

# Package ‘fdid’

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**Type** Package

**Title** Factorial Difference-in-Differences

**Version** 1.0.2

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**Description** Implements the factorial difference-in-differences (FDID) framework for panel data settings where all units are exposed to a universal event but vary in a baseline factor  $G$ . Provides support for various estimators; supports robust, bootstrap, and jackknife variance; returns dynamic, pre/event/post aggregates and raw means; and includes helpers for data preparation and plotting. Methodology follows Xu, Zhao and Ding (2026) <[doi:10.1080/01621459.2026.2628343](https://doi.org/10.1080/01621459.2026.2628343)>.

**URL** <https://yiqingxu.org/packages/fdid/>,  
<https://github.com/xuyiqing/fdid>

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**Encoding** UTF-8

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foreach, doFuture, future, ebal, grf, car, sandwich

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**VignetteBuilder** knitr

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**NeedsCompilation** no

**Author** Yiqing Xu [aut, cre],  
Rivka Lipkowitz [aut],  
Enhan Liu [aut]

**Maintainer** Yiqing Xu <yiqingxu@stanford.edu>

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fdid	<i>Factorial Difference-in-Differences Estimation</i>
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### Description

Performs factorial difference-in-differences (FDID) estimation using various methods and variance estimation techniques.

### Usage

```
fdid(
  s,
  tr_period,
  ref_period,
  entire_period = NULL,
  method = "ols1",
  vartype = "robust",
  missing_data = c("listwise", "available"),
  nsims = 1000,
  parallel = FALSE,
  cores = 2,
  target.pop = c("all", "1", "0")
)
```

### Arguments

s	A data frame prepared using <code>fdid_prepare</code> .
tr_period	A numeric vector specifying the treatment periods.
ref_period	A numeric scalar specifying the reference period.
entire_period	A numeric vector specifying the total range of time periods. If <code>NULL</code> , estimation is performed on all available time periods. Example: <code>c(1958, 1959, 1960, 1961)</code> .
method	A string specifying the estimation method. Options: <code>"ols1"</code> , <code>"ols2"</code> , <code>"did"</code> , <code>"ebal"</code> , <code>"ipw"</code> , <code>"aipw"</code> . Default is <code>"ols1"</code> .

vartype	A string specifying the variance estimation type. Options: "robust", "bootstrap", "jackknife". Default is "robust".
missing_data	How to handle missing data. Two options: <ul style="list-style-type: none"> <li>"listwise": Drop any row missing <b>any</b> relevant column (including outcomes in the periods used).</li> <li>"available": Drop rows only if they are missing in group/covariates/cluster columns, but allow partial usage of outcomes.</li> </ul> Default is "listwise".
nsims	Number of simulations for bootstrap variance estimation. Default is 1000.
parallel	Logical; whether to perform parallel computations. Default is FALSE.
cores	Number of cores for parallel computations. Default is 2.
target.pop	Character; the target population for averaging: "all", "1", or "0". "all" corresponds to the full sample. "1" targets the G=1 population. "0" targets the G=0 population. Default is "all".

### Value

A list with the following components:

est	A list with three elements: \$pre, \$event, and \$post containing aggregated pre-treatment, overall event, and post-treatment FDID estimates, respectively.
dynamic	Dynamic FDID estimates for each time in entire_period.
raw_means	Raw mean outcomes by group for each time in entire_period.
tr_period	Treatment periods used.
ref_period	Reference period used.
entire_period	All time periods for dynamic estimation.
method	Method used.
vartype	Variance type used.
times	All numeric time columns found.
G	Group indicator (0/1).
ps	Propensity scores (if ipw or aipw method used).
call	The matched call.
target.pop	Character indicating the target population used.

### Author(s)

Rivka Lipkowitz, Enhao Liu

## Examples

```
data(fdid)
mortality$uniqueid <- paste(mortality$provid, mortality$countyid, sep = "-")
mortality$G <- ifelse(mortality$pczupu >= median(mortality$pczupu, na.rm = TRUE), 1, 0)
s <- fdid_prepare(
  data = mortality, Y_label = "mortality",
  X_labels = c("avggrain", "lnpop"),
  G_label = "G", unit_label = "uniqueid", time_label = "year"
)
result <- fdid(s, tr_period = 1958:1961, ref_period = 1957)
summary(result)
```

---

fdid\_list

*Create an 'fdid\_list' Object*

---

## Description

Bundles multiple 'fdid' objects into a single list with class "'fdid\_list'" for convenient collective handling.

## Usage

```
fdid_list(..., validate = TRUE)
```

## Arguments

... One or more objects of class "'fdid'", or a single list of them.

validate Logical; if 'TRUE' (default) verify each element inherits from "'fdid'".

## Value

A list with classes 'c("fdid\_list", "list")'.

## Author(s)

Rivka Lipkowitz

**Description**

Prepares a dataset for factorial difference-in-differences (FDID) analysis by reshaping the data into a wide format, averaging time-varying covariates, and renaming columns for consistency in subsequent analysis.

**Usage**

```
fdid_prepare(  
  data,  
  Y_label,  
  X_labels = NULL,  
  G_label,  
  unit_label,  
  time_label,  
  cluster_label = NULL  
)
```

**Arguments**

<code>data</code>	A data frame containing the dataset to be processed.
<code>Y_label</code>	A string specifying the column name of the outcome variable.
<code>X_labels</code>	A character vector specifying the column names of the time-varying covariates.
<code>G_label</code>	A string specifying the column name of the group variable (e.g., treatment vs. control).
<code>unit_label</code>	A string specifying the column name of the unit identifier (e.g., individual or entity).
<code>time_label</code>	A string specifying the column name of the time variable.
<code>cluster_label</code>	An optional string specifying the column name of the clustering variable. Default is 'NULL'.

**Value**

A data frame in wide format with the following: - Outcome variable pivoted to wide format with time columns. - Time-varying covariates averaged across time. - Columns renamed: - Unit identifier -> 'unit' - Covariates -> 'x1', 'x2', ... - Group variable -> 'G' - Clustering variable (if provided) -> 'c'

**Author(s)**

Rivka Lipkowitz

**Examples**

```

data <- data.frame(
  id = rep(1:3, each = 4),
  time = rep(1:4, times = 3),
  outcome = rnorm(12),
  covar1 = runif(12),
  covar2 = runif(12),
  group = c(0, 0, 1, 1, 0, 0, 1, 1, 0, 0, 1, 1)
)
fdid_data <- fdid_prepare(
  data = data,
  Y_label = "outcome",
  X_labels = c("covar1", "covar2"),
  G_label = "group",
  unit_label = "id",
  time_label = "time"
)
head(fdid_data)

```

---

mortality

*FDID example dataset*


---

**Description**

A long-format panel dataset for demonstrating the fdid package.

**Usage**

```
mortality
```

**Format**

A data frame with 11973 rows and 17 columns:

**provid** Province ID

**countyid** County ID

**zupu** Genealogy book count

**pczupu** Genealogy book density (per capita); 45% of counties have zero

**lnpczupu** Log-transformed genealogy density:  $\log(\text{pczupu} + 1)$ ; used as a continuous treatment in Xu, Zhao, and Ding (2026)

**anyzupu** Indicator: any genealogy book present

**avggrain** Average grain output

**nograin** Indicator: no grain data

**urban** Urban population share

**dis\_bj** Distance to Beijing

**dis\_pc** Distance to provincial capital  
**rice** Rice cultivation indicator  
**minority** Minority population share  
**edu** Education level  
**lnpop** Log population  
**year** Year (1954–1966)  
**mortality** Mortality rate

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plot.fdid

*Plot Results from FDID Analysis*


---

### Description

Provides visualisations for FDID results, including raw means, dynamic effects, and propensity-score overlap. The comparison plot of multiple methods has been removed; use `plot.fdid_list()` for that.

### Usage

```
## S3 method for class 'fdid'
plot(
  x,
  type = c("raw", "dynamic", "overlap"),
  connected = FALSE,
  ci = TRUE,
  shade_periods = x$str_period,
  alpha_shade = 0.2,
  palette = "Set2",
  group_labels = c("Group 0", "Group 1"),
  xlab = NULL,
  ylab = NULL,
  main = NULL,
  ylim = NULL,
  ...
)
```

### Arguments

<code>x</code>	An <code>fdid</code> object.
<code>type</code>	One of "raw", "dynamic", or "overlap".
<code>connected</code>	Logical; if TRUE, connects points with lines in the "raw" and "dynamic" plots. Default is FALSE.
<code>ci</code>	Logical; if TRUE, draw 95% CIs when available. Default is TRUE.

shade\_periods Shaded intervals on the time axis. Default uses x\$str\_period, i.e. event periods. Set to NULL to remove shaded area.

alpha\_shade Transparency for shading the treatment period.

palette A palette name from **RColorBrewer**. Default "Set2".

group\_labels Labels for the two groups.

xlab, ylab, main Axis labels and main title.

ylim Y-axis limits. Default NULL (computed automatically).

... Additional graphics parameters.

**Value**

Produces a plot; invisibly returns NULL.

**Author(s)**

Rivka Lipkowitz, Enhan Liu

**Examples**

```
data(fdid)
mortality$uniqueid <- paste(mortality$provid, mortality$countyid, sep = "-")
mortality$G <- ifelse(mortality$pczupu >= median(mortality$pczupu, na.rm = TRUE), 1, 0)
s <- fdid_prepare(
  data = mortality, Y_label = "mortality",
  X_labels = c("avggrain", "lnpop"),
  G_label = "G", unit_label = "uniqueid", time_label = "year"
)
result <- fdid(s, tr_period = 1958:1961, ref_period = 1957)
plot(result, type = "raw")
plot(result, type = "dynamic")
```

---

plot.fdid\_list

*Plot Multiple FDID Estimates*

---

**Description**

Creates a comparison plot of point estimates and confidence intervals for every element of an 'fdid\_list'.

**Usage**

```
## S3 method for class 'fdid_list'
plot(
  x,
  xlab = NULL,
  ylab = NULL,
```

```

    main = NULL,
    ylim = NULL,
    vertical = TRUE,
    show_vartype = TRUE,
    ...
)

```

### Arguments

x	An object of class <code>"fdid_list"</code> .
xlab, ylab, main	Axis labels and title. If <code>'NULL'</code> , sensible defaults are used.
ylim	Optional numeric vector of length two giving the <i>estimate-axis</i> limits. (Backward compatible: for horizontal plots this is the x-limit; for vertical plots this is the y-limit.)
vertical	Logical; default is TRUE.
show_vartype	Logical; include vartype in labels. Default is TRUE.
...	Additional graphics parameters passed to <code>plot()</code> .

### Value

Invisibly returns `'x'`; called for its side-effect of drawing a plot.

### Author(s)

Rivka Lipkowitz, Enhao Liu

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print.fdid

*Print Method for FDID Objects*

---

### Description

Print Method for FDID Objects

### Usage

```

## S3 method for class 'fdid'
print(x, ...)

```

### Arguments

x	An object of class <code>'fdid'</code> .
...	Additional arguments (not used).

### Value

Prints a brief overview of the `'fdid'` object

**Author(s)**

Rivka Lipkovitz.

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summary.fdid

*Summary Method for FDID Objects*

---

**Description**

Summary Method for FDID Objects

**Usage**

```
## S3 method for class 'fdid'  
summary(object, ...)
```

**Arguments**

object            An object of class fdid.  
...                Additional arguments (not used).

**Value**

Prints a summary of the fdid object.

**Author(s)**

Rivka Lipkovitz, Enhao Liu

**Examples**

```
data(fdid)  
mortality$uniqueid <- paste(mortality$provid, mortality$countyid, sep = "-")  
mortality$G <- ifelse(mortality$pczupu >= median(mortality$pczupu, na.rm = TRUE), 1, 0)  
s <- fdid_prepare(  
  data = mortality, Y_label = "mortality",  
  X_labels = c("avggrain", "lnpop"),  
  G_label = "G", unit_label = "uniqueid", time_label = "year"  
)  
result <- fdid(s, tr_period = 1958:1961, ref_period = 1957)  
summary(result)
```

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