

0.1 threesls: Three Stage Least Squares

threesls is a combination of two stage least squares and seemingly unrelated regression. It provides consistent estimates for linear regression models with explanatory variables correlated with the error term. It also extends ordinary least squares analysis to estimate system of linear equations with correlated error terms

Syntax

```
> fml <- list(mu1 = Y1 ~ X1 + Z1, mu2 = Y2 ~ X2 + Z2, inst1 = Z1 ~  
+           W1 + X1, inst2 = Z2 ~ W2 + X2)  
  
> z.out <- zelig(formula = fml, model = "threesls", data = mydata)  
> x.out <- setx(z.out)  
> s.out <- sim(z.out, x = x.out)
```

Inputs

threesls regression specification requires at least two sets of equations. The first set of M equations corresponds to the M dependent variables (Y_1, \dots, Y_M) to be estimated. The second set of equations (Z) corresponds to the instrumental variables in the M equations.

- **formula**: a list of the system of equations and instrumental variable equations. The system of equations is listed first as **mus**. The equations for the instrumental variables are listed next as **insts**. For example:

```
> fml <- list(mu1 = Y1 ~ X1 + Z1, mu2 = Y2 ~ X2 + Z2, inst1 = Z1 ~  
+           W1 + X1, inst2 = Z2 ~ W2 + X2)
```

"**mu1**" is the first equation in the two equation model with **Y1** as the dependent variable and **X1** and **Z1** as the explanatory variables. "**mu2**" is the second equation with **Y2** as the dependent variable and **X2** and **Z2** as the explanatory variables. **Z1** and **Z2** are also problematic endogenous variables, so they are estimated through instruments in the "**inst1**" and "**inst2**" equations.

- **Y**: dependent variables of interest in the system of equations.
- **Z**: the problematic explanatory variables correlated with the error term.
- **W**: exogenous instrument variables used to estimate the problematic explanatory variables (**Z**)

Additional Inputs

threesls takes the following additional inputs for model specifications:

- **TX**: an optional matrix to transform the regressor matrix and, hence, also the coefficient vector (see details). Default is **NULL**.
- **maxiter**: maximum number of iterations.
- **tol**: tolerance level indicating when to stop the iteration.
- **rcovformula**: formula to calculate the estimated residual covariance matrix (see details). Default is equal to 1.
- **formulathreesls**: formula for calculating the threesls estimator, one of “GLS”, “IV”, “GMM”, “Schmidt”, or “Eviews” (see details.)
- **probdfsys**: use the degrees of freedom of the whole system (in place of the degrees of freedom of the single equation to calculate probability values for the t-test of individual parameters.
- **single.eq.sigma**: use different σ^2 for each single equation to calculate the covariance matrix and the standard errors of the coefficients.
- **solvetol**: tolerance level for detecting linear dependencies when inverting a matrix or calculating a determinant. Default is **solvetol=Machine\$double.eps**.
- **saveMemory**: logical. Save memory by omitting some calculation that are not crucial for the basic estimate (e.g McElroy’s R^2).

Details

The matrix **TX** transforms the regressor matrix (X) by $X^* = X \times TX$. Thus, the vector of coefficients is now $b = TX \times b^*$ where b is the original(stacked) vector of all coefficients and b^* is the new coefficient vector that is estimated instead. Thus, the elements of vector b and $b_i = \sum_j TX_{ij} \times b_j^*$. The TX matrix can be used to change the order of the coefficients and also to restrict coefficients (if TX has less columns than it has rows). If iterated (with **maxit**>1), the coverage criterion is

$$\sqrt{\frac{\sum_i (b_{i,g} - b_{i,g-1})^2}{\sum_i b_{i,g-1}^2}} < tol$$

where $b_{i,g}$ is the i th coefficient of the g th iteration step. The formula (**rcovformula** to calculate the estimated covariance matrix of the residuals($\hat{\Sigma}$) can be one of the following (see Judge et al., 1955, p.469): if **rcovformula**= 0:

$$\hat{\sigma}_{ij} = \frac{\hat{e}_i \hat{e}_j}{T}$$

if `rcovformula= 1` or `rcovformula='geomean'`:

$$\hat{\sigma}_{ij} = \frac{\hat{e}_i' \hat{e}_j}{\sqrt{(T - k_i) \times (T - k_j)}}$$

if `rcovformula= 2` or `rcovformula='Theil'`:

$$\hat{\sigma}_{ij} = \frac{\hat{e}_i' \hat{e}_j}{T - k_i - k_j + \text{tr}[X_i(X_i'X_i)^{-1}X_i'X_j(X_j'X_j)^{-1}X_j']}$$

if `rcovformula= 3` or `rcovformula='max'`:

$$\hat{\sigma}_{ij} = \frac{\hat{e}_i' \hat{e}_j}{T - \max(k_i, k_j)}$$

If $i = j$, formula 1, 2, and 3 are equal. All these three formulas yield unbiased estimators for the diagonal elements of the residual covariance matrix. If *ineqj*, only formula 2 yields an unbiased estimator for the residual covariance matrix, but it is not necessarily positive semidefinite. Thus, it is doubtful whether formula 2 is really superior to formula 1 (Theil, 1971, p.322). The formulas to calculate the three-sls estimator lead to identical results if the same instruments are used in all equations. If different instruments are used in the different equations, only the GMM-three-sls estimator (“GMM”) and the three-sls estimator proposed by Schmidt (1990) (“Schmidt”) are consistent, whereas “GMM” is efficient relative to “Schmidt” (see Schmidt, 1990).

Examples

Attaching the example dataset:

```
> data(kmenta)
```

Formula:

```
> formula <- list(mu1 = q ~ p + d, mu2 = q ~ p + f + a, inst = ~d +
+       f + a)
```

Estimating the model using `threesls`:

```
> z.out <- zelig(formula = formula, model = "threesls", data = kmenta)
> summary(z.out)
```

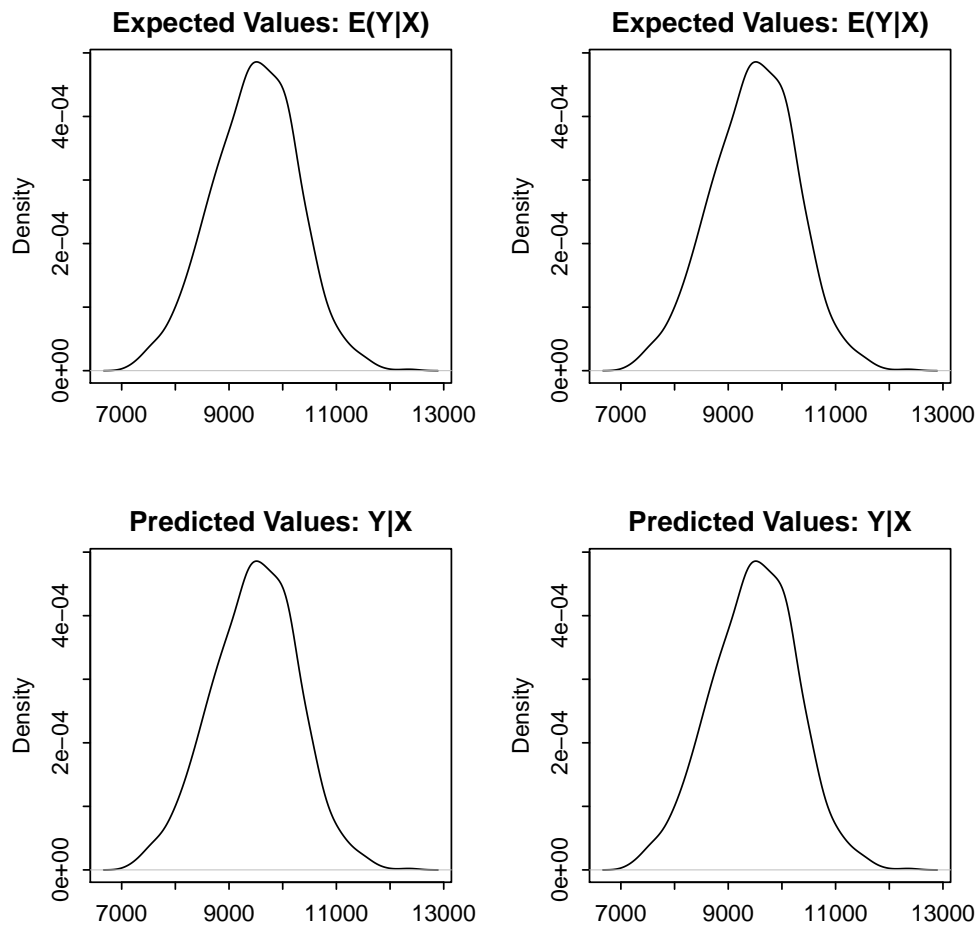
Set explanatory variables to their default (mean/mode) values

```
> x.out <- setx(z.out)
```

Simulate draws from the posterior distribution:

```
> s.out <- sim(z.out, x = x.out)
> summary(s.out)
```

Plot the quantities of interest



Model

See Also

For information about two stage least square regression, see Section ?? and `help(2sls)`. For information about seemingly unrelated regression, see Section ?? and `help(sur)`.

Quantities of Interest

Output Values

The output of each Zelig command contains useful information which you may view. For example, if you run:

```
z.out <- zelig(formula=fml, model = "threesls", data)
```

then you may examine the available information in `z.out` by using `names(z.out)`, see the draws from the posterior distribution of the `coefficients` by using `z.out$coefficients`, and view a default summary of information through `summary(z.out)`. Other elements available through the `$` operator are listed below:

- `rcovest`: residual covariance matrix used for estimation.
- `mcelr2`: McElroys R-squared value for the system.
- `h`: matrix of all (diagonally stacked) instrumental variables.
- `formulathreesls`: formula for calculating the threesls estimator
- `method`: Estimation method.
- `g`: number of equations.
- `n`: total number of observations.
- `k`: total number of coefficients.
- `ki`: total number of linear independent coefficients.
- `df`: degrees of freedom of the whole system.
- `iter`: number of iteration steps.
- `b`: vector of all estimated coefficients.
- `t`: t values for b .
- `se`: estimated standard errors of b .
- `bt`: coefficient vector transformed by TX .

- `p`: p values for b .
- `bcov`: estimated covariance matrix of b .
- `btcov`: covariance matrix of bt .
- `rcov`: estimated residual covariance matrix.
- `drcov`: determinant of `rcov`.
- `rcor`: estimated residual correlation matrix.
- `olsr2`: system OLS R-squared value.
- `y`: vector of all (stacked) endogenous variables.
- `x`: matrix of all (diagonally stacked) regressors.
- `data`: data frame of the whole system (including instruments).
- `TX`: matrix used to transform the regressor matrix.
- `rcovformula`: formula to calculate the estimated residual covariance matrix.
- `probdffsys`: system degrees of freedom to calculate probability values?.
- `solvetol`: tolerance level when inverting a matrix or calculating a determinant.
- `eq`: a list that contains the results that belong to the individual equations.
- `eqnlabel*`: the equation label of the i th equation (from the labels list).
- `formula*`: model formula of the i th equation.
- `n*`: number of observations of the i th equation.
- `k*`: number of coefficients/regressors in the i th equation (including the constant).
- `ki*`: number of linear independent coefficients in the i th equation (including the constant differs from k only if there are restrictions that are not cross equation).
- `df*`: degrees of freedom of the i th equation.
- `b*`: estimated coefficients of the i th equation.
- `se*`: estimated standard errors of b of the i th equation.
- `t*`: t values for b of the i th equation.
- `p*`: p values for b of the i th equation.

- `covb*`: estimated covariance matrix of b of the i th equation.
- `y*`: vector of endogenous variable (response values) of the i th equation.
- `x*`: matrix of regressors (model matrix) of the i th equation.
- `data*`: data frame (including instruments) of the i th equation.
- `fitted*`: vector of fitted values of the i th equation.
- `residuals*`: vector of residuals of the i th equation.
- `ssr*`: sum of squared residuals of the i th equation.
- `mse*`: estimated variance of the residuals (mean of squared errors) of the i th equation.
- `s2*`: estimated variance of the residuals ($\hat{\sigma}^2$) of the i th equation.
- `rmse*`: estimated standard error of the residuals (square root of mse) of the i th equation.
- `s*`: estimated standard error of the residuals ($\hat{\sigma}$) of the i th equation.
- `r2*`: R-squared (coefficient of determination).
- `adjr2*`: adjusted R-squared value.
- `inst*`: instruments of the i th equation.
- `h*`: matrix of instrumental variables of the i th equation.
- `zelig.data`: the input data frame if `save.data = TRUE`.

How to Cite

To cite the *threesls* Zelig model:

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To cite Zelig as a whole, please reference these two sources:

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Imai, Kosuke, Gary King, and Olivia Lau. (2008). "Toward A Common Framework for Statistical Analysis and Development." *Journal of Computational and Graphical Statistics*, Vol. 17, No. 4 (December), pp. 892-913.

See also

The *threesls* function is adapted from the `systemfit` library (Hamann and Henningsen 2005).

Bibliography

Hamann, J. and Henningsen, A. (2005), *systemfit: Simultaneous Equation Systems in R Package*.