

# Package ‘did2s’

September 27, 2021

**Title** Two-Stage Difference-in-Differences Following Gardner (2021)

**Version** 0.4.0

**Description** Estimates Two-way Fixed Effects difference-in-differences/event-study models using the approach proposed by Gardner (2021). To avoid the problems caused by OLS estimation of the Two-way Fixed Effects model, this function first estimates the fixed effects and covariates using untreated observations and then in a second stage, estimates the treatment effects.

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**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**Depends** R (>= 2.10), fixest (>= 0.9.0)

**Imports** dplyr, magrittr, ggplot2, glue, stats, stringr, rlang, tibble, tidy, rsample, purrr, cli, Matrix, Rcpp, did, broom, methods, staggered, didimputation, gt

**URL** <http://kylebutts.com/did2s/>

**Suggests** rmarkdown, knitr, haven, testthat (>= 3.0.0)

**VignetteBuilder** knitr

**LinkingTo** RcppArmadillo, Rcpp

**Config/testthat/edition** 3

**NeedsCompilation** yes

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castle	<i>Data from Cheng and Hoekstra (2013)</i>
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### Description

State-wide panel data from 2000-2010 that has information on castle-doctrine, the so-called "stand-your-ground" laws that were implemented by 20 states.

### Usage

```
castle
```

### Format

A data frame with 550 rows and 5 variables:

**sid** state id, unit of observation

**year** time in panel data

**l\_homicide** log of the number of homicides per capita

**effyear** year that castle doctrine is passed

**post** 0/1 variable for when castle doctrine is active

**time\_til** time relative to castle doctrine being passed into law

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df_het	<i>Simulated data with two treatment groups and heterogenous effects Generated using the following call: did2s::gen_data(panel = c(1990, 2020), g1 = 2000, g2 = 2010, g3 = 0, te1 = 2, te2 = 1, te3 = 0, te_m1 = 0.05, te_m2 = 0.15, te_m3 = 0)</i>
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### Description

Simulated data with two treatment groups and heterogenous effects

Generated using the following call: did2s::gen\_data(panel = c(1990, 2020), g1 = 2000, g2 = 2010, g3 = 0, te1 = 2, te2 = 1, te3 = 0, te\_m1 = 0.05, te\_m2 = 0.15, te\_m3 = 0)

**Usage**

df\_het

**Format**

A data frame with 31000 rows and 15 variables:

**unit** individual in panel data

**year** time in panel data

**g** the year that treatment starts

**dep\_var** outcome variable

**treat** T/F variable for when treatment is on

**rel\_year** year relative to treatment start. Inf = never treated.

**rel\_year\_binned** year relative to treatment start, but  $\leq -6$  and  $\geq 6$  are binned.

**unit\_fe** Unit FE

**year\_fe** Year FE

**error** Random error component

**te** Static treatment effect = te

**te\_dynamic** Dynamic treatment effect = te\_m

**state** State that unit is in

**group** String name for group

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df_hom	<i>Simulated data with two treatment groups and homogenous effects Generated using the following call: did2s::gen_data(panel = c(1990, 2020), g1 = 2000, g2 = 2010, g3 = 0, te1 = 2, te2 = 2, te3 = 0, te_m1 = 0, te_m2 = 0, te_m3 = 0)</i>
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**Description**

Simulated data with two treatment groups and homogenous effects

Generated using the following call: did2s::gen\_data(panel = c(1990, 2020), g1 = 2000, g2 = 2010, g3 = 0, te1 = 2, te2 = 2, te3 = 0, te\_m1 = 0, te\_m2 = 0, te\_m3 = 0)

**Usage**

df\_hom

**Format**

A data frame with 31000 rows and 15 variables:

**unit** individual in panel data

**year** time in panel data

**g** the year that treatment starts

**dep\_var** outcome variable

**treat** T/F variable for when treatment is on

**rel\_year** year relative to treatment start. Inf = never treated.

**rel\_year\_binned** year relative to treatment start, but  $\leq -6$  and  $\geq 6$  are binned.

**unit\_fe** Unit FE

**year\_fe** Year FE

**error** Random error component

**te** Static treatment effect = te

**te\_dynamic** Dynamic treatment effect = te\_m

**group** String name for group

**state** State that unit is in

**weight** Weight from runif()

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did2s	<i>Calculate two-stage difference-in-differences following Gardner (2021)</i>
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**Description**

Calculate two-stage difference-in-differences following Gardner (2021)

**Usage**

```
did2s(
  data,
  yname,
  first_stage,
  second_stage,
  treatment,
  cluster_var,
  weights = NULL,
  bootstrap = FALSE,
  n_bootstraps = 250,
  return_bootstrap = FALSE,
  verbose = TRUE
)
```

## Arguments

<code>data</code>	The dataframe containing all the variables
<code>yname</code>	Outcome variable
<code>first_stage</code>	Fixed effects and other covariates you want to residualize with in first stage. Formula following <code>fixest::feols</code> . Fixed effects specified after " <code> </code> ".
<code>second_stage</code>	Second stage, these should be the treatment indicator(s) (e.g. treatment variable or event-study leads/lags). Formula following <code>fixest::feols</code> . Use <code>i()</code> for factor variables, see <code>fixest::i</code> .
<code>treatment</code>	A variable that = 1 if treated, = 0 otherwise
<code>cluster_var</code>	What variable to cluster standard errors. This can be IDs or a higher aggregate level (state for example)
<code>weights</code>	Optional. Variable name for regression weights.
<code>bootstrap</code>	Optional. Should standard errors be calculated using bootstrap? Default is FALSE.
<code>n_bootstraps</code>	Optional. How many bootstraps to run. Default is 250.
<code>return_bootstrap</code>	Optional. Logical. Will return each bootstrap second-stage estimate to allow for manual use, e.g. percentile standard errors and empirical confidence intervals.
<code>verbose</code>	Optional. Logical. Should information about the two-stage procedure be printed back to the user? Default is TRUE.

## Value

`fixest` object with adjusted standard errors (either by formula or by bootstrap). All the methods from `fixest` package will work, including `fixest::esttable` and `fixest::coefplot`

## Examples

Load example dataset which has two treatment groups and homogeneous treatment effects

```
# Load Example Dataset
data("df_hom")
```

### Static TWFE:

You can run a static TWFE fixed effect model for a simple treatment indicator

```
static <- did2s(df_hom,
  yname = "dep_var", treatment = "treat", cluster_var = "state",
  first_stage = ~ 0 | unit + year,
  second_stage = ~ i(treat, ref=FALSE))
#> Running Two-stage Difference-in-Differences
#> • first stage formula `~ 0 | unit + year`
#> • second stage formula `~ i(treat, ref = FALSE)`
#> • The indicator variable that denotes when treatment is on is `treat`
#> • Standard errors will be clustered by `state`
```

```

fixest::esttable(static)
#>                               static
#> Dependent Var.:                dep_var
#>
#> treat = TRUE      2.025*** (0.0307)
#> -----
#> S.E. type          Custom
#> Observations      31,000
#> R2                 0.31846
#> Adj. R2           0.31846

```

### Event Study:

Or you can use relative-treatment indicators to estimate an event study estimate

```

es <- did2s(df_hom,
  yname = "dep_var", treatment = "treat", cluster_var = "state",
  first_stage = ~ 0 | unit + year,
  second_stage = ~ i(rel_year, ref=c(-1, Inf)))
#> Running Two-stage Difference-in-Differences
#> • first stage formula `~ 0 | unit + year`
#> • second stage formula `~ i(rel_year, ref = c(-1, Inf))`
#> • The indicator variable that denotes when treatment is on is `treat`
#> • Standard errors will be clustered by `state`

```

```

fixest::esttable(es)
#>                               es
#> Dependent Var.:                dep_var
#>
#> rel_year = -20   -0.0335 (0.0697)
#> rel_year = -19    0.0581 (0.0588)
#> rel_year = -18    0.0348 (0.0578)
#> rel_year = -17    0.0236 (0.0670)
#> rel_year = -16    0.0115 (0.0542)
#> rel_year = -15   -0.0148 (0.0769)
#> rel_year = -14    0.1150. (0.0613)
#> rel_year = -13   -0.0108 (0.0720)
#> rel_year = -12   -0.0727 (0.0635)
#> rel_year = -11    0.0666 (0.0559)
#> rel_year = -10    0.0396 (0.0382)
#> rel_year = -9     -0.0109 (0.0379)
#> rel_year = -8     0.0105 (0.0388)
#> rel_year = -7     -0.0001 (0.0445)
#> rel_year = -6    -0.0829* (0.0388)
#> rel_year = -5     0.0189 (0.0429)
#> rel_year = -4    -0.0664 (0.0437)
#> rel_year = -3    -0.0144 (0.0302)
#> rel_year = -2     0.0223 (0.0442)
#> rel_year = 0     2.117*** (0.0622)
#> rel_year = 1     1.857*** (0.0720)

```

```

#> rel_year = 2    1.986*** (0.0665)
#> rel_year = 3    2.005*** (0.0708)
#> rel_year = 4    1.950*** (0.0748)
#> rel_year = 5    2.038*** (0.0797)
#> rel_year = 6    2.032*** (0.0732)
#> rel_year = 7    2.025*** (0.0858)
#> rel_year = 8    1.976*** (0.0621)
#> rel_year = 9    2.121*** (0.0619)
#> rel_year = 10   2.088*** (0.0913)
#> rel_year = 11   1.943*** (0.1148)
#> rel_year = 12   1.941*** (0.1131)
#> rel_year = 13   1.965*** (0.1076)
#> rel_year = 14   2.023*** (0.0972)
#> rel_year = 15   2.235*** (0.1174)
#> rel_year = 16   2.178*** (0.1282)
#> rel_year = 17   1.936*** (0.1044)
#> rel_year = 18   2.135*** (0.1115)
#> rel_year = 19   2.112*** (0.1038)
#> rel_year = 20   1.925*** (0.1071)
#> -----
#> S.E. type          Custom
#> Observations      31,000
#> R2                 0.31957
#> Adj. R2            0.31872

# plot rel_year coefficients and standard errors
fixest::coefplot(es, keep = "rel_year::(.*)")

```

### Example from Cheng and Hoekstra (2013):

Here's an example using data from Cheng and Hoekstra (2013)

```

# Castle Data
castle <- haven::read_dta("https://github.com/scunning1975/mixtape/raw/master/castle.dta")

did2s(
  data = castle,
  yname = "l_homicide",
  first_stage = ~ 0 | sid + year,
  second_stage = ~ i(post, ref=0),
  treatment = "post",
  cluster_var = "state", weights = "popwt"
)
#> Running Two-stage Difference-in-Differences
#> • first stage formula `~ 0 | sid + year`
#> • second stage formula `~ i(post, ref = 0)`
#> • The indicator variable that denotes when treatment is on is `post`
#> • Standard errors will be clustered by `state`
#> OLS estimation, Dep. Var.: l_homicide
#> Observations: 550
#> Standard-errors: Custom

```

```
#>           Estimate Std. Error t value Pr(>|t|)
#> post::1 0.075142    0.03538  2.1239 0.034127 *
#> ---
#> Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
#> RMSE: 263.4   Adj. R2: 0.052465
```

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event_study	<i>Estimate event-study coefficients using TWFE and 5 proposed improvements.</i>
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## Description

Uses the estimation procedures recommended from Borusyak, Jaravel, Spiess (2021); Callaway and Sant'Anna (2020); Gardner (2021); Roth and Sant'Anna (2021); Sun and Abraham (2020)

## Usage

```
event_study(
  data,
  yname,
  idname,
  gname,
  tname,
  xformula = NULL,
  horizon = NULL,
  weights = NULL
)

plot_event_study(out, separate = TRUE, horizon = NULL)
```

## Arguments

data	The dataframe containing all the variables
yname	Variable name for outcome variable
idname	Variable name for unique unit id
gname	Variable name for unit-specific date of initial treatment (never-treated should be zero or NA)
tname	Variable name for calendar period
xformula	A formula for the covariates to include in the model. It should be of the form $\sim X1 + X2$ . Default is NULL.
horizon	Numeric. Vector of length 2. First element is min and second element is max of event_time to plot
weights	Variable name for estimation weights. This is used in estimating $Y(0)$ and also augments treatment effect weights
out	Output from <a href="#">event_study()</a>
separate	Logical. Should the estimators be on separate plots? Default is TRUE.



**Value**

event\_study returns a data.frame of point estimates for each estimator  
 plot\_event\_study returns a ggplot object that can be fully customized

**Examples**

```
out = event_study(
  data = did2s::df_het, yname = "dep_var", idname = "unit",
  tname = "year", gname = "g"
)
plot_event_study(out)
```

---

 gen\_data

*Generate TWFE data*


---

**Description**

Generate TWFE data

**Usage**

```
gen_data(
  g1 = 2000,
  g2 = 2010,
  g3 = 0,
  panel = c(1990, 2020),
  te1 = 2,
  te2 = 2,
  te3 = 2,
  te_m1 = 0,
  te_m2 = 0,
  te_m3 = 0
)
```

**Arguments**

g1	treatment date for group 1. For no treatment, set g = 0.
g2	treatment date for group 2. For no treatment, set g = 0.
g3	treatment date for group 3. For no treatment, set g = 0.
panel	numeric vector of size 2, start and end years for panel
te1	treatment effect for group 1. Will ignore for that group if g = 0.
te2	treatment effect for group 1. Will ignore for that group if g = 0.
te3	treatment effect for group 1. Will ignore for that group if g = 0.

te_m1	treatment effect slope per year
te_m2	treatment effect slope per year
te_m3	treatment effect slope per year

**Value**

Dataframe of generated data

**Examples**

```
# Homogeneous treatment effect
df_hom <- gen_data(panel = c(1990, 2020),
  g1 = 2000, g2 = 2010, g3 = 0,
  te1 = 2, te2 = 2, te3 = 0,
  te_m1 = 0, te_m2 = 0, te_m3 = 0)
# Heterogeneous treatment effect
df_het <- gen_data(panel = c(1990, 2020),
  g1 = 2000, g2 = 2010, g3 = 0,
  te1 = 2, te2 = 1, te3 = 0,
  te_m1 = 0.05, te_m2 = 0.15, te_m3 = 0)
```

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