

NDK_PCR_PRFTest

Last Modified on 03/14/2016 11:39 am CDT

- [C/C++](#)
- [.Net](#)

```
int __stdcall NDK_PCR_PRFTest ( double ** X,  
                               size_t   nXSize,  
                               size_t   nXVars,  
                               double *  Y,  
                               size_t   nYSize,  
                               double   intercept,  
                               LPBYTE   mask1,  
                               size_t   nMaskLen1,  
                               LPBYTE   mask2,  
                               size_t   nMaskLen2,  
                               double   alpha,  
                               WORD     nRetType,  
                               double *  retVal  
                               )
```

Returns an array of cells for the i-th principal component (or residuals).

Returns

status code of the operation

Return values

NDK_SUCCESS Operation successful

NDK_FAILED Operation unsuccessful. See [Macros](#) for full list.

Parameters

- | | |
|-----------------------|--|
| [in] X | is the independent variables data matrix, such that each column represents one variable |
| [in] nXSize | is the number of observations (i.e. rows) in X |
| [in] nXVars | is the number of variables (i.e. columns) in X |
| [in] Y | is the response or the dependent variable data array (one dimensional array) |
| [in] nYSize | is the number of elements in Y |
| [in] intercept | is the constant or the intercept value to fix (e.g. zero). If missing (NaN), an intercept will not be fixed and is computed normally |

- [in] **mask1** is the boolean array to select a subset of the input variables in X. If missing (i.e. NULL), all variables in X are included.
- [in] **nMaskLen1** is the number of elements in mask1
- [in] **mask2** is the boolean array to select a subset of the input variables in X. If missing (i.e. NULL), all variables in X are included.
- [in] **nMaskLen2** is the number of elements in mask2
- [in] **alpha** is the statistical significance of the test (i.e. alpha)
- [in] **nRetType** is a switch to select the return output (1 = P-Value (default), 2 = Test Stats, 3 = Critical Value.)
- [out] **retVal** is the calculated test statistics/

Remarks

1. The underlying model is described [here](#).
2. Model 1 must be a sub-model of Model 2. In other words, all variables included in Model 1 must be included in Model 2.
3. The coefficient of determination (i.e. (R^2)) increases in value as we add variables to the regression model, but we often wish to test whether the improvement in R square by adding those variables is statistically significant.
4. To do so, we developed an inclusion/exclusion test for those variables. First, let's start with a regression model with (K_1) variables: $Y_t = \alpha + \beta_1 X_1 + \dots + \beta_{K_1} X_{K_1}$ Now, let's add few more variables $(\left(X_{K_1+1} \dots X_{K_2}\right))$ $Y_t = \alpha + \beta_1 X_1 + \dots + \beta_{K_1} X_{K_1} + \dots + \beta_{K_1+1} X_{K_1+1} + \dots + \beta_{K_2} X_{K_2}$
5. The test of hypothesis is as follows: $[H_0 : \beta_{K_1+1} = \beta_{K_1+2} = \dots = \beta_{K_2} = 0]$ $[H_1 : \exists \beta_i \neq 0, i \in \left[K_1+1 \dots K_2\right)]$
6. Using the change in the coefficient of determination (i.e. (R^2)) as we added new variables, we can calculate the test statistics: $\left[\mathrm{f} = \frac{(R^2_f - R^2_r)/(K_2 - K_1)}{(1 - R^2_f)/(N - K_2 - 1)} \sim \mathrm{F}_{K_2 - K_1, N - K_2 - 1}\right]$ Where:
 - (R^2_f) is the (R^2) of the full model (with added variables).
 - (R^2_r) is the (R^2) of the reduced model (without the added variables).
 - (K_1) is the number of variables in the reduced model.
 - (K_2) is the number of variables in the full model.
 - (N) is the number of observations in the sample data.
7. The sample data may include missing values.
8. Each column in the input matrix corresponds to a separate variable.
9. Each row in the input matrix corresponds to an observation.
10. Observations (i.e. row) with missing values in X or Y are removed.
11. The number of rows of the response variable (Y) must be equal to the number of rows of the explanatory variables (X).
12. The MLR_ANOVA function is available starting with version 1.60 APACHE.

Requirements

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References

Hamilton, J .D.; [Time Series Analysis](#), Princeton University Press (1994), ISBN 0-691-04289-6
Tsay, Ruey S.; [Analysis of Financial Time Series](#) John Wiley & SONS. (2005), ISBN 0-471-690740

See Also

[template("related")]