714	0.693	0.747	0.429	0.504	0.602				
AR	2.080	1.872	1.719	1.976	2.722	2.500				
PCRC	0.553	0.585	0.711	0.665	0.498	1.024				
		Relativ	/e Mean	Absolute	e Error					
PCR	1	1	1	1	1	1				
Phillips	1.331	1.353	1.489	1.712	1.687	1.793				
PLS	0.888	0.851	0.770	0.634	0.681	0.788				
AR	1.576	1.376	1.353	1.458	1.643	1.597				
PCRC	0.758	0.773	0.798	0.822	0.668	1.030				
				0.022						
			PUN							
			PUN							
sample size	50		PUN	IEW		300				
	50 1	Relati 100 1	PUN ve Mear 150 1	IEW Square 200 1	Error 250 1	300 1				
sample size PCR Phillips	50 1 1.615	Relati 100 1 1.967	PUN ve Mean 150 1.984	IEW Square 200 1 3.079	Error 250 1 3.146	300 1 4.159				
sample size PCR	50 1	Relati 100 1	PUN ve Mear 150 1	IEW Square 200 1	Error 250 1	300 1				
sample size PCR Phillips PLS AR	50 1 1.615	Relati 100 1 1.967	PUN ve Mean 150 1.984	IEW Square 200 1 3.079	Error 250 1 3.146	300 1 4.159				
sample size PCR Phillips PLS	50 1 1.615 0.703	Relati 100 1.967 0.711	PUN ve Mean 150 1 1.984 0.802	IEW Square 200 1 3.079 0.707	Error 250 1 3.146 0.683	300 1 4.159 0.823				
sample size PCR Phillips PLS AR	50 1 1.615 0.703 2.345	Relati 100 1 1.967 0.711 1.928 0.580	PUN ve Mean 150 1.984 0.802 1.557 0.831	IEW 200 1 3.079 0.707 1.993	Error 250 1 3.146 0.683 2.893 0.680	300 1 4.159 0.823 3.182				
sample size PCR Phillips PLS AR	50 1 1.615 0.703 2.345 0.519	Relati 100 1.967 0.711 1.928 0.580 Relativ 1	PUN ve Mean 150 1.984 0.802 1.557 0.831 ve Mean 1	IEW Square 200 1 3.079 0.707 1.993 0.766 Absolute 1	Error 250 1 3.146 0.683 2.893 0.680 e Error	300 1 4.159 0.823 3.182 1.018 1.018				
sample size PCR Phillips PLS AR PCRC PCR PCR	50 1 1.615 0.703 2.345 0.519 1 1.343	Relati 100 1 1.967 0.711 1.928 0.580	PUN ve Mean 150 1.984 0.802 1.557 0.831 ve Mean 1.479	IEW Square 200 1 3.079 0.707 1.993 0.766 Absolute 1 1.828	Error 250 1 3.146 0.683 2.893 0.680	300 1 4.159 0.823 3.182				
sample size PCR Phillips PLS AR PCRC PCR Phillips PLS	50 1 1.615 0.703 2.345 0.519	Relati 100 1.967 0.711 1.928 0.580 Relativ 1	PUN ve Mean 150 1.984 0.802 1.557 0.831 ve Mean 1	IEW Square 200 1 3.079 0.707 1.993 0.766 Absolute 1	Error 250 1 3.146 0.683 2.893 0.680 e Error	300 1 4.159 0.823 3.182 1.018 1.018				
sample size PCR Phillips PLS AR PCRC PCR PCR	50 1 1.615 0.703 2.345 0.519 1 1.343	Relati 100 1 1.967 0.711 1.928 0.580 Relativ 1 1.377	PUN ve Mean 150 1.984 0.802 1.557 0.831 ve Mean 1.479	IEW Square 200 1 3.079 0.707 1.993 0.766 Absolute 1 1.828	Error 250 1 3.146 0.683 2.893 0.680 e Error 1 1.898	300 1 4.159 0.823 3.182 1.018 1.018 1 2.133				

Table 5: two years inflation

<u></u>	<u>e one month inflation forecasts — PCC vs PCR</u>								
		GMDC							
sample size	50	100	150	200	250	300			
Sign statistic	0.641	-1.270	-0.506	-1.556	-1.796	-0.600			
p-value	0.261	0.102	0.306	0.060	0.036	0.274			
DM statistic (MSE)	0.049	-0.342	-0.640	-1.268	-0.864	-0.727			
p-value	0.481	0.366	0.261	0.102	0.194	0.234			
DM statistic (MAE)	-0.420	-0.392	-1.065	-1.616	-1.794	-1.391			
p-value	0.337	0.347	0.144	0.053	0.036	0.082			
			PUN	IEW					
Sign statistic	-0.855	-0.115	-0.759	-1.697	-1.306	-1.200			
p-value	0.196	0.454	0.224	0.045	0.096	0.115			
DM statistic (MSE)	-0.674	1.069	-1.086	-0.904	-0.816	-1.524			
p-value	0.250	0.142	0.139	0.183	0.207	0.064			
DM statistic (MAE)	-0.284	0.472	-1.142	-0.953	-1.053	-0.739			
p-value	0.388	0.319	0.127	0.170	0.146	0.230			

Table 6: tests for the one month inflation forecasts — PCC vs PCB

Table 7: tests for the three months inflation forecasts — PCC vs PCR

		GMDC							
sample size	50	100	150	200	250	300			
Sign statistic	-1.608	0.000	-0.127	-1.137	-1.644	-0.202			
p-value	0.054	0.500	0.449	0.128	0.050	0.420			
DM statistic (MSE)	-0.077	-0.904	-1.098	-1.372	-1.163	-1.061			
p-value	0.469	0.183	0.136	0.085	0.122	0.144			
DM statistic (MAE)	-0.943	-0.448	-0.510	-1.048	-1.387	-0.888			
p-value	0.173	0.327	0.305	0.147	0.083	0.187			
			PUN	JEW					
Sign statistic	-1.823	0.116	0.000	-0.711	-1.151	-1.818			
p-value	0.034	0.454	0.500	0.239	0.125	0.035			
DM statistic (MSE)	-2.006	-0.598	-0.081	-0.751	-0.609	-1.404			
p-value	0.022	0.275	0.468	0.226	0.271	0.080			
DM statistic (MAE)	-2.258	-0.306	0.009	-0.290	-0.884	-1.381			
p-value	0.012	0.380	0.497	0.386	0.188	0.084			

Table 6. tests for the	$\frac{1}{100}$ SIX months milation forecasts – 1 CC vs 1 CN							
			GM	DC				
sample size	50	100	150	200	250	300		
Sign statistic	-4.899	-2.154	-0.447	-0.645	-1.080	0.103		
p-value	0.000	0.016	0.327	0.260	0.140	0.459		
DM statistic (MSE)	-2.027	-0.631	-1.526	-0.941	-0.714	0.394		
p-value	0.021	0.264	0.063	0.173	0.238	0.347		
DM statistic (MAE)	-3.116	-1.049	-1.227	-0.614	-0.913	0.060		
p-value	0.001	0.147	0.110	0.270	0.181	0.476		
			PUN	IEW				
Sign statistic	-3.499	-2.154	0.831	0.931	-0.415	0.718		
p-value	0.000	0.016	0.203	0.176	0.339	0.236		
DM statistic (MSE)	-2.539	-1.544	1.185	0.010	-0.231	0.021		
p-value	0.006	0.061	0.118	0.496	0.409	0.492		
DM statistic (MAE)	-3.048	-1.412	1.297	1.133	-0.398	0.851		
p-value	0.001	0.079	0.097	0.129	0.345	0.197		

Table 8: tests for the six months inflation forecasts — PCC vs PCR

Table 9: tests for the twelve months inflation forecasts — PCC vs PCR

		GMDC							
sample size	50	100	150	200	250	300			
Sign statistic	-2.879	-4.294	-1.100	-2.255	-3.986	0.742			
p-value	0.002	0.000	0.136	0.012	0.000	0.229			
DM statistic (MSE)	-2.625	-1.491	-1.622	-1.481	-2.371	0.321			
p-value	0.004	0.068	0.052	0.069	0.009	0.374			
DM statistic (MSE)	-3.694	-2.100	-1.534	-1.403	-2.548	0.040			
p-value	0.000	0.018	0.063	0.080	0.005	0.484			
	PUNEW								
Sign statistic	-4.182	-3.000	1.100	0.800	-1.272	-1.378			
p-value	0.000	0.001	0.136	0.212	0.102	0.084			
DM statistic (MSE)	-3.975	-1.520	-0.520	-0.643	-0.596	-0.914			
p-value	0.000	0.064	0.301	0.260	0.276	0.180			
DM statistic (MSE)	-4.669	-1.766	0.476	0.141	-0.763	-1.053			
p-value	0.000	0.039	0.317	0.444	0.223	0.146			

		GMDC							
sample size	50	100	150	200	250	300			
Sign statistic	-5.807	-4.867	-4.580	-2.330	-4.348	0.798			
p-value	0.000	0.000	0.000	0.010	0.000	0.213			
DM statistic (MSE)	-5.095	-2.217	-2.198	-2.477	-5.801	0.272			
p-value	0.000	0.013	0.014	0.007	0.000	0.393			
DM statistic (MSE)	-6.123	-3.071	-3.286	-3.445	-4.309	0.675			
p-value	0.000	0.001	0.001	0.000	0.000	0.250			
			PUN	JEW					
Sign statistic	-6.470	-5.468	-3.252	-2.931	-1.863	-0.114			
p-value	0.000	0.000	0.001	0.002	0.031	0.455			
DM statistic (MSE)	-2.817	-2.240	-2.607	-1.683	-2.005	0.598			
p-value	0.002	0.013	0.005	0.046	0.022	0.275			
DM statistic (MSE)	-4.482	-10.55	-2.507	-1.457	-1.506	-0.099			
p-value	0.000	0.000	0.006	0.073	0.066	0.461			

Table 10: tests for the two years inflation forecasts — PCC vs PCR

Table 11: tests for the one month inflation forecasts — PCC vs PLS

		GMDC								
sample size	50	100	150	200	250	300				
Sign statistic	-3.100	-3.233	-2.403	-1.414	-1.633	-1.000				
p-value	0.001	0.001	0.008	0.079	0.051	0.159				
DM statistic (MSE)	-2.161	-2.406	-2.550	-1.807	-1.023	-1.110				
p-value	0.015	0.008	0.005	0.035	0.153	0.134				
DM statistic (MAE)	-2.863	-3.444	-3.020	-1.798	-1.906	-1.829				
p-value	0.002	0.000	0.001	0.036	0.028	0.034				
			PUN	JEW						
Sign statistic	-1.604	-1.386	-2.403	-4.101	-3.266	-1.400				
p-value	0.054	0.083	0.008	0.000	0.001	0.081				
DM statistic (MSE)	0.504	0.926	-2.542	-3.940	-3.496	-2.311				
p-value	0.307	0.177	0.006	0.000	0.000	0.010				
DM statistic (MAE)	-0.010	0.065	-2.758	-3.652	-3.513	-2.380				
p-value	0.496	0.474	0.003	0.000	0.000	0.009				

		GMDC							
sample size	50	100	150	200	250	300			
Sign statistic	0.322	-1.738	0.000	0.711	-2.466	0.000			
p-value	0.374	0.041	0.500	0.239	0.007	0.500			
DM statistic (MSE)	1.055	-0.621	-1.080	-0.163	-1.760	0.229			
p-value	0.146	0.267	0.140	0.435	0.039	0.409			
DM statistic (MAE)	1.216	-1.524	-1.232	0.352	-2.205	-0.275			
p-value	0.112	0.064	0.109	0.362	0.014	0.392			
			PUN	JEW					
Sign statistic	-1.501	-3.244	-0.381	-0.284	-3.124	-2.424			
p-value	0.067	0.001	0.352	0.388	0.001	0.008			
DM statistic (MSE)	-0.841	0.627	-0.331	-1.476	-3.345	-3.067			
p-value	0.200	0.265	0.370	0.070	0.000	0.001			
DM statistic (MAE)	-1.130	-0.646	-1.014	-0.928	-2.984	-2.798			
p-value	0.129	0.259	0.155	0.177	0.001	0.003			

Table 12: tests for the three months inflation forecasts — PCC vs PLS

Table 13: tests for the six months inflation forecasts — PCC vs PLS

	GMDC						
sample size	50	100	150	200	250	300	
Sign statistic	-1.669	-3.086	0.958	0.501	-3.239	1.949	
p-value	0.048	0.001	0.169	0.308	0.001	0.026	
DM statistic (MSE)	-0.562	-1.490	-0.307	-0.127	-2.047	0.670	
p-value	0.287	0.068	0.379	0.449	0.020	0.251	
DM statistic (MAE)	-1.565	-2.408	0.155	0.404	-2.388	0.361	
p-value	0.059	0.008	0.438	0.343	0.008	0.359	
	PUNEW						
Sign statistic	-0.700	-1.572	1.086	1.217	-1.246	1.129	
p-value	0.242	0.058	0.139	0.112	0.106	0.130	
DM statistic (MSE)	-1.573	-0.948	0.285	-0.049	-1.731	-0.451	
p-value	0.058	0.172	0.388	0.480	0.042	0.326	
DM statistic (MAE)	-0.953	-0.540	0.723	0.454	-1.860	-0.065	
p-value	0.170	0.295	0.235	0.325	0.031	0.474	

	GMDC						
sample size	50	100	150	200	250	300	
Sign statistic	-1.684	-5.588	2.652	0.800	-2.120	3.922	
p-value	0.046	0.000	0.004	0.212	0.017	0.000	
DM statistic (MSE)	-1.570	-2.536	0.577	0.053	-1.784	1.462	
p-value	0.058	0.006	0.282	0.479	0.037	0.072	
DM statistic (MAE)	-1.765	-3.332	1.312	0.380	-2.142	1.430	
p-value	0.039	0.000	0.095	0.352	0.016	0.076	
	PUNEW						
Sign statistic	-3.313	-2.529	1.488	0.364	-1.612	1.378	
p-value	0.000	0.006	0.068	0.358	0.054	0.084	
DM statistic (MSE)	-2.840	-1.450	-0.305	0.007	-0.184	0.745	
p-value	0.002	0.074	0.380	0.497	0.427	0.228	
DM statistic (MAE)	-2.219	-2.155	0.774	1.445	-0.844	0.598	
p-value	0.013	0.016	0.220	0.074	0.199	0.275	

Table 14: tests for the twelve months inflation forecasts — PCC vs PLS

Table 15: tests for the two years inflation forecasts — PCC vs PLS

	GMDC						
sample size	50	100	150	200	250	300	
Sign statistic	-4.479	-1.742	1.792	3.533	-0.266	4.900	
p-value	0.000	0.041	0.037	0.000	0.395	0.000	
DM statistic (MSE)	-1.674	-1.127	-0.361	2.387	-0.046	2.220	
p-value	0.047	0.130	0.359	0.008	0.481	0.013	
DM statistic (MAE)	-1.729	-1.082	0.383	2.387	-0.152	2.247	
p-value	0.042	0.140	0.351	0.009	0.439	0.012	
	PUNEW						
Sign statistic	-4.258	-3.305	1.261	2.029	0.089	2.165	
p-value	0.000	0.000	0.104	0.021	0.465	0.015	
DM statistic (MSE)	-1.903	-1.190	0.361	0.858	-0.034	0.814	
p-value	0.029	0.117	0.359	0.195	0.486	0.208	
DM statistic (MAE)	-2.308	-2.04	0.489	1.520	0.070	0.780	
p-value	0.011	0.021	0.312	0.064	0.472	0.218	