

# Package: pgsc (via r-universe)

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**Type** Package

**Title** Computes Powell's Generalized Synthetic Control Estimator

**Version** 1.0.0

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**Author** Philip Barrett

**Maintainer** Philip Barrett <pobarrett@gmail.com>

**Description** Computes the generalized synthetic control estimator described in Powell (2017) <doi:10.7249/WR1142>. Provides both point estimates, and hypothesis testing.

**License** GPL-2

**Depends** R (>= 2.10), Rcpp (>= 0.12.18), nloptr, reshape2

**LinkingTo** Rcpp, RcppArmadillo

**RoxygenNote** 6.1.0

**Suggests** knitr, rmarkdown, plm

**VignetteBuilder** knitr

**URL** <https://github.com/philipbarrett/pgsc>

**Encoding** UTF-8

**Repository** <https://philipbarrett.r-universe.dev>

**RemoteUrl** <https://github.com/philipbarrett/pgsc>

**RemoteRef** HEAD

**RemoteSha** 2abfdd9ceaa3470a36dd024450a959ceff17d0b6

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 pgsc

*Wrapper function for GSC estimation*


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### Description

Wrapper function for GSC estimation

### Usage

```
pgsc(dta, dep.var, indep.var, b.init, method, sol.it = NULL,
     wt.init = NULL, print.level = 0, g.i = NULL, g.i.grad = NULL,
     ...)
```

### Arguments

<code>dta</code>	A data frame
<code>dep.var</code>	A string defining the dependent variable
<code>indep.var</code>	A vector of strings defining the independent (treatment) variables
<code>b.init</code>	An initial value for the treatment variable coefficients. Must have same length as ‘ <code>indep.var</code> ’
<code>method</code>	The GSC iteration method to be used. Must be one of: <ul style="list-style-type: none"> <li>• <code>onestep</code>: "Plain" GSC solution, without weights</li> <li>• <code>twostep.aggte</code>: Observations weighted by unit MSEs from the one-step solution.</li> <li>• <code>twostep.indiv</code>: Observations weighted by unit MSEs from individual, unit-by-unit unweighted solutions.</li> </ul>
<code>sol.it</code>	The first step solution used in the two-step methods. If omitted, a new one-step solution is computed.
<code>wt.init</code>	An initial value for the weighting matrix
<code>print.level</code>	The level of detail provided in the printed output
<code>g.i</code>	A function defining a restriction on the parameters. Used in hypothesis testing.
<code>g.i.grad</code>	The gradient of <code>g.i</code> .
<code>...</code>	Other arguments to be passed to the optimization

### Details

See the vignette "Using pgsc" for an extended example.

**Value**

Returns the point estimate of the model as a gsc object, a list with entries:

**b** The point estimate of the coefficients on the dependent variables

**diff** The difference between successive iterations

**err** The maximum error on the within-iteration optimization problems

**it** Number of iterations require to solve

**sig.i** The unit-specific MSEs from the solution

**W** The "full" weighting matrix for counterfactuals, containing own-unit weights (all zero) and unit-N weights

**wt** The "minimal" weighting matrix, omitting own-unit weights and weights on unit N (which can be computed as one-minus-rowsum)

**Examples**

```
data("pgsc.dta")
sol <- pgsc(pgsc.dta, dep.var = 'y', indep.var = c('D1','D2'),
b.init = c(0,0), method='onestep' )
summary(sol)
g.i <- function(b) b[1] ; g.i.grad <- function(b) c(1,0)
sol.r <- pgsc(pgsc.dta, dep.var = 'y', indep.var = c('D1','D2'),
b.init = sol$b, method='onestep', g.i=g.i, g.i.grad=g.i.grad )
summary(sol.r)
```

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pgsc.dta

*Synthetic data for PGSC testing.*

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**Description**

A dataset with an outcome given by a treatment and a set of factors.

**Usage**

pgsc.dta

**Format**

A data frame with 750 rows and 8 variables:

**n** The unit, here labeled as a US state

**t** The time period

**y** The outcome variable

**D1** The first treatment variable

**D2** The second treatment variable

**X1** The first observed confounding factor

**X2** The second observed confounding factor

**X3** The third observed confounding factor

**Source**

Generated by code in the package vignette "Using pgsc".

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pgsc.wald.test	<i>A wrapper for the wald test of a restricted solution</i>
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**Description**

A wrapper for the wald test of a restricted solution

**Usage**

```
pgsc.wald.test(dta, dep.var, indep.var, sol.rest, n.boot = 10000,
  seed = 42)
```

**Arguments**

dta	A dataframe
dep.var	A vector of strings of names of dependent variables.
indep.var	A vector of strings of names of independent (treatment) variables.
sol.rest	A restricted solution which is being tested
n.boot	The number of bootstrapped samples for the variance calculation. Default is 10000.
seed	Randomization seed. Default is 42.

**Details**

See the vignette "Using pgsc" for an extended example.

**Value**

Returns the wald test as gsc.wald object, a list with entries:

**b** The point estimate of the coefficients on the dependent variables

**S** The Wald statistic

**s.boot** The bootstrapped Wald statistic

**p.value** The p-value for the Wald statistic.

**Examples**

```
data("pgsc.dta")
g.i <- function(b) b[1] ; g.i.grad <- function(b) c(1,0)
sol.r <- pgsc(pgsc.dta, dep.var = 'y', indep.var = c('D1','D2'),
  b.init = c(0,1), method='onestep', g.i=g.i, g.i.grad=g.i.grad )
wald <- pgsc.wald.test( pgsc.dta, 'y', indep.var = c('D1','D2'), sol.r )
summary(wald)
plot(wald)
```

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