

# Package ‘Zelig’

April 22, 2019

**License** GPL (>= 3)

**Title** Everyone's Statistical Software

**Description** A framework that brings together an abundance of common statistical models found across packages into a unified interface, and provides a common architecture for estimation and interpretation, as well as bridging functions to absorb increasingly more models into the package. Zelig allows each individual package, for each statistical model, to be accessed by a common uniformly structured call and set of arguments. Moreover, Zelig automates all the surrounding building blocks of a statistical work-flow--procedures and algorithms that may be essential to one user's application but which the original package developer did not use in their own research and might not themselves support. These include bootstrapping, jackknifing, and re-weighting of data. In particular, Zelig automatically generates predicted and simulated quantities of interest (such as relative risk ratios, average treatment effects, first differences and predicted and expected values) to interpret and visualize complex models.

**URL** <https://cran.r-project.org/package=Zelig>

**BugReports** <https://github.com/IQSS/Zelig/issues>

**Version** 5.1.6.1

**Date** 2018-02-27

**Depends** survival

**Imports** AER, Amelia, coda, dplyr (>= 0.3.0.2), Formula, geopack, jsonlite, sandwich, MASS, MatchIt, maxLik, MCMCpack, methods, quantreg, survey, VGAM

**Suggests** ei, eiPack, knitr, networkD3, optmatch, rmarkdown, testthat, tidyverse, ZeligChoice, ZeligEI, zeligverse

**Collate** 'assertions.R' 'model-zelig.R' 'model-timeseries.R'  
'model-ma.R' 'model-ar.R' 'model-arma.R' 'model-weibull.R'  
'model-tobit.R' 'model-bayes.R' 'model-tobit-bayes.R'  
'model-glm.R' 'model-binchoice.R' 'model-probit.R'  
'model-probit-bayes.R' 'model-poisson.R'

'model-poisson-bayes.R' 'model-oprobit-bayes.R'  
 'model-normal.R' 'model-normal-bayes.R' 'model-mlogit-bayes.R'  
 'model-gamma.R' 'model-gee.R' 'model-logit.R'  
 'model-logit-bayes.R' 'model-factor-bayes.R'  
 'model-poisson-gee.R' 'model-normal-gee.R' 'model-gamma-gee.R'  
 'model-binchoice-gee.R' 'model-probit-gee.R'  
 'model-logit-gee.R' 'model-relogit.R' 'model-quantile.R'  
 'model-lognorm.R' 'model-exp.R' 'model-negbinom.R'  
 'model-ivreg.R' 'model-ls.R' 'utils.R' 'create-json.R'  
 'datasets.R' 'interface.R' 'model-survey.R'  
 'model-binchoice-survey.R' 'model-gamma-survey.R'  
 'model-logit-survey.R' 'model-normal-survey.R'  
 'model-poisson-survey.R' 'model-probit-survey.R' 'plots.R'  
 'wrappers.R'

**RoxygenNote** 6.0.1

**NeedsCompilation** no

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approval *U.S. Presidential Approval Data*

**Description**

Monthly public opinion data for 2001-2006.

**Usage**

data(approval)

**Format**

A table containing 8 variables ("month", "year", "approve", "disapprove", "unsure", "sept.oct.2001", "iraq.war", and "avg.price") and 65 observations.

**Source**

ICPSR

**References**

Stuff here

---

ATT	<i>Compute simulated (sample) average treatment effects on the treated from a Zelig model estimation</i>
-----	--

---

**Description**

Compute simulated (sample) average treatment effects on the treated from a Zelig model estimation

**Usage**

```
ATT(object, treatment, treated = 1, num = NULL)
```

**Arguments**

object	an object of class Zelig
treatment	character string naming the variable that denotes the treatment and non-treated groups.
treated	value of treatment variable indicating treatment
num	number of simulations to run. Default is 1000.

**Author(s)**

Christopher Gandrud

**Examples**

```
library(dplyr)
data(sanction)
z.att <- zelig(num ~ target + coop + mil, model = "poisson",
              data = sanction) %>%
  ATT(treatment = "mil") %>%
  get_qi(qi = "ATT", xvalue = "TE")
```

---

avg	<i>Compute central tendency as appropriate to data type</i>
-----	---

---

**Description**

Compute central tendency as appropriate to data type

**Usage**

```
avg(val)
```

**Arguments**

`val` a vector of values

**Value**

a mean (if numeric) or a median (if ordered) or mode (otherwise)

---

bivariate	<i>Sample data for bivariate probit regression</i>
-----------	--

---

**Description**

Sample data for the bivariate probit regression.

**Usage**

```
data(bivariate)
```

**Format**

A table containing 6 variables ("y1", "y2", "x1", "x2", "x3", and "x4") and 78 observations.

**Source**

This is a cleaned and relabelled version of the sanction data set, available in Zelig.

**References**

Martin, Lisa (1992). *Coercive Cooperation: Explaining Multilateral Economic Sanctions*, Princeton: Princeton University Press.

---

<code>ci.plot</code>	<i>Method for plotting qi simulations across a range within a variable, with confidence intervals</i>
----------------------	---

---

**Description**

Method for plotting qi simulations across a range within a variable, with confidence intervals

**Usage**

```
ci.plot(obj, qi="ev", var=NULL, ..., main = NULL, sub =
  NULL, xlab = NULL, ylab = NULL, xlim = NULL, ylim =
  NULL, legcol="gray20", col=NULL, leg=1, legpos=
  NULL, ci = c(80, 95, 99.9), discount=NULL)
```

**Arguments**

obj	A reference class zelig5 object
qi	a character-string specifying the quantity of interest to plot
var	The variable to be used on the x-axis. Default is the variable across all the chosen values with smallest nonzero variance
...	Parameters to be passed to the ‘truehist’ function which is implicitly called for numeric simulations
main	a character-string specifying the main heading of the plot
sub	a character-string specifying the sub heading of the plot
xlab	a character-string specifying the label for the x-axis
ylab	a character-string specifying the label for the y-axis
xlim	Limits to the x-axis
ylim	Limits to the y-axis
legcol	“legend color”, an valid color used for plotting the line colors in the legend
col	a valid vector of colors of at least length 3 to use to color the confidence intervals
leg	“legend position”, an integer from 1 to 4, specifying the position of the legend. 1 to 4 correspond to “SE”, “SW”, “NW”, and “NE” respectively. Setting to 0 or “n” turns off the legend.
legpos	“legend type”, exact coordinates and sizes for legend. Overrides argment “leg.type”
ci	vector of length three of confidence interval levels to draw.
discont	optional point of discontinuity along the x-axis at which to interrupt the graph

**Value**

the current graphical parameters. This is subject to change in future implementations of Zelig

**Author(s)**

James Honaker

---

CigarettesSW

*Cigarette Consumption Panel Data*

---

**Description**

Cigarette Consumption Panel Data

**Format**

A data set with 96 observations and 9 variables

**Source**

From Christian Kleiber and Achim Zeileis (2008). Applied Econometrics with R. New York: Springer-Verlag. ISBN 978-0-387-77316-2. URL <https://CRAN.R-project.org/package=AER>



---

cluster.formula	<i>Generate Formulae that Consider Clustering</i>
-----------------	---

---

**Description**

This method is used internally by the "Zelig" Package to interpret clustering in GEE models.

**Usage**

```
cluster.formula(formula, cluster)
```

**Arguments**

formula	a formula object
cluster	a vector

**Value**

a formula object describing clustering

---

coalition	<i>Coalition Dissolution in Parliamentary Democracies</i>
-----------	---

---

**Description**

This data set contains survival data on government coalitions in parliamentary democracies (Belgium, Canada, Denmark, Finland, France, Iceland, Ireland, Israel, Italy, Netherlands, Norway, Portugal, Spain, Sweden, and the United Kingdom) for the period 1945-1987. For parsimony, country indicator variables are omitted in the sample data.

**Usage**

```
data(coalition)
```

**Format**

A table containing 7 variables ("duration", "ciep12", "invest", "fract", "polar", "numst2", "crisis") and 314 observations. For variable descriptions, please refer to King, Alt, Burns and Laver (1990).

**Source**

ICPSR

**References**

King, Gary, James E. Alt, Nancy Elizabeth Burns and Michael Laver (1990). "A Unified Model of Cabinet Dissolution in Parliamentary Democracies," *American Journal of Political Science*, vol. 34, no. 3, pp. 846-870.

Gary King, James E. Alt, Nancy Burns, and Michael Laver. ICPSR Publication Related Archive, 1115.

---

coalition2

*Coalition Dissolution in Parliamentary Democracies, Modified Version*

---

**Description**

This data set contains survival data on government coalitions in parliamentary democracies (Belgium, Canada, Denmark, Finland, France, Iceland, Ireland, Israel, Italy, Netherlands, Norway, Portugal, Spain, Sweden, and the United Kingdom) for the period 1945-1987. Country indicator variables are included in the sample data.

**Usage**

```
data(coalition2)
```

**Format**

A data frame containing 8 variables ("duration", "ciep12", "invest", "fract", "polar", "numst2", "crisis", "country") and 314 observations. For variable descriptions, please refer to King, Alt, Burns and Laver (1990).

**Source**

ICPSR

**References**

King, Gary, James E. Alt, Nancy Elizabeth Burns and Michael Laver (1990). "A Unified Model of Cabinet Dissolution in Parliamentary Democracies," *American Journal of Political Science*, vol. 34, no. 3, pp. 846-870.

Gary King, James E. Alt, Nancy Burns, and Michael Laver. ICPSR Publication Related Archive, 1115.

---

coef, Zelig-method      *Method for extracting estimated coefficients from Zelig objects*

---

### **Description**

Method for extracting estimated coefficients from Zelig objects

### **Usage**

```
## S4 method for signature 'Zelig'  
coef(object)
```

### **Arguments**

object              An Object of Class Zelig

---

coefficients,Zelig-method  
*Method for extracting estimated coefficients from Zelig objects*

---

### **Description**

Method for extracting estimated coefficients from Zelig objects

### **Usage**

```
## S4 method for signature 'Zelig'  
coefficients(object)
```

### **Arguments**

object              An Object of Class Zelig

---

combine_coef_se	<i>Combines estimated coefficients and associated statistics from models estimated with multiply imputed data sets or bootstrapped</i>
-----------------	--

---

### Description

Combines estimated coefficients and associated statistics from models estimated with multiply imputed data sets or bootstrapped

### Usage

```
combine_coef_se(obj, out_type = "matrix", bagging = FALSE,  
  messages = TRUE)
```

### Arguments

obj	a zelig object with an estimated model
out_type	either "matrix" or "list" specifying whether the results should be returned as a matrix or a list.
bagging	logical whether or not to bag the bootstrapped coefficients
messages	logical whether or not to return messages for what is being returned

### Value

If the model uses multiply imputed or bootstrapped data then a matrix (default) or list of combined coefficients (coef), standard errors (se), z values (zvalue), p-values (p) is returned. Rubin's Rules are used to combine output from multiply imputed data. An error is returned if no imputations were included or there wasn't bootstrapping. Please use `get_coef`, `get_se`, and `get_pvalue` methods instead in cases where there are no imputations or bootstrap.

### Author(s)

Christopher Gandrud and James Honaker

### Source

Partially based on [mi.meld](#) from Amelia.

### Examples

```
set.seed(123)  
  
## Multiple imputation example  
# Create fake imputed data  
n <- 100  
x1 <- runif(n)  
x2 <- runif(n)
```

```
y <- rnorm(n)
data.1 <- data.frame(y = y, x = x1)
data.2 <- data.frame(y = y, x = x2)

# Estimate model
mi.out <- to_zelig_mi(data.1, data.2)
z.out.mi <- zelig(y ~ x, model = "ls", data = mi.out)

# Combine and extract coefficients and standard errors
combine_coef_se(z.out.mi)

## Bootstrap example
z.out.boot <- zelig(y ~ x, model = "ls", data = data.1, bootstrap = 20)
combine_coef_se(z.out.boot)
```

---

createJSON

*Utility function for constructing JSON file that encodes the hierarchy of available statistical models in Zelig*

---

## Description

Utility function for construction a JSON file that encodes the hierarchy of available statistical models.

## Usage

```
createJSON(movefile=TRUE)
```

## Arguments

`movefile` Logical of whether to (TRUE) move the JSON file into path `./inst/JSON` or (FALSE) leave in working directory.

## Value

Returns TRUE on successful completion of json file

## Author(s)

Christine Choirat, Vito D'Orazio, James Honaker

df.residual,Zelig-method

*Method for extracting residual degrees-of-freedom from Zelig objects*

---

### **Description**

Method for extracting residual degrees-of-freedom from Zelig objects

### **Usage**

```
## S4 method for signature 'Zelig'  
df.residual(object)
```

### **Arguments**

object            An Object of Class Zelig

---

eidat

*Simulation Data for Ecological Inference*

---

### **Description**

This dataframe contains a simulated data set to illustrate the models for ecological inference.

### **Usage**

```
data(eidat)
```

### **Format**

A table containing 4 variables ("t0", "t1", "x0", "x1") and 10 observations.

---

fitted,Zelig-method     *Method for extracting estimated fitted values from Zelig objects*

---

**Description**

Method for extracting estimated fitted values from Zelig objects

**Usage**

```
## S4 method for signature 'Zelig'
fitted(object, ...)
```

**Arguments**

object	An Object of Class Zelig
...	Additional parameters to be passed to fitted

---

free1                     *Freedom of Speech Data*

---

**Description**

Selection of individual-level survey data for freedom of speech.

**Usage**

```
data(free1)
```

**Details**

A table with 150 observations and 12 variables.

- sex1 for men and 0 for women
- ageAge of respondent in years
- educLevels of education, coded as a numeric variable with
  - 1No formal education
  - 2Less than primary school education
  - 3Completed primary school
  - 4Completed secondary school
  - 5Completed high school
  - 6Completed college
  - 7Completed post-graduate degree
- countryCharacter strings consisting of "Oceania", "Eurasia", and "Eastasia", after Orwell's 1984.

- ySelf assessment (see below).
- v1-v6Response to vignettes (see below).

Survey respondents were asked in almost the same language for a self-assessment and for an assessment of several hypothetical persons described by written vignettes. The self assessment (self, in the data set), "How free do you think [name/you] [is/are] to express [him-her/your]self without fear of government reprisal?" was first asked of the survey respondent with respect to him or herself, and then after each of vignette. The possible response categories are:

- 1Completely free
- 2Very free
- 3Moderately free
- 4Slightly free
- 5Not free at all

The vignettes, ordered from most free to least free, are:

- vign1[Kay] does not like many of the government's policies. She frequently publishes her opinion in newspapers, criticizing decisions by officials and calling for change. She sees little reason these actions could lead to government reprisal.
- vign2[Michael] disagrees with many of the government's policies. Though he knows criticism is frowned upon, he doesn't believe the government would punish someone for expressing critical views. He makes his opinion known on most issues without regard to who is listening.
- vign3[Bob] has political views at odds with the government. He has heard of people occasionally being arrested for speaking out against the government, and government leaders sometimes make political speeches condemning those who criticize. He sometimes writes letters to newspapers about politics, but he is careful not to use his real name.
- vign4[Connie] does not like the government's stance on many issues. She has a friend who was arrested for being too openly critical of governmental leaders, and so she avoids voicing her opinions in public places.
- vign5[Vito] disagrees with many of the government's policies, and is very careful about whom he says this to, reserving his real opinions for family and close friends only. He knows several men who have been taken away by government officials for saying negative things in public.
- vign6[Sonny] lives in fear of being harassed for his political views. Everyone he knows who has spoken out against the government has been arrested or taken away. He never says a word about anything the government does, not even when he is at home alone with his family.

## References

*WHO's World Health Survey* by Lydia Bendib, Somnath Chatterji, Alena Petrakova, Ritu Sadana, Joshua A. Salomon, Margie Schneider, Bedirhan Ustun, Maria Villanueva

Jonathan Wand, Gary King and Olivia Lau. (2007) "Anchors: Software for Anchoring Vignettes". *Journal of Statistical Software*. Forthcoming. copy at <http://wand.stanford.edu/research/anchors-jss.pdf>

Gary King and Jonathan Wand. "Comparing Incomparable Survey Responses: New Tools for Anchoring Vignettes," *Political Analysis*, 15, 1 (Winter, 2007): Pp. 46-66, copy at <http://gking.harvard.edu/files/abs/c-abs.shtml>.



---

free2

*Freedom of Speech Data*

---

### Description

Selection of individual-level survey data for freedom of speech.

### Usage

```
data(free2)
```

### Details

A table with 150 observations and 12 variables.

- sex1 for men and 0 for women
- ageAge of respondent in years
- educLevels of education, coded as a numeric variable with
  - 1No formal education
  - 2Less than primary school education
  - 3Completed primary school
  - 4Completed secondary school
  - 5Completed high school
  - 6Completed college
  - 7Completed post-graduate degree
- countryCharacter strings consisting of "Oceania", "Eurasia", and "Eastasia", after Orwell's *1984*.
- ySelf assessment (see below).
- v1-v6Response to vignettes (see below).

Survey respondents were asked in almost the same language for a self-assessment and for an assessment of several hypothetical persons described by written vignettes. The self assessment (`self`, in the data set), "How free do you think [name/you] [is/are] to express [him-her/your]self without fear of government reprisal?" was first asked of the survey respondent with respect to him or herself, and then after each of vignette. The possible response categories are:

- 1Completely free
- 2Very free
- 3Moderately free
- 4Slightly free
- 5Not free at all

The vignettes, ordered from most free to least free, are:

- vign1[Kay] does not like many of the government's policies. She frequently publishes her opinion in newspapers, criticizing decisions by officials and calling for change. She sees little reason these actions could lead to government reprisal.
- vign2[Michael] disagrees with many of the government's policies. Though he knows criticism is frowned upon, he doesn't believe the government would punish someone for expressing critical views. He makes his opinion known on most issues without regard to who is listening.
- vign3[Bob] has political views at odds with the government. He has heard of people occasionally being arrested for speaking out against the government, and government leaders sometimes make political speeches condemning those who criticize. He sometimes writes letters to newspapers about politics, but he is careful not to use his real name.
- vign4[Connie] does not like the government's stance on many issues. She has a friend who was arrested for being too openly critical of governmental leaders, and so she avoids voicing her opinions in public places.
- vign5[Vito] disagrees with many of the government's policies, and is very careful about whom he says this to, reserving his real opinions for family and close friends only. He knows several men who have been taken away by government officials for saying negative things in public.
- vign6[Sonny] lives in fear of being harassed for his political views. Everyone he knows who has spoken out against the government has been arrested or taken away. He never says a word about anything the government does, not even when he is at home alone with his family.

## References

*WHO's World Health Survey* by Lydia Bendib, Somnath Chatterji, Alena Petrakova, Ritu Sadana, Joshua A. Salomon, Margie Schneider, Bedirhan Ustun, Maria Villanueva

Jonathan Wand, Gary King and Olivia Lau. (2007) "Anchors: Software for Anchoring Vignettes". *Journal of Statistical Software*. Forthcoming. copy at <http://wand.stanford.edu/research/anchors-jss.pdf>

Gary King and Jonathan Wand. "Comparing Incomparable Survey Responses: New Tools for Anchoring Vignettes," *Political Analysis*, 15, 1 (Winter, 2007): Pp. 46-66, copy at <http://gking.harvard.edu/files/abs/c-abs.shtml>.

---

friendship

*Simulated Example of Schoolchildren Friendship Network*

---

## Description

This data set contains six sociomatrices of simulated data on friendship ties among schoolchildren.

## Usage

`data(friendship)`

**Format**

Each variable in the dataset is a 15 by 15 matrix representing some form of social network tie held by the fictitious children. The matrices are labeled "friends", "advice", "prestige", "authority", "perpower" and "per".

The sociomatrices were combined into the friendship dataset using the `format.network.data` function from the `netglm` package by Skyler Cranmer as shown in the example.

**Source**

fictitious

**Examples**

```
## Not run:  
friendship <- format.network.data(friends, advice, prestige, authority, perpower, per)  
  
## End(Not run)
```

---

<code>from_zelig_model</code>	<i>Extract the original fitted model object from a zelig estimation</i>
-------------------------------	---

---

**Description**

Extract the original fitted model object from a zelig estimation

**Usage**

```
from_zelig_model(obj)
```

**Arguments**

`obj` a zelig object with an estimated model

**Details**

Extracts the original fitted model object from a zelig estimation. This can be useful for passing output to non-Zelig post-estimation functions and packages such as `texreg` and `stargazer` for creating well-formatted presentation document tables.

**Author(s)**

Christopher Gandrud

**Examples**

```
z5 <- zls$new()  
z5$zelig(Fertility ~ Education, data = swiss)  
from_zelig_model(z5)
```

---

get_pvalue	<i>Extract p-values from a Zelig estimated model</i>
------------	--

---

**Description**

Extract p-values from a Zelig estimated model

**Usage**

```
get_pvalue(object)
```

**Arguments**

object            an object of class Zelig

**Author(s)**

Christopher Gandrud

---

get_qi	<i>Extract quantities of interest from a Zelig simulation</i>
--------	---

---

**Description**

Extract quantities of interest from a Zelig simulation

**Usage**

```
get_qi(object, qi = "ev", xvalue = "x", subset = NULL)
```

**Arguments**

object            an object of class Zelig

qi                character string with the name of quantity of interest desired: "ev" for expected values, "pv" for predicted values or "fd" for first differences.

xvalue            character string stating which of the set of values of x should be used for getting the quantity of interest.

subset            subset for multiply imputed data (only relevant if multiply imputed data is supplied in the original call.)

**Author(s)**

Christopher Gandrud

---

`get_se`*Extract standard errors from a Zelig estimated model*

---

**Description**

Extract standard errors from a Zelig estimated model

**Usage**

```
get_se(object)
```

**Arguments**

`object` an object of class Zelig

**Author(s)**

Christopher Gandrud

---

`grunfeld`*Simulation Data for model Seemingly Unrelated Regression (sur) that corresponds to method SUR of systemfit*

---

**Description**

Dataframe contains 20 annual observations from 1935 to 1954 of 7 variables for two firms General Electric and Westinghouse. Columns are Year; Ige and Iw = Gross investment for GE and W, respectively; Fge and Fw=Market value of Firm as of begin of the year; Cge and Cw= Capital stock measure as of begin of the year.

**Usage**

```
data(grunfeld)
```

**Format**

A table containing 7 variables ("Year", "Ige", "Fge", "Cge", "Iw", "Fw", "Cw") and 20 observations.

---

hoff

*Social Security Expenditure Data*

---

### Description

This data set contains annual social security expenditure (as percent of budget lagged by two years), the relative frequency of mentions social justice received in the party's platform in each year, and whether the president is Republican or Democrat.

### Usage

data(hoff)

### Format

A table containing 5 variables ("year", "L2SocSec", "Just503D", "Just503R", "RGovDumy") and 36 observations.

### Source

ICPSR (replication dataset s1109)

### References

Gary King and Michael Laver. "On Party Platforms, Mandates, and Government Spending." *American Political Science Review*, Vol. 87, No. 3 (September, 1993): pp. 744-750.

---

homerun

*Sample Data on Home Runs Hit By Mark McGwire and Sammy Sosa in 1998.*

---

### Description

Game-by-game information for the 1998 season for Mark McGwire and Sammy Sosa. Data are a subset of the dataset provided in Simonoff (1998).

### Usage

data(homerun)

**Format**

A data frame containing 5 variables ("gameno", "month", "homeruns", "playerstatus", "player") and 326 observations.

gameno an integer variable denoting the game number

month a factor variable taking with levels "March" through "September" denoting the month of the game

homeruns an integer vector denoting the number of homeruns hit in that game for that player

playerstatus an integer vector equal to "0" if the player played in the game, and "1" if they did not.

player an integer vector equal to "0" (McGwire) or "1" (Sosa)

**Source**

<https://ww2.amstat.org/publications/jse/v6n3/datasets.simonoff.html>

**References**

Simonoff, Jeffrey S. 1998. "Move Over, Roger Maris: Breaking Baseball's Most Famous Record." *Journal of Statistics Education* 6(3). Data used are a subset of the data in the article.

---

immigration

*Individual Preferences Over Immigration Policy*

---

**Description**

These five datasets are part of a larger set of 10 multiply imputed data sets describing individual preferences toward immigration policy. Imputation was performed via Amelia.

**Format**

Each multiply-imputed data set consists of a table with 7 variables ("ipip", "wage1992", "prtyid", "ideol", "gender") and 2,485 observations. For variable descriptions, please refer to Scheve and Slaughter, 2001.

**Source**

National Election Survey

**References**

Scheve, Kenneth and Matthew Slaughter (2001). "Labor Market Competition and Individual Preferences Over Immigration Policy," *The Review of Economics and Statistics*, vol. 83, no. 1, pp. 133-145.

---

is_length_not_1	<i>Check if an object has a length greater than 1</i>
-----------------	---

---

**Description**

Check if an object has a length greater than 1

**Usage**

```
is_length_not_1(x, msg = "Length is 1.", fail = TRUE)
```

**Arguments**

x	an object
msg	character string with the error message to return if fail = TRUE.
fail	logical whether to return an error if length is not greater than 1.

---

is_simsrange	<i>Check if simulations for a range of fitted values are present in sim.out</i>
--------------	---

---

**Description**

Check if simulations for a range of fitted values are present in sim.out

**Usage**

```
is_simsrange(x, fail = TRUE)
```

**Arguments**

x	a sim.out method
fail	logical whether to return an error if simulation range is not present.



---

is_simsrange1	<i>Check if simulations for a range1 of fitted values are present in sim.out</i>
---------------	--

---

**Description**

Check if simulations for a range1 of fitted values are present in sim.out

**Usage**

```
is_simsrange1(x, fail = TRUE)
```

**Arguments**

x	a sim.out method
fail	logical whether to return an error if simulation range is not present.

---

is_simsx	<i>Check if simulations for individual values are present in sim.out</i>
----------	--

---

**Description**

Check if simulations for individual values are present in sim.out

**Usage**

```
is_simsx(x, fail = TRUE)
```

**Arguments**

x	a sim.out method
fail	logical whether to return an error if simulation range is not present.

---

is_simsx1	<i>Check if simulations for individual values for x1 are present in sim.out</i>
-----------	---

---

**Description**

Check if simulations for individual values for x1 are present in sim.out

**Usage**

```
is_simsx1(x, fail = TRUE)
```

**Arguments**

x	a sim.out method
fail	logical whether to return an error if simulation range is not present.

---

is_sims_present	<i>Check if any simulations are present in sim.out</i>
-----------------	--

---

**Description**

Check if any simulations are present in sim.out

**Usage**

```
is_sims_present(x, fail = TRUE)
```

**Arguments**

x	a sim.out method
fail	logical whether to return an error if no simulations are present.

---

is_timeseries	<i>Check if a zelig object contains a time series model</i>
---------------	---

---

**Description**

Check if a zelig object contains a time series model

**Usage**

```
is_timeseries(x, msg = "Not a timeseries object.", fail = FALSE)
```

**Arguments**

x	a zelig object
msg	character string with the error message to return if fail = TRUE.
fail	logical whether to return an error if x is not a timeseries.

---

is\_uninitializedField *Check if uninitializedField*

---

**Description**

Check if uninitializedField

**Usage**

```
is_uninitializedField(x, msg = "Zelig model has not been estimated.",  
  fail = TRUE)
```

**Arguments**

x	a zelig.out method
msg	character string with the error message to return if fail = TRUE.
fail	logical whether to return an error if x uninitialized.

---

is\_varying *Check if the values in a vector vary*

---

**Description**

Check if the values in a vector vary

**Usage**

```
is_varying(x, msg = "Vector does not vary.", fail = TRUE)
```

**Arguments**

x	a vector
msg	character string with the error message to return if fail = TRUE.
fail	logical whether to return an error if x does not vary.

---

is_zelig	<i>Check if is a zelig object</i>
----------	-----------------------------------

---

**Description**

Check if is a zelig object

**Usage**

```
is_zelig(x, fail = TRUE)
```

**Arguments**

x	an object
fail	logical whether to return an error if x is not a Zelig object.

---

is_zeligei	<i>Check if an object was created with ZeligEI</i>
------------	--

---

**Description**

Check if an object was created with ZeligEI

**Usage**

```
is_zeligei(x, msg = "Function is not relevant for ZeligEI objects.",  
           fail = TRUE)
```

**Arguments**

x	a zelig object
msg	character string with the error message to return if fail = TRUE.
fail	logical whether to return an error if x is not a timeseries.

---

klein	<i>Simulation Data for model Two-Stage Least Square (twosls) that corresponds to method 2SLS of systemfit</i>
-------	---

---

**Description**

Dataframe contains annual observations of US economy from 1920 to 1940. The columns are, Year, C=Consumption, P=Corporate profits, P1=Previous year corporate profit, Wtot=Total wage, Wp=Private wage bill, Wg=Government wage bill, I=Investment, K1=Previous year capital stock, X=GNP, G=Government spending, T=Taxes, X1=Previous year GNP, Tm=Year-1931.

**Usage**

```
data(klein)
```

**Format**

A table containing 14 variables ("year", "C", "P", "P1", "Wtot", "Wp", "Wg", "I", "K1", "X", "G", "T", "X1", "Tm") and 21 observations.

**Source**

<http://pages.stern.nyu.edu/~wgreene/Text/econometricanalysis.htm>

---

kmenta	<i>Simulation Data for model Three-Stage Least Square (threesls) that corresponds to method 3SLS of systemfit</i>
--------	---

---

**Description**

Dataframe contains 20 annual observations of a supply/demand model with 5 variables. Columns are q=Food consumption per capita, p=Ratio of food price to general consumer prices, d=Disposable income in constant dollars, f=Ratio of preceding year's prices received by farmers to general consumer prices, a=Time index.

**Usage**

```
data(kmenta)
```

**Format**

A table containing 5 variables ("q", "p", "d", "f", "a") and 20 observations.

---

macro

*Macroeconomic Data*

---

### **Description**

Selected macroeconomic indicators for Austria, Belgium, Canada, Denmark, Finland, France, Italy, Japan, the Netherlands, Norway, Sweden, the United Kingdom, the United States, and West Germany for the period 1966-1990.

### **Usage**

data(macro)

### **Format**

A table containing 6 variables ("country", "year", "gdp", "unem", "capmob", and "trade") and 350 observations.

### **Source**

ICPSR

### **References**

King, Gary, Michael Tomz and Jason Wittenberg. ICPSR Publication Related Archive, 1225.

King, Gary, Michael Tomz and Jason Wittenberg (2000). "Making the Most of Statistical Analyses: Improving Interpretation and Presentation," *American Journal of Political Science*, vol. 44, pp. 341-355.

---

MatchIt.url

*Table of links for Zelig*

---

### **Description**

Table of links for help.zelig for the companion MatchIt package.

---

Median	<i>Compute the Statistical Median of a Vector</i>
--------	---

---

**Description**

Compute the Statistical Median of a Vector

**Usage**

```
Median(x, na.rm = NULL)
```

**Arguments**

x	a vector of numeric or ordered values
na.rm	ignored

**Value**

the median of the vector

**Author(s)**

Matt Owen

---

mexico	<i>Voting Data from the 1988 Mexican Presidential Election</i>
--------	--

---

**Description**

This dataset contains voting data for the 1988 Mexican presidential election.

**Usage**

```
data(mexico)
```

**Format**

A table containing 33 variables and 1,359 observations.

**Source**

ICPSR

**References**

King, Gary, Michael Tomz and Jason Wittenberg (2000). "Making the Most of Statistical Analyses: Improving Interpretation and Presentation," *American Journal of Political Science*, vol. 44, pp. 341-355.

King, Tomz and Wittenberg. ICPSR Publication Related Archive, 1255.

---

mi	<i>Enables backwards compatibility for preparing non-amelia imputed data sets for zelig.</i>
----	--

---

**Description**

See [to\\_zelig\\_mi](#)

**Usage**

```
mi(...)
```

**Arguments**

... a set of data.frame's

**Value**

an mi object composed of a list of data frames.

---

mid	<i>Militarized Interstate Disputes</i>
-----	--

---

**Description**

A small sample from the militarized interstate disputes (MID) database.

**Usage**

```
data(mid)
```

**Format**

A table containing 6 variables ("conflict", "major", "contig", "power", "maxdem", "mindem", and "years") and 3,126 observations. For full variable descriptions, please see King and Zeng, 2001.

**Source**

Militarized Interstate Disputes database



**References**

King, Gary, and Lanche Zeng (2001). "Explaining Rare Events in International Relations," *International Organization*, vol. 55, no. 3, pp. 693-715.

Jones, Daniel M., Stuart A. Bremer and David Singer (1996). "Militarized Interstate Disputes, 1816-1992: Rationale, Coding Rules, and Empirical Patterns," *Conflict Management and Peace Science*, vol. 15, no. 2, pp. 163-213.

---

Mode	<i>Compute the Statistical Mode of a Vector</i>
------	---

---

**Description**

Compute the Statistical Mode of a Vector

**Usage**

Mode(x)

**Arguments**

x                      a vector of numeric, factor, or ordered values

**Value**

the statistical mode of the vector. If more than one mode exists, the last one in the factor order is arbitrarily chosen (by design)

**Author(s)**

Christopher Gandrud and Matt Owen

---

model_lookup_df	<i>Instructions for how to convert non-Zelig fitted model objects to Zelig. Used in to_zelig</i>
-----------------	--

---

**Description**

Instructions for how to convert non-Zelig fitted model objects to Zelig. Used in to\_zelig

**Usage**

model\_lookup\_df

**Format**

An object of class `data.frame` with 9 rows and 4 columns.

---

names,Zelig-method      *Names method for Zelig objects*

---

### Description

Names method for Zelig objects

### Usage

```
## S4 method for signature 'Zelig'
names(x)
```

### Arguments

x                      An Object of Class Zelig

---

newpainters              *The Discretized Painter's Data of de Piles*

---

### Description

The original painters data contain the subjective assessment, on a 0 to 20 integer scale, of 54 classical painters. The newpainters data discretizes the subjective assessment by quartiles with thresholds 25%, 50%, 75%. The painters were assessed on four characteristics: composition, drawing, colour and expression. The data is due to the Eighteenth century art critic, de Piles.

### Usage

```
data(newpainters)
```

### Format

A table containing 5 variables ("Composition", "Drawing", "Colour", "Expression", and "School") and 54 observations.

### Source

A. J. Weekes (1986). "A Genstat Primer". Edward Arnold.

M. Davenport and G. Studdert-Kennedy (1972). "The statistical analysis of aesthetic judgement: an exploration." *Applied Statistics*, vol. 21, pp. 324–333.

I. T. Jolliffe (1986) "Principal Component Analysis." Springer.

### References

Venables, W. N. and Ripley, B. D. (2002) "Modern Applied Statistics with S," Fourth edition. Springer.

---

or_summary	<i>Find odds ratios for coefficients and standard errors for glm.summary class objects</i>
------------	--

---

### Description

Find odds ratios for coefficients and standard errors for glm.summary class objects

### Usage

```
or_summary(obj, label_mod_coef = "(OR)", label_mod_se = "(OR)")
```

### Arguments

obj                    a glm.summary class object  
label\_mod\_coef        character string for how to modify the coefficient label.  
label\_mod\_se         character string for how to modify the standard error label.

---

PERisk	<i>Political Economic Risk Data from 62 Countries in 1987</i>
--------	---

---

### Description

Political Economic Risk Data from 62 Countries in 1987.

### Usage

```
data(PERisk)
```

### Format

A data frame with 62 observations on the following 6 variables. All data points are from 1987. See Quinn (2004) for more details.

country: a factor with levels 'Argentina' 'Australia' 'Austria' 'Bangladesh' 'Belgium' 'Bolivia' 'Botswana' 'Brazil' 'Burma' 'Cameroon' 'Canada' 'Chile' 'Colombia' 'Congo-Kinshasa' 'Costa Rica' 'Cote d'Ivoire' 'Denmark' 'Dominican Republic' 'Ecuador' 'Finland' 'Gambia, The' 'Ghana' 'Greece' 'Hungary' 'India' 'Indonesia' 'Iran' 'Ireland' 'Israel' 'Italy' 'Japan' 'Kenya' 'Korea, South' 'Malawi' 'Malaysia' 'Mexico' 'Morocco' 'New Zealand' 'Nigeria' 'Norway' 'Papua New Guinea' 'Paraguay' 'Philippines' 'Poland' 'Portugal' 'Sierra Leone' 'Singapore' 'South Africa' 'Spain' 'Sri Lanka' 'Sweden' 'Switzerland' 'Syria' 'Thailand' 'Togo' 'Tunisia' 'Turkey' 'United Kingdom' 'Uruguay' 'Venezuela' 'Zambia' 'Zimbabwe'

courts: an ordered factor with levels '0' < '1'. 'courts' is an indicator of whether the country in question is judged to have an independent judiciary. From Henisz (2002).

barb2: a numeric vector giving the natural log of the black market premium in each country. The black market premium is coded as the black market exchange rate (local currency per dollar) divided by the official exchange rate minus 1. From Marshall, Gurr, and Harff (2002).

prsexp2: an ordered factor with levels '0' < '1' < '2' < '3' < '4' < '5', giving the lack of expropriation risk. From Marshall, Gurr, and Harff (2002).

prscorr2: an ordered factor with levels '0' < '1' < '2' < '3' < '4' < '5', measuring the lack of corruption. From Marshall, Gurr, and Harff (2002).

gdpw2: a numeric vector giving the natural log of real GDP per worker in 1985 international prices. From Alvarez et al. (1999).

### Source

Mike Alvarez, Jose Antonio Cheibub, Fernando Limongi, and Adam Przeworski. 1999. "ACLP Political and Economic Database." <URL: <http://www.ssc.upenn.edu/~cheibub/data/>>.

Witold J. Henisz. 2002. "The Political Constraint Index (POLCON) Dataset." \<URL: <http://www-management.wharton.upenn.edu/henisz/POLCON/ContactInfo.html>>.

Monty G. Marshall, Ted Robert Gurr, and Barbara Harff. 2002. "State Failure Task Force Problem Set." <URL: <http://www.cidcm.umd.edu/inscr/stfail/index.htm>>.

### References

Kevin M. Quinn. 2004. "Bayesian Factor Analysis for Mixed Ordinal and Continuous Response." *Political Analysis*. Vol. 12, pp.338–353.

---

*plot,Zelig,ANY-method* *Plot method for Zelig objects*

---

### Description

Plot method for Zelig objects

### Usage

```
## S4 method for signature 'Zelig,ANY'
plot(x, y, ...)
```

### Arguments

x	An Object of Class Zelig
y	unused
...	Additional parameters to be passed to plot

---

predict,Zelig-method    *Method for getting predicted values from Zelig objects*

---

**Description**

Method for getting predicted values from Zelig objects

**Usage**

```
## S4 method for signature 'Zelig'
predict(object, ...)
```

**Arguments**

object	An Object of Class Zelig
...	Additional parameters to be passed to predict

---

qi.plot                      *Default Plot Design For Zelig Model QI's*

---

**Description**

Default Plot Design For Zelig Model QI's

**Usage**

```
qi.plot(obj, ...)
```

**Arguments**

obj	A reference class zelig5 object
...	Parameters to be passed to the 'truehist' function which is implicitly called for numeric simulations

**Author(s)**

James Honaker with panel layouts from Matt Owen

---

qi_slimmer	<i>Find the median and a central interval of simulated quantity of interest distributions</i>
------------	---

---

## Description

Find the median and a central interval of simulated quantity of interest distributions

## Usage

```
qi_slimmer(df, qi_type = "ev", ci = 0.95)
```

## Arguments

df	a tidy-formatted data frame of simulated quantities of interest created by <a href="#">zelig_qi_to_df</a> .
qi_type	character string either ev or pv for returning the central intervals for the expected value or predicted value, respectively.
ci	numeric. The central interval to return, expressed on the $(0, 100]$ or the equivalent $(0, 1]$ interval.

## Details

A tidy-formatted data frame with the following columns:

- The values fitted with [setx](#)
- qi\_ci\_min: the minimum value of the central interval specified with ci
- qi\_ci\_median: the median of the simulated quantity of interest distribution
- qi\_ci\_max: the maximum value of the central interval specified with ci

## Author(s)

Christopher Gandrud

## See Also

[zelig\\_qi\\_to\\_df](#)

## Examples

```
library(dplyr)
qi.central.interval <- zelig(Petal.Width ~ Petal.Length + Species,
  data = iris, model = "ls") %>%
  setx(Petal.Length = 2:4, Species = "setosa") %>%
  sim() %>%
  zelig_qi_to_df() %>%
  qi_slimmer()
```

---

 residuals,Zelig-method

*Method for extracting residuals from Zelig objects*


---

**Description**

Method for extracting residuals from Zelig objects

**Usage**

```
## S4 method for signature 'Zelig'
residuals(object)
```

**Arguments**

object            An Object of Class Zelig

---

rocplot

*Receiver Operator Characteristic Plots*


---

**Description**

The 'rocplot' command generates a receiver operator characteristic plot to compare the in-sample (default) or out-of-sample fit for two logit or probit regressions.

**Usage**

```
rocplot(z1, z2,
  cutoff = seq(from=0, to=1, length=100), lty1="solid",
  lty2="dashed", lwd1=par("lwd"), lwd2=par("lwd"),
  col1=par("col"), col2=par("col"),
  main="ROC Curve",
  xlab = "Proportion of 1's Correctly Predicted",
  ylab="Proportion of 0's Correctly Predicted",
  plot = TRUE,
  ...
)
```

**Arguments**

z1                first model  
 z2                second model  
 cutoff            A vector of cut-off values between 0 and 1, at which to evaluate the proportion of 0s and 1s correctly predicted by the first and second model. By default, this is 100 increments between 0 and 1 inclusive

lty1	the line type of the first model (defaults to 'line')
lty2	the line type of the second model (defaults to 'dashed')
lwd1	the line width of the first model (defaults to 1)
lwd2	the line width of the second model (defaults to 1)
col1	the color of the first model (defaults to 'black')
col2	the color of the second model (defaults to 'black')
main	a title for the plot (defaults to "ROC Curve")
xlab	a label for the X-axis
ylab	a label for the Y-axis
plot	whether to generate a plot to the selected device
...	additional parameters to be passed to the plot

**Value**

if plot is TRUE, rocplot simply generates a plot. Otherwise, a list with the following is produced:

roc1	a matrix containing a vector of x-coordinates and y-coordinates corresponding to the number of ones and zeros correctly predicted for the first model.
roc2	a matrix containing a vector of x-coordinates and y-coordinates corresponding to the number of ones and zeros correctly predicted for the second model.
area1	the area under the first ROC curve, calculated using Reimann sums.
area2	the area under the second ROC curve, calculated using Reimann sums.

---

sanction

*Multilateral Economic Sanctions*

---

**Description**

Data on bilateral sanctions behavior for selected years during the general period 1939-1983. This data contains errors that have since been corrected. Please contact Lisa Martin before using this data for publication.

**Usage**

```
data(sanction)
```

**Format**

A table containing 8 variables ("mil", "coop", "target", "import", "export", "cost", "num", and "ncost") and 78 observations. For full variable description, see Martin, 1992.

**Source**

Martin, 1992



**References**

Martin, Lisa (1992). *Coercive Cooperation: Explaining Multilateral Economic Sanctions*, Princeton: Princeton University Press.

---

 seatshare

*Left Party Seat Share in 11 OECD Countries*


---

**Description**

This data set contains time-series data of the seat shares in the lower legislative house of left leaning parties over time, as well as the level of unemployment. Data follows the style used in Hibbs (1977).

**Usage**

```
data(seatshare)
```

**Format**

A table containing N variables ("country","year","unemp","leftseat") and 384 observations split across 11 countries.

**Source**

OECD data and Mackie and Rose (1991), extended to further years.

**References**

Douglas A. Hibbs. (1977). *Political Parties and Macroeconomic Policy*. American Political Science Review 71(4):1467-1487.

Thomas T. Mackie and Richard Rose. (1991). *The International Almanac of Electoral History* Macmillan: London.

---

 setx

*Setting Explanatory Variable Values*


---

**Description**

The setx function uses the variables identified in the formula generated by zelig and sets the values of the explanatory variables to the selected values. Use setx after zelig and before sim to simulate quantities of interest.

**Usage**

```
setx(obj, fn = NULL, data = NULL, cond = FALSE, ...)
```

## Arguments

<code>obj</code>	output object from <code>zelig</code>
<code>fn</code>	a list of functions to apply to the data frame
<code>data</code>	a new data frame used to set the values of explanatory variables. If <code>data = NULL</code> (the default), the data frame called in <code>zelig</code> is used
<code>cond</code>	a logical value indicating whether unconditional (default) or conditional (choose <code>cond = TRUE</code> ) prediction should be performed. If you choose <code>cond = TRUE</code> , <code>setx</code> will coerce <code>fn = NULL</code> and ignore the additional arguments in <code>...</code> . If <code>cond = TRUE</code> and <code>data = NULL</code> , <code>setx</code> will prompt you for a data frame.
<code>...</code>	user-defined values of specific variables for overwriting the default values set by the function <code>fn</code> . For example, adding <code>var1 = mean(data\$var1)</code> or <code>x1 = 12</code> explicitly sets the value of <code>x1</code> to 12. In addition, you may specify one explanatory variable as a range of values, creating one observation for every unique value in the range of values

## Details

This documentation describes the `setx` Zelig 4 compatibility wrapper function.

## Value

The output is returned in a field to the Zelig object. For unconditional prediction, `x.out` is a model matrix based on the specified values for the explanatory variables. For multiple analyses (i.e., when choosing the `by` option in `zelig`, `setx` returns the selected values calculated over the entire data frame. If you wish to calculate values over just one subset of the data frame, the 5th subset for example, you may use: `x.out <- setx(z.out[[5]])`

## Author(s)

Matt Owen, Olivia Lau and Kosuke Imai

## See Also

The full Zelig manual may be accessed online at <http://docs.zeligproject.org/articles/>

## Examples

```
# Unconditional prediction:
data(turnout)
z.out <- zelig(vote ~ race + educate, model = 'logit', data = turnout)
x.out <- setx(z.out)
s.out <- sim(z.out, x = x.out)
```

---

 setx1

*Setting Explanatory Variable Values for First Differences*


---

## Description

This documentation describes the `setx1` Zelig 4 compatibility wrapper function. The wrapper is primarily useful for setting fitted values for creating first differences in piped workflows.

## Usage

```
setx1(obj, fn = NULL, data = NULL, cond = FALSE, ...)
```

## Arguments

<code>obj</code>	output object from <code>zelig</code>
<code>fn</code>	a list of functions to apply to the data frame
<code>data</code>	a new data frame used to set the values of explanatory variables. If <code>data = NULL</code> (the default), the data frame called in <code>zelig</code> is used
<code>cond</code>	a logical value indicating whether unconditional (default) or conditional (choose <code>cond = TRUE</code> ) prediction should be performed. If you choose <code>cond = TRUE</code> , <code>setx1</code> will coerce <code>fn = NULL</code> and ignore the additional arguments in <code>...</code> . If <code>cond = TRUE</code> and <code>data = NULL</code> , <code>setx1</code> will prompt you for a data frame.
<code>...</code>	user-defined values of specific variables for overwriting the default values set by the function <code>fn</code> . For example, adding <code>var1 = mean(data\$var1)</code> or <code>x1 = 12</code> explicitly sets the value of <code>x1</code> to 12. In addition, you may specify one explanatory variable as a range of values, creating one observation for every unique value in the range of values

## Value

The output is returned in a field to the Zelig object. For unconditional prediction, `x.out` is a model matrix based on the specified values for the explanatory variables. For multiple analyses (i.e., when choosing the `by` option in `zelig`, `setx1` returns the selected values calculated over the entire data frame. If you wish to calculate values over just one subset of the data frame, the 5th subset for example, you may use: `x.out <- setx(z.out[[5]])`

## Author(s)

Christopher Gandrud, Matt Owen, Olivia Lau, Kosuke Imai

## See Also

The full Zelig manual may be accessed online at <http://docs.zeligproject.org/articles/>

**Examples**

```

library(dplyr) # contains pipe operator %>%
data(turnout)

# plot first differences
zelig(Fertility ~ Education, data = swiss, model = 'ls') %>%
  setx(z4, Education = 10) %>%
  setx1(z4, Education = 30) %>%
  sim() %>%
  plot()

```

---

sim	<i>Generic Method for Computing and Organizing Simulated Quantities of Interest</i>
-----	---

---

**Description**

Simulate quantities of interest from the estimated model output from `zelig()` given specified values of explanatory variables established in `setx()`. For classical *maximum likelihood* models, `sim()` uses asymptotic normal approximation to the log-likelihood. For *Bayesian models*, Zelig simulates quantities of interest from the posterior density, whenever possible. For *robust Bayesian models*, simulations are drawn from the identified class of Bayesian posteriors. Alternatively, you may generate quantities of interest using bootstrapped parameters.

**Usage**

```

sim(obj, x, x1, y = NULL, num = 1000, bootstrap = F, bootfn = NULL,
    cond.data = NULL, ...)

```

**Arguments**

<code>obj</code>	output object from <code>zelig</code>
<code>x</code>	values of explanatory variables used for simulation, generated by <code>setx</code> . Not if omitted, then <code>sim</code> will look for values in the reference class object
<code>x1</code>	optional values of explanatory variables (generated by a second call of <code>setx</code> ) particular computations of quantities of interest
<code>y</code>	a parameter reserved for the computation of particular quantities of interest (average treatment effects). Few models currently support this parameter
<code>num</code>	an integer specifying the number of simulations to compute
<code>bootstrap</code>	currently unsupported
<code>bootfn</code>	currently unsupported
<code>cond.data</code>	currently unsupported
<code>...</code>	arguments reserved future versions of Zelig

## Details

This documentation describes the `sim` Zelig 4 compatibility wrapper function.

## Value

The output stored in `s.out` varies by model. Use the `names` function to view the output stored in `s.out`. Common elements include:

<code>x</code>	the <code>setx</code> values for the explanatory variables, used to calculate the quantities of interest (expected values, predicted values, etc.).
<code>x1</code>	the optional <code>setx</code> object used to simulate first differences, and other model-specific quantities of interest, such as risk-ratios.
<code>call</code>	the options selected for <code>sim</code> , used to replicate quantities of interest.
<code>zelig.call</code>	the original function and options for <code>zelig</code> , used to replicate analyses.
<code>num</code>	the number of simulations requested.
<code>par</code>	the parameters (coefficients, and additional model-specific parameters). You may wish to use the same set of simulated parameters to calculate quantities of interest rather than simulating another set.
<code>qi\sev</code>	simulations of the expected values given the model and <code>x</code> .
<code>qi\spr</code>	simulations of the predicted values given by the fitted values.
<code>qi\afd</code>	simulations of the first differences (or risk difference for binary models) for the given <code>x</code> and <code>x1</code> . The difference is calculated by subtracting the expected values given <code>x</code> from the expected values given <code>x1</code> . (If do not specify <code>x1</code> , you will not get first differences or risk ratios.)
<code>qi\rr</code>	simulations of the risk ratios for binary and multinomial models. See specific models for details.
<code>qi\state.ev</code>	simulations of the average expected treatment effect for the treatment group, using conditional prediction. Let $t_i$ be a binary explanatory variable defining the treatment ( $t_i = 1$ ) and control ( $t_i = 0$ ) groups. Then the average expected treatment effect for the treatment group is

$$\frac{1}{n} \sum_{i=1}^n [Y_i(t_i = 1) - E[Y_i(t_i = 0) \mid t_i = 1]],$$

where  $Y_i(t_i = 1)$  is the value of the dependent variable for observation  $i$  in the treatment group. Variation in the simulations are due to uncertainty in simulating  $E[Y_i(t_i = 0)]$ , the counterfactual expected value of  $Y_i$  for observations in the treatment group, under the assumption that everything stays the same except that the treatment indicator is switched to  $t_i = 0$ .

<code>qi\state.pr</code>	simulations of the average predicted treatment effect for the treatment group, using conditional prediction. Let $t_i$ be a binary explanatory variable defining the treatment ( $t_i = 1$ ) and control ( $t_i = 0$ ) groups. Then the average predicted treatment effect for the treatment group is
--------------------------	---

$$\frac{1}{n} \sum_{i=1}^n [Y_i(t_i = 1) - Y_i(\widehat{t_i = 0}) \mid t_i = 1],$$

where  $Y_i(t_i = 1)$  is the value of the dependent variable for observation  $i$  in the treatment group. Variation in the simulations are due to uncertainty in simulating  $Y_i(\widehat{t_i = 0})$ , the counterfactual predicted value of  $Y_i$  for observations in the treatment group, under the assumption that everything stays the same except that the treatment indicator is switched to  $t_i = 0$ .

### Author(s)

Christopher Gandrud, Matt Owen, Olivia Lau and Kosuke Imai

---

simulations.plot

*Plot Quantities of Interest in a Zelig-fashion*

---

### Description

Various graph generation for different common types of simulated results from Zelig

### Usage

```
simulations.plot(y, y1=NULL, xlab="", ylab="", main="", col=NULL, line.col=NULL,
axisnames=TRUE)
```

### Arguments

y	A matrix or vector of simulated results generated by Zelig, to be graphed.
y1	For comparison of two sets of simulated results at different choices of covariates, this should be an object of the same type and dimension as y. If no comparison is to be made, this should be NULL.
xlab	Label for the x-axis.
ylab	Label for the y-axis.
main	Main plot title.
col	A vector of colors. Colors will be used in turn as the graph is built for main plot objects. For nominal/categorical data, this colors renders as the bar color, while for numeric data it renders as the background color.
line.col	A vector of colors. Colors will be used in turn as the graph is built for line color shading of plot objects.
axisnames	a character-vector, specifying the names of the axes

### Value

nothing

### Author(s)

James Honaker

---

`sna.ex`*Simulated Example of Social Network Data*

---

**Description**

This data set contains five sociomatrices of simulated data social network data.

**Usage**

```
data(sna.ex)
```

**Format**

Each variable in the dataset is a 25 by 25 matrix of simulated social network data. The matrices are labeled "Var1", "Var2", "Var3", "Var4", and "Var5".

**Source**

fictitious

---

`summary,Zelig-method` *Summary method for Zelig objects*

---

**Description**

Summary method for Zelig objects

**Usage**

```
## S4 method for signature 'Zelig'  
summary(object, ...)
```

**Arguments**

<code>object</code>	An Object of Class Zelig
<code>...</code>	Additional parameters to be passed to summary

---

summary.Arima	<i>Summary of an object of class Arima</i>
---------------	--

---

**Description**

Summary of an object of class Arima

**Usage**

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

**Arguments**

object	An object of class Arima
...	Additional parameters

**Value**

The original object

---

SupremeCourt	<i>U.S. Supreme Court Vote Matrix</i>
--------------	---------------------------------------

---

**Description**

This dataframe contains a matrix votes cast by U.S. Supreme Court justices in all cases in the 2000 term.

**Usage**

```
data(SupremeCourt)
```

**Format**

The dataframe has contains data for justices Rehnquist, Stevens, O'Connor, Scalia, Kennedy, Souter, Thomas, Ginsburg, and Breyer for the 2000 term of the U.S. Supreme Court. It contains data from 43 non-unanimous cases. The votes are coded liberal (1) and conservative (0) using the protocol of Spaeth (2003). The unit of analysis is the case citation (ANALU=0). We are concerned with formally decided cases issued with written opinions, after full oral argument and cases decided by an equally divided vote (DECTYPE=1,5,6,7).

**Source**

Harold J. Spaeth (2005). "Original United States Supreme Court Database: 1953-2004 Terms." <URL:<http://www.as.uky.edu/polisci/ulmerproject/sctdata.htm>>.



---

 swiss

*Swiss Fertility and Socioeconomic Indicators (1888) Data*


---

**Description**

Standardized fertility measure and socio-economic indicators for each of 47 French-speaking provinces of Switzerland at about 1888.

**Usage**

```
data(swiss)
```

**Format**

A data frame with 47 observations on 6 variables, each of which is in percent, i.e., in [0,100].

[,1] Fertility Ig, "common standardized fertility measure" [,2] Agriculture [,3] Examination nation  
[,4] Education [,5] Catholic [,6] Infant.Mortality live births who live less than 1 year.

All variables but 'Fert' give proportions of the population.

**Source**

Project "16P5", pages 549-551 in

Mosteller, F. and Tukey, J. W. (1977) "Data Analysis and Regression: A Second Course in Statistics". Addison-Wesley, Reading Mass.

indicating their source as "Data used by permission of Franice van de Walle. Office of Population Research, Princeton University, 1976. Unpublished data assembled under NICHD contract number No 1-HD-O-2077."

**References**

Becker, R. A., Chambers, J. M. and Wilks, A. R. (1988) "The New S Language". Wadsworth & Brooks/Cole.

---

 table.levels

*Create a table, but ensure that the correct columns exist. In particular, this allows for entires with zero as a value, which is not the default for standard tables*

---

**Description**

Create a table, but ensure that the correct columns exist. In particular, this allows for entires with zero as a value, which is not the default for standard tables

**Usage**

```
table.levels(x, levels, ...)
```

**Arguments**

x	a vector
levels	a vector of levels
...	parameters for table

**Value**

a table

**Author(s)**

Matt Owen

---

tobin

*Tobin's Tobit Data*

---

**Description**

Economists fit a parametric censored data model called the 'tobit'. These data are from Tobin's original paper.

**Usage**

```
data(tobin)
```

**Format**

A data frame with 20 observations on the following 3 variables.

durable: Durable goods purchase

age: Age in years

quant: Liquidity ratio (x 1000)

**Source**

J. Tobin, Estimation of relationships for limited dependent variables, *Econometrica*, v26, 24-36, 1958.

---

to_zelig	<i>Coerce a non-Zelig fitted model object to a Zelig class object</i>
----------	---

---

**Description**

Coerce a non-Zelig fitted model object to a Zelig class object

**Usage**

```
to_zelig(obj)
```

**Arguments**

obj                    a fitted model object fitted using `lm` and many using `glm`. Note: more intended in future Zelig releases.

**Author(s)**

Christopher Gandrud and Ista Zhan

**Examples**

```
library(dplyr)
lm.out <- lm(Fertility ~ Education, data = swiss)

z.out <- to_zelig(lm.out)

# to_zelig called from within setx
setx(z.out) %>% sim() %>% plot()
```

---

to_zelig_mi	<i>Bundle Multiply Imputed Data Sets into an Object for Zelig</i>
-------------	---

---

**Description**

This object prepares multiply imputed data sets so they can be used by `zelig`.

**Usage**

```
to_zelig_mi(...)
```

**Arguments**

...                    a set of `data.frame`'s or a single list of `data.frame`'s

**Value**

an `mi` object composed of a list of data frames.

**Note**

This function creates a list of `data.frame` objects, which resembles the storage of imputed data sets in the `amelia` object.

**Author(s)**

Matt Owen, James Honaker, and Christopher Gandrud

**Examples**

```
# create datasets
n <- 100
x1 <- runif(n)
x2 <- runif(n)
y <- rnorm(n)
data.1 <- data.frame(y = y, x = x1)
data.2 <- data.frame(y = y, x = x2)

# merge datasets into one object as if imputed datasets

mi.out <- to_zelig_mi(data.1, data.2)

# pass object in place of data argument
z.out <- zelig(y ~ x, model = "ls", data = mi.out)
```

---

turnout

*Turnout Data Set from the National Election Survey*

---

**Description**

This data set contains individual-level turnout data. It pools several American National Election Surveys conducted during the 1992 presidential election year. Only the first 2,000 observations (from a total of 15,837 observations) are included in the sample data.

**Usage**

```
data(turnout)
```

**Format**

A table containing 5 variables ("race", "age", "educate", "income", and "vote") and 2,000 observations.

**Source**

National Election Survey

**References**

King, Gary, Michael Tomz, Jason Wittenberg (2000). "Making the Most of Statistical Analyses: Improving Interpretation and Presentation," *American Journal of Political Science*, vol. 44, pp.341–355.

---

vcov,Zelig-method	<i>Variance-covariance method for Zelig objects</i>
-------------------	---

---

**Description**

Variance-covariance method for Zelig objects

**Usage**

```
## S4 method for signature 'Zelig'
vcov(object)
```

**Arguments**

object            An Object of Class Zelig

---

vcov_gee	<i>Find vcov for GEE models</i>
----------	---------------------------------

---

**Description**

Find vcov for GEE models

**Usage**

```
vcov_gee(obj)
```

**Arguments**

obj                a geeglm class object.

---

vcov_rq	<i>Find vcov for quantile regression models</i>
---------	---

---

**Description**

Find vcov for quantile regression models

**Usage**

```
vcov_rq(obj)
```

**Arguments**

obj            a rq class object.

---

voteincome	<i>Sample Turnout and Demographic Data from the 2000 Current Population Survey</i>
------------	--

---

**Description**

This data set contains turnout and demographic data from a sample of respondents to the 2000 Current Population Survey (CPS). The states represented are South Carolina and Arkansas. The data represent only a sample and results from this example should not be used in publication.

**Usage**

```
data(voteincome)
```

**Format**

A data frame containing 7 variables ("state", "year", "vote", "income", "education", "age", "female") and 1500 observations.

state a factor variable with levels equal to "AR" (Arkansas) and "SC" (South Carolina)

year an integer vector

vote an integer vector taking on values "1" (Voted) and "0" (Did Not Vote)

income an integer vector ranging from "4" (Less than \$5000) to "17" (Greater than \$75000) denoting family income. See the CPS codebook for more information on variable coding

education an integer vector ranging from "1" (Less than High School Education) to "4" (More than a College Education). See the CPS codebook for more information on variable coding

age an integer vector ranging from "18" to "85"

female an integer vector taking on values "1" (Female) and "0" (Male)

**Source**

Census Bureau Current Population Survey

**References**

<http://www.census.gov/cps>

---

Weimar

*1932 Weimar election data*

---

**Description**

This data set contains election results for 10 kreise (equivalent to precincts) from the 1932 Weimar (German) election.

**Usage**

`data(Weimar)`

**Format**

A table containing 11 variables and 10 observations. The variables are

**Nazi** Number of votes for the Nazi party

**Government** Number of votes for the Government

**Communists** Number of votes for the Communist party

**FarRight** Number of votes for far right parties

**Other** Number of votes for other parties, and non-voters

**shareunemployed** Proportion unemployed

**shareblue** Proportion working class

**sharewhite** Proportion white-collar workers

**sharedomestic** Proportion domestic servants

**shareprotestants** Proportion Protestant

**Source**

ICPSR

**Description**

The `zelig` function estimates a variety of statistical models. Use `zelig` output with `setx` and `sim` to compute quantities of interest, such as predicted probabilities, expected values, and first differences, along with the associated measures of uncertainty (standard errors and confidence intervals).

**Usage**

```
zelig(formula, model, data, ..., by = NULL, cite = TRUE)
```

**Arguments**

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
<code>model</code>	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <code>Amelia</code> or <code>to_zelig_mi</code> ).
<code>...</code>	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
<code>by</code>	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. For example, to run the same model on all fifty states, you could use: <code>z.out &lt;- zelig(y ~ x1 + x2, data = mydata, model = 'ls', by = 's')</code> . You may also use <code>by</code> to run models using <code>MatchIt</code> subclasses.
<code>cite</code>	If is set to 'TRUE' (default), the model citation will be printed to the console.

**Details**

This documentation describes the `zelig` Zelig 4 compatibility wrapper function.

Additional parameters available to many models include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.



- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

### Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

### Author(s)

Matt Owen, Kosuke Imai, Olivia Lau, and Gary King

### See Also

<http://docs.zeligproject.org/articles/>

---

Zelig-ar-class

*Time-Series Model with Autoregressive Disturbance*

---

### Description

Warning: `summary` does not work with timeseries models after simulation.

### Arguments

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
<code>model</code>	the name of a statistical model to estimate. For a list of supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <code>to_zelig_mi</code> ).
<code>...</code>	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.

by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. For example, to run the same model on all fifty states, you could use: <code>z.out &lt;- zelig(y ~ x1 + x2, data = mydata, model = 'ls', by = 's')</code> . You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
ts	The name of the variable containing the time indicator. This should be passed in as a string. If this variable is not provided, Zelig will assume that the data is already ordered by time.
cs	Name of a variable that denotes the cross-sectional element of the data, for example, country name in a dataset with time-series across different countries. As a variable name, this should be in quotes. If this is not provided, Zelig will assume that all observations come from the same unit over time, and should be pooled, but if provided, individual models will be run in each cross-section. If cs is given as an argument, ts must also be provided. Additionally, by must be NULL.
order	A vector of length 3 passed in as <code>c(p, d, q)</code> where p represents the order of the autoregressive model, d represents the number of differences taken in the model, and q represents the order of the moving average model.

### Details

Currently only the Reference class syntax for time series. This model does not accept Bootstraps or weights.

### Value

Depending on the class of model selected, zelig will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_ar.html](http://docs.zeligproject.org/articles/zelig_ar.html)

### Examples

```
data(seatshare)
subset <- seatshare[seatshare$country == "UNITED KINGDOM",]
ts.out <- zelig(formula = unemp ~ leftseat, model = "ar", ts = "year", data = subset)
summary(ts.out)
```

---

Zelig-arma-class	<i>Autoregressive and Moving-Average Models with Integration for Time-Series Data</i>
------------------	---

---

## Description

Warning: summary does not work with timeseries models after simulation.

## Arguments

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to zelig, relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. For example, to run the same model on all fifty states, you could use: <code>z.out &lt;- zelig(y ~ x1 + x2, data = mydata, model = 'ls', by = 's</code> . You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
ts	The name of the variable containing the time indicator. This should be passed in as a string. If this variable is not provided, Zelig will assume that the data is already ordered by time.
cs	Name of a variable that denotes the cross-sectional element of the data, for example, country name in a dataset with time-series across different countries. As a variable name, this should be in quotes. If this is not provided, Zelig will assume that all observations come from the same unit over time, and should be pooled, but if provided, individual models will be run in each cross-section. If cs is given as an argument, ts must also be provided. Additionally, by must be NULL.
order	A vector of length 3 passed in as <code>c(p, d, q)</code> where p represents the order of the autoregressive model, d represents the number of differences taken in the model, and q represents the order of the moving average model.

**Details**

Currently only the Reference class syntax for time series. This model does not accept Bootstraps or weights.

**Value**

Depending on the class of model selected, zelig will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_arma.html](http://docs.zeligproject.org/articles/zelig_arma.html)

**Examples**

```
data(seatshare)
subset <- seatshare[seatshare$country == "UNITED KINGDOM",]
ts.out <- zarima$new()
ts.out$zelig(unemp ~ leftseat, order = c(1, 0, 1), data = subset)

# Set fitted values and simulate quantities of interest
ts.out$setx(leftseat = 0.75)
ts.out$setx1(leftseat = 0.25)
ts.out$sim()
```

---

Zelig-bayes-class

*Bayes Model object for inheritance across models in Zelig*


---

**Description**

Bayes Model object for inheritance across models in Zelig

**Methods**

`get_coef(nonlist = FALSE)` Get estimated model coefficients  
`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The zelig function estimates a variety of statistical models

---

Zelig-binchoice-class

*Binary Choice object for inheritance across models in Zelig*


---

**Description**

Binary Choice object for inheritance across models in Zelig

---

Zelig-binchoice-gee-class

*Object for Binary Choice outcomes in Generalized Estimating Equations for inheritance across models in Zelig*

---

### Description

Object for Binary Choice outcomes in Generalized Estimating Equations for inheritance across models in Zelig

---

Zelig-binchoice-survey-class

*Object for Binary Choice outcomes with Survey Weights for inheritance across models in Zelig*

---

### Description

Object for Binary Choice outcomes with Survey Weights for inheritance across models in Zelig

---

Zelig-class

*Zelig reference class*

---

### Description

Zelig website: <http://zeligproject.org/>

### Fields

fn R function to call to wrap  
 formula Zelig formula  
 weights [forthcoming]  
 name name of the Zelig model  
 data data frame or matrix  
 by split the data by factors  
 mi work with imputed dataset  
 idx model index  
 zelig.call Zelig function call  
 model.call wrapped function call  
 zelig.out estimated zelig model(s)  
 setx.out set values

setx.labels pretty-print qi  
 bsetx is x set?  
 bsetx1 is x1 set?  
 bsetrange is range set?  
 bsetrange1 is range1 set?  
 range range  
 range1 range1  
 test.statistics list of test statistics  
 sim.out simulated qi's  
 simparam simulated parameters  
 num number of simulations  
 authors Zelig model authors  
 zeligauthors Zelig authors  
 modelauthors wrapped model authors  
 packageauthors wrapped package authors  
 refs citation information  
 year model is released  
 description model description  
 url model URL  
 url.docs model documentation URL  
 category model category  
 vignette.url vignette URL  
 json JSON export  
 ljson JSON export  
 outcome JSON export  
 wrapper JSON export  
 explanatory JSON export  
 mcunit.test unit testing  
 with.feedback Feedback  
 robust.se return robust standard errors

### Methods

ATT(treatment, treated = 1, quietly = TRUE, num = NULL) Generic Method for Computing Simulated (Sample) Average Treatment Effects on the Treated  
 cite() Provide citation information about Zelig and Zelig model, and about wrapped package and wrapped model  
 feedback() Send feedback to the Zelig team

`from_zelig_model()` Extract the original fitted model object from a zelig call. Note only works for models using directly wrapped functions.  
`get_coef(nonlist = FALSE)` Get estimated model coefficients  
`get_df_residual()` Get residual degrees-of-freedom  
`get_fitted(...)` Get estimated fitted values  
`get_model_data()` Get data used to estimate the model  
`get_names()` Return Zelig object field names  
`get_predict(...)` Get predicted values  
`get_pvalue()` Get estimated model p-values  
`get_qi(qi = "ev", xvalue = "x", subset = NULL)` Get quantities of interest  
`get_residuals(...)` Get estimated model residuals  
`get_se()` Get estimated model standard errors  
`get_vcov()` Get estimated model variance-covariance matrix  
`graph(...)` Plot the quantities of interest  
`help()` Open the model vignette from <http://zeligproject.org/>  
`packagename()` Automatically retrieve wrapped package name  
`references(style = "sphinx")` Construct a reference list specific to a Zelig model.  
`set(..., fn = list(numeric = mean, ordered = Median))` Setting Explanatory Variable Values  
`sim(num = NULL)` Generic Method for Computing and Organizing Simulated Quantities of Interest  
`simATT(simparam, data, depvar, treatment, treated)` Simulate an Average Treatment on the Treated  
`summarise(...)` Display a Zelig object  
`summarize(...)` Display a Zelig object  
`toJSON()` Convert Zelig object to JSON format  
`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The zelig function estimates a variety of statistical models

## Description

Exponential Regression for Duration Dependent Variables

## Arguments

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <code>to_zelig_mi</code> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. For example, to run the same model on all fifty states, you could use: <code>z.out &lt;- zelig(y ~ x1 + x2, data = mydata, model = 'ls', by = 's</code> . You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
robust	defaults to FALSE. If TRUE, <code>zelig()</code> computes robust standard errors based on sandwich estimators and the options selected in <code>cluster</code> .
if	<code>robust = TRUE</code> , you may select a variable to define groups of correlated observations. Let $x3$ be a variable that consists of either discrete numeric values, character strings, or factors that define strata. Then <code>z.out &lt;- zelig(y ~ x1 + x2, robust = TRUE, cluster = "x3", model = "exp", data = mydata)</code> means that the observations can be correlated within the strata defined by the variable $x3$ , and that robust standard errors should be calculated according to those clusters. If <code>robust = TRUE</code> but <code>cluster</code> is not specified, <code>zelig()</code> assumes that each observation falls into its own cluster.

## Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.



**Value**

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**Methods**

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_exp.html](http://docs.zeligproject.org/articles/zelig_exp.html)

**Examples**

```
library(Zelig)
data(coalition)
library(survival)
z.out <- zelig(Surv(duration, ciepv12) ~ fract + numst2, model = "exp",
              data = coalition)
summary(z.out)
```

---

Zelig-factor-bayes-class

*Bayesian Factor Analysis*

---

**Description**

Bayesian Factor Analysis

**Arguments**

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $\sim Y1 + Y2 + Y3$ , where $Y1$ , $Y2$ , and $Y3$ are variables of interest in factor analysis (manifest variables), assumed to be normally distributed. The model requires a minimum of three manifest variables contained in the same dataset. The $+$ symbol means “inclusion” not “addition.”
<code>factors</code>	number of the factors to be fitted (defaults to 2).
<code>model</code>	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <code>Amelia</code> or <code>to_zelig_mi</code> ).

...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
<code>by</code>	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using <code>MatchIt</code> subclasses.
<code>cite</code>	If is set to 'TRUE' (default), the model citation will be printed to the console.

## Details

In addition, `zelig()` accepts the following additional arguments for model specification:

- `lambda.constraints`: list containing the equality or inequality constraints on the factor loadings. Choose from one of the following forms:
- `varname = list()`: by default, no constraints are imposed.
- `varname = list(d, c)`: constrains the `d`th loading for the variable named `varname` to be equal to `c`.
- `varname = list(d, +)`: constrains the `d`th loading for the variable named `varname` to be positive;
- `varname = list(d, -)`: constrains the `d`th loading for the variable named `varname` to be negative.
- `std.var`: defaults to FALSE (manifest variables are rescaled to zero mean, but retain observed variance). If TRUE, the manifest variables are rescaled to be mean zero and unit variance.

In addition, `zelig()` accepts the following additional inputs for `bayes.factor`:

- `burnin`: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- `mcmc`: number of the MCMC iterations after burnin (defaults to 20,000).
- `thin`: thinning interval for the Markov chain. Only every `thin`-th draw from the Markov chain is kept. The value of `mcmc` must be divisible by this value. The default value is 1.
- `verbose`: defaults to FALSE. If TRUE, the progress of the sampler (every 10
- `seed`: seed for the random number generator. The default is NA which corresponds to a random seed 12345.
- `Lambda.start`: starting values of the factor loading matrix  $\Lambda$ , either a scalar (all unconstrained loadings are set to that value), or a matrix with compatible dimensions. The default is NA, where the start value are set to be 0 for unconstrained factor loadings, and 0.5 or - 0.5 for constrained factor loadings (depending on the nature of the constraints).
- `Psi.start`: starting values for the uniquenesses, either a scalar (the starting values for all diagonal elements of  $\Psi$  are set to be this value), or a vector with length equal to the number of manifest variables. In the latter case, the starting values of the diagonal elements of  $\Psi$  take the values of `Psi.start`. The default value is NA where the starting values of the all the uniquenesses are set to be 0.5.
- `store.lambda`: defaults to TRUE, which stores the posterior draws of the factor loadings.
- `store.scores`: defaults to FALSE. If TRUE, stores the posterior draws of the factor scores. (Storing factor scores may take large amount of memory for a large number of draws or observations.)

The model also accepts the following additional arguments to specify prior parameters:

- $\mu_0$ : mean of the Normal prior for the factor loadings, either a scalar or a matrix with the same dimensions as  $\Lambda$ . If a scalar value, that value will be the prior mean for all the factor loadings. Defaults to 0.
- $L_0$ : precision parameter of the Normal prior for the factor loadings, either a scalar or a matrix with the same dimensions as  $\Lambda$ . If  $L_0$  takes a scalar value, then the precision matrix will be a diagonal matrix with the diagonal elements set to that value. The default value is 0, which leads to an improper prior.
- $a_0$ : the shape parameter of the Inverse Gamma prior for the uniquenesses is  $a_0/2$ . It can take a scalar value or a vector. The default value is 0.001.
- $b_0$ : the scale parameter of the Inverse Gamma prior for the uniquenesses is  $b_0/2$ . It can take a scalar value or a vector. The default value is 0.001.

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

## Methods

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

## See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_factorbayes.html](http://docs.zeligproject.org/articles/zelig_factorbayes.html)

## Examples

```
## Not run:
data(swiss)
names(swiss) <- c("Fert", "Agr", "Exam", "Educ", "Cath", "InfMort")
z.out <- zelig(~ Agr + Exam + Educ + Cath + InfMort,
model = "factor.bayes", data = swiss,
factors = 2, verbose = FALSE,
a0 = 1, b0 = 0.15, burnin = 500, mcmc = 5000)

z.out$geweke.diag()
z.out <- zelig(~ Agr + Exam + Educ + Cath + InfMort,
model = "factor.bayes", data = swiss, factors = 2,
```

```

lambda.constraints =
  list(Exam = list(1,"+"),
       Exam = list(2,"-"),
       Educ = c(2, 0),
       InfMort = c(1, 0)),
verbose = FALSE, a0 = 1, b0 = 0.15,
burnin = 500, mcmc = 5000)

z.out$geweke.diag()
z.out$heidel.diag()
z.out$raftery.diag()
summary(z.out)

## End(Not run)

```

---

Zelig-gamma-class

*Gamma Regression for Continuous, Positive Dependent Variables*


---

## Description

Gamma Regression for Continuous, Positive Dependent Variables

## Arguments

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

**Details**

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

**Value**

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_gamma.html](http://docs.zeligproject.org/articles/zelig_gamma.html)

**Examples**

```
library(Zelig)
data(coalition)
z.out <- zelig(duration ~ fract + numst2, model = "gamma", data = coalition)
summary(z.out)
```

---

Zelig-gamma-gee-class *Generalized Estimating Equation for Gamma Regression*

---

**Description**

Generalized Estimating Equation for Gamma Regression

**Arguments**

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The <code>+</code> symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
----------------------	--

model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <code>to_zelig_mi</code> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
corstr: character	string specifying the correlation structure: "independence", "exchangeable", "ar1", "unstructured" and "userdefined"
See	<code>geeglm</code> in package <code>geepack</code> for other function arguments.
id:	where <code>id</code> is a variable which identifies the clusters. The data should be sorted by <code>id</code> and should be ordered within each cluster when appropriate
corstr:	character string specifying the correlation structure: "independence", "exchangeable", "ar1", "unstructured" and "userdefined"
geeglm:	See <code>geeglm</code> in package <code>geepack</code> for other function arguments

## Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

## See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_gammagee.html](http://docs.zeligproject.org/articles/zelig_gammagee.html)

**Examples**

```
library(Zelig)
data(coalition)
coalition$cluster <- c(rep(c(1:62), 5), rep(c(63), 4))
sorted.coalition <- coalition[order(coalition$cluster),]
z.out <- zelig(duration ~ fract + numst2, model = "gamma.gee", id = "cluster",
              data = sorted.coalition, corstr = "exchangeable")
summary(z.out)
```

---

Zelig-gamma-survey-class

*Gamma Regression with Survey Weights*


---

**Description**

Gamma Regression with Survey Weights

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <i>Amelia</i> or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <i>zelig</i> , relevant for the model to be estimated.
by	a factor variable contained in <i>data</i> . If supplied, <i>zelig</i> will subset the data frame based on the levels in the <i>by</i> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <i>by</i> to run models using <i>MatchIt</i> subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

**Details**

Additional parameters available to this model include:

- **weights**: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.

- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

## See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_gammasurvey.html](http://docs.zeligproject.org/articles/zelig_gammasurvey.html)

## Examples

```
library(Zelig)
data(api, package="survey")
z.out1 <- zelig(api00 ~ meals + yr.rnd, model = "gamma.survey",
weights = ~pw, data = apistrat)
summary(z.out1)
```

---

Zelig-gee-class	<i>Generalized Estimating Equations Model object for inheritance across models in Zelig</i>
-----------------	---

---

## Description

Generalized Estimating Equations Model object for inheritance across models in Zelig

## Methods

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models



---

Zelig-glm-class	<i>Generalized Linear Model object for inheritance across models in Zelig</i>
-----------------	---

---

**Description**

Generalized Linear Model object for inheritance across models in Zelig

**Methods**

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

---

Zelig-ivreg-class	<i>Instrumental-Variable Regression</i>
-------------------	---

---

**Description**

Instrumental-Variable Regression

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means inclusion ' ' not addition." You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
formula	specification(s) of the regression relationship

<code>instruments</code>	the instruments. Either <code>instruments</code> is missing and formula has three parts as in $y \sim x1 + x2 \mid z1 + z2 + z3$ (recommended) or formula is $y \sim x1 + x2$ and <code>instruments</code> is a one-sided formula $\sim z1 + z2 + z3$ . Using <code>instruments</code> is not recommended with <code>zelig</code> .
<code>model, x, y</code>	logicals. If TRUE the corresponding components of the fit (the model frame, the model matrices, the response) are returned.
<code>...</code>	further arguments passed to methods. See also <a href="#">zelig</a> .

## Details

Additional parameters available to many models include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

Regressors and instruments for `ivreg` are most easily specified in a formula with two parts on the right-hand side, e.g.,  $y \sim x1 + x2 \mid z1 + z2 + z3$ , where  $x1$  and  $x2$  are the regressors and  $z1$ ,  $z2$ , and  $z3$  are the instruments. Note that exogenous regressors have to be included as instruments for themselves. For example, if there is one exogenous regressor `ex` and one endogenous regressor `en` with instrument `in`, the appropriate formula would be  $y \sim ex + en \mid ex + in$ . Equivalently, this can be specified as  $y \sim ex + en \mid . - en + in$ , i.e., by providing an update formula with a `.` in the second part of the formula. The latter is typically more convenient, if there is a large number of exogenous regressors.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

## Methods

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

## Source

`ivreg` is from Christian Kleiber and Achim Zeileis (2008). Applied Econometrics with R. New York: Springer-Verlag. ISBN 978-0-387-77316-2. URL <https://CRAN.R-project.org/package=AER>

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_ivreg.html](http://docs.zeligproject.org/articles/zelig_ivreg.html) Fit instrumental-variable regression by two-stage least squares. This is equivalent to direct instrumental-variables estimation when the number of instruments is equal to the number of predictors.

`zelig`, Greene, W. H. (1993) *Econometric Analysis*, 2nd ed., Macmillan.

**Examples**

```
library(Zelig)
library(dplyr) # for the pipe operator %>%
# load and transform data
data("CigarettesSW")
CigarettesSW$rprice <- with(CigarettesSW, price/cpi)
CigarettesSW$rincome <- with(CigarettesSW, income/population/cpi)
CigarettesSW$tdiff <- with(CigarettesSW, (taxs - tax)/cpi)
# log second stage independent variables, as logging internally for ivreg is
# not currently supported
CigarettesSW$log_rprice <- log(CigarettesSW$rprice)
CigarettesSW$log_rincome <- log(CigarettesSW$rincome)
z.out1 <- zelig(log(packs) ~ log_rprice + log_rincome |
log_rincome + tdiff + I(tax/cpi), data = CigarettesSW, subset = year == "1995", model = "ivreg")
summary(z.out1)
library(Zelig)
library(AER) # for sandwich vcov
library(dplyr) # for the pipe operator %>%

# load and transform data
data("CigarettesSW")
CigarettesSW$rprice <- with(CigarettesSW, price/cpi)
CigarettesSW$rincome <- with(CigarettesSW, income/population/cpi)
CigarettesSW$tdiff <- with(CigarettesSW, (taxs - tax)/cpi)

# log second stage independent variables, as logging internally for ivreg is
# not currently supported
CigarettesSW$log_rprice <- log(CigarettesSW$rprice)
CigarettesSW$log_rincome <- log(CigarettesSW$rincome)

# estimate model
z.out1 <- zelig(log(packs) ~ log_rprice + log_rincome |
log_rincome + tdiff + I(tax/cpi),
data = CigarettesSW,
model = "ivreg")
summary(z.out1)
```

**Description**

Bayesian Logit Regression

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to zelig, relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

**Details**

Additional parameters available to this model include:

- **weights**: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- **burnin**: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- **mcmc**: number of the MCMC iterations after burnin (defaults to 10,000).
- **thin**: thinning interval for the Markov chain. Only every thin-th draw from the Markov chain is kept. The value of mcmc must be divisible by this value. The default value is 1.
- **verbose**: defaults to FALSE. If TRUE, the progress of the sampler (every 10%) is printed to the screen.
- **seed**: seed for the random number generator. The default is NA which corresponds to a random seed of 12345.
- **beta.start**: starting values for the Markov chain, either a scalar or vector with length equal to the number of estimated coefficients. The default is NA, such that the maximum likelihood estimates are used as the starting values.

Use the following parameters to specify the model's priors:

- $b_0$ : prior mean for the coefficients, either a numeric vector or a scalar. If a scalar value, that value will be the prior mean for all the coefficients. The default is 0.
- $B_0$ : prior precision parameter for the coefficients, either a square matrix (with the dimensions equal to the number of the coefficients) or a scalar. If a scalar value, that value times an identity matrix will be the prior precision parameter. The default is 0, which leads to an improper prior.

Use the following arguments to specify optional output for the model:

- `bayes.resid`: defaults to FALSE. If TRUE, the latent Bayesian residuals for all observations are returned. Alternatively, users can specify a vector of observations for which the latent residuals should be returned.

### Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_logitbayes.html](http://docs.zeligproject.org/articles/zelig_logitbayes.html)

### Examples

```
data(turnout)
z.out <- zelig(vote ~ race + educate, model = "logit.bayes", data = turnout, verbose = FALSE)
```

---

Zelig-logit-class

*Logistic Regression for Dichotomous Dependent Variables*

---

### Description

Logistic Regression for Dichotomous Dependent Variables

### Arguments

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
----------------------	---

model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using <code>MatchIt</code> subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
below	(defaults to 0) The point at which the dependent variable is censored from below. If any values in the dependent variable are observed to be less than the censoring point, it is assumed that that particular observation is censored from below at the observed value. (See for a Bayesian implementation that supports both left and right censoring.)
robust	defaults to FALSE. If TRUE, <code>zelig()</code> computes robust standard errors based on sandwich estimators (see <a href="#">and</a> ) and the options selected in <code>cluster</code> .
if	<code>robust = TRUE</code> , you may select a variable to define groups of correlated observations. Let <code>x3</code> be a variable that consists of either discrete numeric values, character strings, or factors that define strata. Then <code>z.out &lt;- zelig(y ~ x1 + x2, robust = TRUE, cluster = "x3", model = "tobit", data = mydata)</code> means that the observations can be correlated within the strata defined by the variable <code>x3</code> , and that robust standard errors should be calculated according to those clusters. If <code>robust = TRUE</code> but <code>cluster</code> is not specified, <code>zelig()</code> assumes that each observation falls into its own cluster.

## Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

**Methods**

`show(signif.stars = FALSE, subset = NULL, bagging = FALSE)` Display a Zelig object

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_logit.html](http://docs.zeligproject.org/articles/zelig_logit.html)

**Examples**

```
library(Zelig)
data(turnout)
z.out1 <- zelig(vote ~ age + race, model = "logit", data = turnout,
               cite = FALSE)
summary(z.out1)
summary(z.out1, odds_ratios = TRUE)
x.out1 <- setx(z.out1, age = 36, race = "white")
s.out1 <- sim(z.out1, x = x.out1)
summary(s.out1)
plot(s.out1)
```

---

Zelig-logit-gee-class *Generalized Estimating Equation for Logit Regression*

---

**Description**

Generalized Estimating Equation for Logit Regression

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.

<code>by</code>	a factor variable contained in data. If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using <code>MatchIt</code> subclasses.
<code>cite</code>	If is set to 'TRUE' (default), the model citation will be printed to the console.
<code>id:</code>	where <code>id</code> is a variable which identifies the clusters. The data should be sorted by <code>id</code> and should be ordered within each cluster when appropriate
<code>corstr:</code>	character string specifying the correlation structure: "independence", "exchangeable", "ar1", "unstructured" and "userdefined"
<code>geeglm:</code>	See <code>geeglm</code> in package <code>geepack</code> for other function arguments

### Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

### Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_logitgee.html](http://docs.zeligproject.org/articles/zelig_logitgee.html)

### Examples

```
data(turnout)
turnout$cluster <- rep(c(1:200), 10)
sorted.turnout <- turnout[order(turnout$cluster),]

z.out1 <- zelig(vote ~ race + educate, model = "logit.gee",
id = "cluster", data = sorted.turnout)

summary(z.out1)
x.out1 <- setx(z.out1)
s.out1 <- sim(z.out1, x = x.out1)
summary(s.out1)
```



```
plot(s.out1)
```

---

Zelig-logit-survey-class

*Logit Regression with Survey Weights*

---

## Description

Logit Regression with Survey Weights

## Arguments

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to zelig, relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
below	(defaults to 0) The point at which the dependent variable is censored from below. If any values in the dependent variable are observed to be less than the censoring point, it is assumed that that particular observation is censored from below at the observed value. (See for a Bayesian implementation that supports both left and right censoring.)
robust	defaults to FALSE. If TRUE, zelig() computes robust standard errors based on sandwich estimators (see <a href="#">sandwich</a> ) and the options selected in cluster.
if	robust = TRUE, you may select a variable to define groups of correlated observations. Let $x3$ be a variable that consists of either discrete numeric values, character strings, or factors that define strata. Then <code>z.out &lt;- zelig(y ~ x1 + x2, robust = TRUE, cluster = "x3", model = "tobit", data = mydata)</code> means that the observations can be correlated within the strata defined by the variable $x3$ , and that robust standard errors should be calculated according to those clusters. If robust = TRUE but cluster is not specified, zelig() assumes that each observation falls into its own cluster.

**Details**

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

**Value**

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_logitsurvey.html](http://docs.zeligproject.org/articles/zelig_logitsurvey.html)

**Examples**

```
data(api, package = "survey")
apistrat$yr.rnd.numeric <- as.numeric(apistrat$yr.rnd == "Yes")
z.out1 <- zelig(yr.rnd.numeric ~ meals + mobility, model = "logit.survey",
               weights = apistrat$pw, data = apistrat)

summary(z.out1)
x.low <- setx(z.out1, meals= quantile(apistrat$meals, 0.2))
x.high <- setx(z.out1, meals= quantile(apistrat$meals, 0.8))
s.out1 <- sim(z.out1, x = x.low, x1 = x.high)
summary(s.out1)
plot(s.out1)
```

---

Zelig-lognorm-class    *Log-Normal Regression for Duration Dependent Variables*

---

**Description**

Log-Normal Regression for Duration Dependent Variables

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
robust	defaults to FALSE. If TRUE, <code>zelig()</code> computes robust standard errors based on sandwich estimators (see <a href="#">and</a> ) based on the options in <code>cluster</code> .
cluster	if <code>robust = TRUE</code> , you may select a variable to define groups of correlated observations. Let $x3$ be a variable that consists of either discrete numeric values, character strings, or factors that define strata. Then means that the observations can be correlated within the strata defined by the variable $x3$ , and that robust standard errors should be calculated according to those clusters. If <code>robust = TRUE</code> but <code>cluster</code> is not specified, <code>zelig()</code> assumes that each observation falls into its own cluster.

**Details**

Additional parameters available to many models include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

**Value**

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or

individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

## Methods

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

## See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_lognorm.html](http://docs.zeligproject.org/articles/zelig_lognorm.html)

## Examples

```
library(Zelig)
data(coalition)
z.out <- zelig(Surv(duration, ciepl2) ~ fract + numst2, model = "lognorm", data = coalition)
summary(z.out)
```

---

Zelig-ls-class

*Least Squares Regression for Continuous Dependent Variables*

---

## Description

Least Squares Regression for Continuous Dependent Variables

## Arguments

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
<code>model</code>	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <code>to_zelig_mi</code> ).
<code>...</code>	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
<code>by</code>	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.

`cite` If is set to 'TRUE' (default), the model citation will be printed to the console.

### Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

### Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

### Methods

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_ls.html](http://docs.zeligproject.org/articles/zelig_ls.html)

### Examples

```
library(Zelig)
data(macro)
z.out1 <- zelig(unem ~ gdp + capmob + trade, model = "ls", data = macro,
cite = FALSE)
summary(z.out1)
```

### Description

Warning: `summary` does not work with timeseries models after simulation.

## Arguments

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
ts	The name of the variable containing the time indicator. This should be passed in as a string. If this variable is not provided, Zelig will assume that the data is already ordered by time.
cs	Name of a variable that denotes the cross-sectional element of the data, for example, country name in a dataset with time-series across different countries. As a variable name, this should be in quotes. If this is not provided, Zelig will assume that all observations come from the same unit over time, and should be pooled, but if provided, individual models will be run in each cross-section. If <code>cs</code> is given as an argument, <code>ts</code> must also be provided. Additionally, <code>by</code> must be NULL.
order	A vector of length 3 passed in as <code>c(p, d, q)</code> where <code>p</code> represents the order of the autoregressive model, <code>d</code> represents the number of differences taken in the model, and <code>q</code> represents the order of the moving average model.

## Details

Currently only the Reference class syntax for time series. This model does not accept Bootstraps or weights.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_ma.html](http://docs.zeligproject.org/articles/zelig_ma.html)

**Examples**

```
data(seatshare)
subset <- seatshare[seatshare$country == "UNITED KINGDOM",]
ts.out <- zelig(formula = unemp ~ leftseat, model = "ma", ts = "year", data = subset)
summary(ts.out)
```

---

Zelig-mlogit-bayes-class

*Bayesian Multinomial Logistic Regression*

---

**Description**

Bayesian Multinomial Logistic Regression

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to zelig, relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

## Details

`zelig()` accepts the following arguments for `mlogit.bayes`:

- `baseline`: either a character string or numeric value (equal to one of the observed values in the dependent variable) specifying a baseline category. The default value is `NA` which sets the baseline to the first alphabetical or numerical unique value of the dependent variable.

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `burnin`: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- `mcmc`: number of the MCMC iterations after burnin (defaults to 10,000).
- `mcmc.method`: either "MH" or "slice", specifying whether to use Metropolis Algorithm or slice sampler. The default value is MH.
- `thin`: thinning interval for the Markov chain. Only every thin-th draw from the Markov chain is kept. The value of `mcmc` must be divisible by this value. The default value is 1.
- `tune`: tuning parameter for the Metropolis-Hasting step, either a scalar or a numeric vector (for `kk` coefficients, enter a `kk` vector). The tuning parameter should be set such that the acceptance rate is satisfactory (between 0.2 and 0.5). The default value is 1.1.
- `verbose`: defaults to `FALSE`. If `TRUE`, the progress of the sampler (every 10%) is printed to the screen.
- `seed`: seed for the random number generator. The default is `NA` which corresponds to a random seed of 12345.
- `beta.start`: starting values for the Markov chain, either a scalar or vector with length equal to the number of estimated coefficients. The default is `NA`, such that the maximum likelihood estimates are used as the starting values.

Use the following parameters to specify the model's priors:

- $b_0$ : prior mean for the coefficients, either a numeric vector or a scalar. If a scalar value, that value will be the prior mean for all the coefficients. The default is 0.
- $B_0$ : prior precision parameter for the coefficients, either a square matrix (with the dimensions equal to the number of the coefficients) or a scalar. If a scalar value, that value times an identity matrix will be the prior precision parameter. The default is 0, which leads to an improper prior.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

## See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_mlogitbayes.html](http://docs.zeligproject.org/articles/zelig_mlogitbayes.html)



**Examples**

```
data(mexico)
z.out <- zelig(vote88 ~ pristr + othcok + othsocok,model = "mlogit.bayes",
data = mexico,verbose = FALSE)
```

---

Zelig-negbin-class      *Negative Binomial Regression for Event Count Dependent Variables*

---

**Description**

Negative Binomial Regression for Event Count Dependent Variables

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to zelig, relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

**Details**

Additional parameters available to this model include:

- **weights**: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- **bootstrap**: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

**Value**

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**Methods**

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_negbin.html](http://docs.zeligproject.org/articles/zelig_negbin.html)

**Examples**

```
library(Zelig)
data(sanction)
z.out <- zelig(num ~ target + coop, model = "negbin", data = sanction)
summary(z.out)
```

---

Zelig-normal-bayes-class

*Bayesian Normal Linear Regression*

---

**Description**

Bayesian Normal Linear Regression

**Arguments**

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
<code>model</code>	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <code>Amelia</code> or <code>to_zelig_mi</code> ).

...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
<code>by</code>	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using <code>MatchIt</code> subclasses.
<code>cite</code>	If is set to 'TRUE' (default), the model citation will be printed to the console.

## Details

Additional parameters available to many models include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `burnin`: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- `mcmc`: number of the MCMC iterations after burnin (defaults to 10,000).
- `thin`: thinning interval for the Markov chain. Only every `thin`-th draw from the Markov chain is kept. The value of `mcmc` must be divisible by this value. The default value is 1.
- `verbose`: defaults to FALSE. If TRUE, the progress of the sampler (every 10%) is printed to the screen.
- `seed`: seed for the random number generator. The default is NA which corresponds to a random seed of 12345.
- `beta.start`: starting values for the Markov chain, either a scalar or vector with length equal to the number of estimated coefficients. The default is NA, such that the maximum likelihood estimates are used as the starting values.

Use the following parameters to specify the model's priors:

- `b0`: prior mean for the coefficients, either a numeric vector or a scalar. If a scalar value, that value will be the prior mean for all the coefficients. The default is 0.
- `B0`: prior precision parameter for the coefficients, either a square matrix (with the dimensions equal to the number of the coefficients) or a scalar. If a scalar value, that value times an identity matrix will be the prior precision parameter. The default is 0, which leads to an improper prior.
- `c0`: `c0/2` is the shape parameter for the Inverse Gamma prior on the variance of the disturbance terms.
- `d0`: `d0/2` is the scale parameter for the Inverse Gamma prior on the variance of the disturbance terms.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_normalbayes.html](http://docs.zeligproject.org/articles/zelig_normalbayes.html)

**Examples**

```
data(macro)
z.out <- zelig(unem ~ gdp + capmob + trade, model = "normal.bayes", data = macro, verbose = FALSE)

data(macro)
z.out <- zelig(unem ~ gdp + capmob + trade, model = "normal.bayes",
data = macro, verbose = FALSE)

z.out$geweke.diag()
z.out$heidel.diag()
z.out$raftery.diag()
summary(z.out)

x.out <- setx(z.out)
s.out1 <- sim(z.out, x = x.out)
summary(s.out1)

x.high <- setx(z.out, trade = quantile(macro$trade, prob = 0.8))
x.low <- setx(z.out, trade = quantile(macro$trade, prob = 0.2))

s.out2 <- sim(z.out, x = x.high, x1 = x.low)
summary(s.out2)
```

---

Zelig-normal-class      *Normal Regression for Continuous Dependent Variables*

---

**Description**

Normal Regression for Continuous Dependent Variables

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .

data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <i>Amelia</i> or <code>to_zelig_mi</code> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using <i>MatchIt</i> subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
below	(defaults to 0) The point at which the dependent variable is censored from below. If any values in the dependent variable are observed to be less than the censoring point, it is assumed that that particular observation is censored from below at the observed value. (See for a Bayesian implementation that supports both left and right censoring.)
robust	defaults to FALSE. If TRUE, <code>zelig()</code> computes robust standard errors based on sandwich estimators (see <code>and</code> ) and the options selected in <code>cluster</code> .
if	<code>robust = TRUE</code> , you may select a variable to define groups of correlated observations. Let <code>x3</code> be a variable that consists of either discrete numeric values, character strings, or factors that define strata. Then <code>z.out &lt;- zelig(y ~ x1 + x2, robust = TRUE, cluster = "x3", model = "tobit", data = mydata)</code> means that the observations can be correlated within the strata defined by the variable <code>x3</code> , and that robust standard errors should be calculated according to those clusters. If <code>robust = TRUE</code> but <code>cluster</code> is not specified, <code>zelig()</code> assumes that each observation falls into its own cluster.
formula	a model fitting formula

## Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_normal.html](http://docs.zeligproject.org/articles/zelig_normal.html)

**Examples**

```
data(macro)
z.out1 <- zelig(unem ~ gdp + capmob + trade, model = "normal",
data = macro)
summary(z.out1)
x.high <- setx(z.out1, trade = quantile(macro$trade, 0.8))
x.low <- setx(z.out1, trade = quantile(macro$trade, 0.2))
s.out1 <- sim(z.out1, x = x.high, x1 = x.low)
summary(s.out1)
plot(s.out1)
```

---

Zelig-normal-gee-class

*Generalized Estimating Equation for Normal Regression*

---

**Description**

Generalized Estimating Equation for Normal Regression

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <i>Amelia</i> or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using <i>MatchIt</i> subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

robust	defaults to TRUE. If TRUE, consistent standard errors are estimated using a "sandwich" estimator.
corstr	defaults to "independence". It can take on the following arguments:
Independence	(corstr = independence): $\text{cor}(y_{it}, y_{it'})=0$ , for all $t, t'$ with $t$ not equal to $t'$ . It assumes that there is no correlation within the clusters and the model becomes equivalent to standard normal regression. The "working" correlation matrix is the identity matrix.
Fixed	corstr = fixed): If selected, the user must define the "working" correlation matrix with the R argument rather than estimating it from the model.
id:	where id is a variable which identifies the clusters. The data should be sorted by id and should be ordered within each cluster when appropriate
corstr:	character string specifying the correlation structure: "independence", "exchangeable", "ar1", "unstructured" and "userdefined"
geeglm:	See geeglm in package geepack for other function arguments
Mv:	defaults to 1. It specifies the number of periods of correlation and only needs to be specified when corstr is stat_M_dep, non_stat_M_dep, or AR-M.
R:	defaults to NULL. It specifies a user-defined correlation matrix rather than estimating it from the data. The argument is used only when corstr is "fixed". The input is a TxT matrix of correlations, where T is the size of the largest cluster.

## Details

Additional parameters available to this model include:

- **weights:** vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.

## Value

Depending on the class of model selected, zelig will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

## See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_normalgee.html](http://docs.zeligproject.org/articles/zelig_normalgee.html)

## Examples

```
library(Zelig)
data(macro)
z.out <- zelig(unem ~ gdp + capmob + trade, model = "normal.gee", id = "country",
              data = macro, corstr = "AR-M")
summary(z.out)
```

---

Zelig-normal-survey-class

*Normal Regression for Continuous Dependent Variables with Survey Weights*


---

## Description

Normal Regression for Continuous Dependent Variables with Survey Weights

## Arguments

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

## Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.



**Value**

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_normalsurvey.html](http://docs.zeligproject.org/articles/zelig_normalsurvey.html)

**Examples**

```
library(Zelig)
data(api, package = "survey")
z.out1 <- zelig(api00 ~ meals + yr.rnd, model = "normal.survey", eights = ~pw, data = apistrat)
summary(z.out1)
```

---

Zelig-oprobit-bayes-class

*Bayesian Ordered Probit Regression*

---

**Description**

Bayesian Ordered Probit Regression

**Arguments**

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
<code>model</code>	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <code>Amelia</code> or <code>to_zelig_mi</code> ).
<code>...</code>	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
<code>by</code>	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using <code>MatchIt</code> subclasses.
<code>cite</code>	If is set to 'TRUE' (default), the model citation will be printed to the console.

## Details

Additional parameters available to many models include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `burnin`: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- `mcmc`: number of the MCMC iterations after burnin (defaults to 10,000).
- `thin`: thinning interval for the Markov chain. Only every thin-th draw from the Markov chain is kept. The value of `mcmc` must be divisible by this value. The default value is 1.
- `verbose`: defaults to FALSE. If TRUE, the progress of the sampler (every 10%) is printed to the screen.
- `seed`: seed for the random number generator. The default is NA which corresponds to a random seed of 12345.
- `beta.start`: starting values for the Markov chain, either a scalar or vector with length equal to the number of estimated coefficients. The default is NA, such that the maximum likelihood estimates are used as the starting values.

Use the following parameters to specify the model's priors:

- `b0`: prior mean for the coefficients, either a numeric vector or a scalar. If a scalar value, that value will be the prior mean for all the coefficients. The default is 0.
- `B0`: prior precision parameter for the coefficients, either a square matrix (with the dimensions equal to the number of the coefficients) or a scalar. If a scalar value, that value times an identity matrix will be the prior precision parameter. The default is 0, which leads to an improper prior.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

Vignette: [http://docs.zeligproject.org/articles/zelig\\_oprobitbayes.html](http://docs.zeligproject.org/articles/zelig_oprobitbayes.html)

---

Zelig-poisson-bayes-class

*Bayesian Poisson Regression*

---

## Description

Bayesian Poisson Regression

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to zelig, relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

**Details**

Additional parameters available to this model include:

- **weights**: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- **burnin**: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- **mcmc**: number of the MCMC iterations after burnin (defaults to 10,000).
- **tune**: Metropolis tuning parameter, either a positive scalar or a vector of length  $kk$ , where  $kk$  is the number of coefficients. The tuning parameter should be set such that the acceptance rate of the Metropolis algorithm is satisfactory (typically between 0.20 and 0.5). The default value is 1.1.
- **thin**: thinning interval for the Markov chain. Only every thin-th draw from the Markov chain is kept. The value of mcmc must be divisible by this value. The default value is 1.
- **verbose**: defaults to FALSE. If TRUE, the progress of the sampler (every 10%) is printed to the screen.
- **seed**: seed for the random number generator. The default is NA which corresponds to a random seed of 12345.
- **beta.start**: starting values for the Markov chain, either a scalar or vector with length equal to the number of estimated coefficients. The default is NA, such that the maximum likelihood estimates are used as the starting values.

Use the following parameters to specify the model's priors:

- $b_0$ : prior mean for the coefficients, either a numeric vector or a scalar. If a scalar value, that value will be the prior mean for all the coefficients. The default is 0.
- $B_0$ : prior precision parameter for the coefficients, either a square matrix (with the dimensions equal to the number of the coefficients) or a scalar. If a scalar value, that value times an identity matrix will be the prior precision parameter. The default is 0, which leads to an improper prior.

### Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_poissonbayes.html](http://docs.zeligproject.org/articles/zelig_poissonbayes.html)

### Examples

```
data(sanction)
z.out <- zelig(num ~ target + coop, model = "poisson.bayes", data = sanction, verbose = FALSE)
```

---

Zelig-poisson-class     *Poisson Regression for Event Count Dependent Variables*

---

### Description

Poisson Regression for Event Count Dependent Variables

### Arguments

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
<code>model</code>	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <code>Amelia</code> or <code>to_zelig_mi</code> ).
<code>...</code>	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.

by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
id:	where id is a variable which identifies the clusters. The data should be sorted by id and should be ordered within each cluster when appropriate
corstr:	character string specifying the correlation structure: "independence", "exchangeable", "ar1", "unstructured" and "userdefined"
geeglm:	See geeglm in package geepack for other function arguments

## Details

Additional parameters available to this model include:

- **weights**: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- **bootstrap**: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

## Value

Depending on the class of model selected, zelig will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zeig_model`.

## See Also

Vignette: [http://docs.zeligproject.org/articles/zeig\\_poisson.html](http://docs.zeligproject.org/articles/zeig_poisson.html)

## Examples

```
library(Zelig)
data(sanction)
z.out <- zelig(num ~ target + coop, model = "poisson", data = sanction)
summary(z.out)
```

---

 Zelig-poisson-gee-class

*Generalized Estimating Equation for Poisson Regression*


---

## Description

Generalized Estimating Equation for Poisson Regression

## Arguments

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <i>Amelia</i> or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <i>zelig</i> , relevant for the model to be estimated.
by	a factor variable contained in <i>data</i> . If supplied, <i>zelig</i> will subset the data frame based on the levels in the <i>by</i> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <i>by</i> to run models using <i>MatchIt</i> subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
id:	where <i>id</i> is a variable which identifies the clusters. The data should be sorted by <i>id</i> and should be ordered within each cluster when appropriate
corstr:	character string specifying the correlation structure: "independence", "exchangeable", "ar1", "unstructured" and "userdefined"
geeglm:	See <i>geeglm</i> in package <i>geepack</i> for other function arguments

## Details

Additional parameters available to this model include:

- **weights**: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- **bootstrap**: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

**Value**

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_poissongee.html](http://docs.zeligproject.org/articles/zelig_poissongee.html)

**Examples**

```
library(Zelig)
data(sanction)
sanction$cluster <- c(rep(c(1:15), 5), rep(c(16), 3))
sorted.sanction <- sanction[order(sanction$cluster),]
z.out <- zelig(num ~ target + coop, model = "poisson.gee", id = "cluster", data = sorted.sanction)
summary(z.out)
```

---

Zelig-poisson-survey-class

*Poisson Regression with Survey Weights*

---

**Description**

Poisson Regression with Survey Weights

**Arguments**

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
<code>model</code>	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <code>Amelia</code> or <code>to_zelig_mi</code> ).
<code>...</code>	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.

by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

### Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

### Value

Depending on the class of model selected, zelig will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_poissonsurvey.html](http://docs.zeligproject.org/articles/zelig_poissonsurvey.html)

### Examples

```
library(Zelig)
data(api, package="survey")
z.out1 <- zelig(enroll ~ api99 + yr.rnd, model = "poisson.survey", data = apistrat)
summary(z.out1)
```

---

Zelig-probit-bayes-class

*Bayesian Probit Regression*

---

### Description

Bayesian Probit Regression



**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. For example, to run the same model on all fifty states, you could use: <code>z.out &lt;- zelig(y ~ x1 + x2, data = mydata, model = 'ls', by = 's')</code> . You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

**Details**

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `burnin`: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- `mcmc`: number of the MCMC iterations after burnin (defaults to 10,000).
- `thin`: thinning interval for the Markov chain. Only every thin-th draw from the Markov chain is kept. The value of `mcmc` must be divisible by this value. The default value is 1.
- `verbose`: defaults to FALSE. If TRUE, the progress of the sampler (every 10%) is printed to the screen.
- `seed`: seed for the random number generator. The default is NA which corresponds to a random seed of 12345.
- `beta.start`: starting values for the Markov chain, either a scalar or vector with length equal to the number of estimated coefficients. The default is NA, such that the maximum likelihood estimates are used as the starting values.

Use the following parameters to specify the model's priors:

- `b0`: prior mean for the coefficients, either a numeric vector or a scalar. If a scalar value, that value will be the prior mean for all the coefficients. The default is 0.

- $B_0$ : prior precision parameter for the coefficients, either a square matrix (with the dimensions equal to the number of the coefficients) or a scalar. If a scalar value, that value times an identity matrix will be the prior precision parameter. The default is 0, which leads to an improper prior.

Use the following arguments to specify optional output for the model:

- `bayes.resid`: defaults to FALSE. If TRUE, the latent Bayesian residuals for all observations are returned. Alternatively, users can specify a vector of observations for which the latent residuals should be returned.

### Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_probitbayes.html](http://docs.zeligproject.org/articles/zelig_probitbayes.html)

### Examples

```
data(turnout)
z.out <- zelig(vote ~ race + educate, model = "probit.bayes", data = turnout, verbose = FALSE)
summary(z.out)
```

---

Zelig-probit-class      *Probit Regression for Dichotomous Dependent Variables*

---

### Description

Probit Regression for Dichotomous Dependent Variables

### Arguments

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The + symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .

data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <code>to_zelig_mi</code> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

### Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_probit.html](http://docs.zeligproject.org/articles/zelig_probit.html)

### Examples

```
data(turnout)
z.out <- zelig(vote ~ race + educate, model = "probit", data = turnout)
summary(z.out)
x.out <- setx(z.out)
s.out <- sim(z.out, x = x.out)
summary(s.out)
plot(s.out)
```

---

Zelig-probit-gee-class

*Generalized Estimating Equation for Probit Regression*

---

### Description

Generalized Estimating Equation for Probit Regression

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <code>to_zelig_mi</code> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
constr: character	string specifying the correlation structure: "independence", "exchangeable", "ar1", "unstructured" and "userdefined"
See	<code>geeglm</code> in package <code>geepack</code> for other function arguments.
id:	where <code>id</code> is a variable which identifies the clusters. The data should be sorted by <code>id</code> and should be ordered within each cluster when appropriate
constr:	character string specifying the correlation structure: "independence", "exchangeable", "ar1", "unstructured" and "userdefined"
geeglm:	See <code>geeglm</code> in package <code>geepack</code> for other function arguments

**Details**

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

**Value**

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_probitgee.html](http://docs.zeligproject.org/articles/zelig_probitgee.html)

**Examples**

```
data(turnout)
turnout$cluster <- rep(c(1:200), 10)
sorted.turnout <- turnout[order(turnout$cluster),]
z.out1 <- zelig(vote ~ race + educate, model = "probit.gee",
id = "cluster", data = sorted.turnout)
summary(z.out1)
```

---

Zelig-probit-survey-class

*Probit Regression with Survey Weights*

---

**Description**

@param formula a symbolic representation of the model to be estimated, in the form  $y \sim x_1 + x_2$ , where  $y$  is the dependent variable and  $x_1$  and  $x_2$  are the explanatory variables, and  $y$ ,  $x_1$ , and  $x_2$  are contained in the same dataset. (You may include more than two explanatory variables, of course.) The  $+$  symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form  $x_1*x_2$  without computing them in prior steps;  $I(x_1*x_2)$  to include only the interaction term and exclude the main effects; and quadratic terms in the form  $I(x_1^2)$ .

**Arguments**

<code>model</code>	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
<code>data</code>	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <code>Amelia</code> or <code>to_zelig_mi</code> ).
<code>...</code>	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
<code>by</code>	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using <code>MatchIt</code> subclasses.

<code>cite</code>	If is set to 'TRUE' (default), the model citation will be printed to the console.
<code>below:</code>	point at which the dependent variable is censored from below. If the dependent variable is only censored from above, set <code>below = -Inf</code> . The default value is 0.
<code>above:</code>	point at which the dependent variable is censored from above. If the dependent variable is only censored from below, set <code>above = Inf</code> . The default value is <code>Inf</code> .

## Details

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `burnin`: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- `mcmc`: number of the MCMC iterations after burnin (defaults to 10,000).
- `thin`: thinning interval for the Markov chain. Only every thin-th draw from the Markov chain is kept. The value of `mcmc` must be divisible by this value. The default value is 1.
- `verbose`: defaults to FALSE. If TRUE, the progress of the sampler (every 10%) is printed to the screen.
- `seed`: seed for the random number generator. The default is NA which corresponds to a random seed of 12345.
- `beta.start`: starting values for the Markov chain, either a scalar or vector with length equal to the number of estimated coefficients. The default is NA, such that the maximum likelihood estimates are used as the starting values.

Use the following parameters to specify the model's priors:

- `b0`: prior mean for the coefficients, either a numeric vector or a scalar. If a scalar value, that value will be the prior mean for all the coefficients. The default is 0.
- `B0`: prior precision parameter for the coefficients, either a square matrix (with the dimensions equal to the number of the coefficients) or a scalar. If a scalar value, that value times an identity matrix will be the prior precision parameter. The default is 0, which leads to an improper prior.
- `c0`:  $c0/2$  is the shape parameter for the Inverse Gamma prior on the variance of the disturbance terms.
- `d0`:  $d0/2$  is the scale parameter for the Inverse Gamma prior on the variance of the disturbance terms.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_probitsurvey.html](http://docs.zeligproject.org/articles/zelig_probitsurvey.html)

**Examples**

```
data(api, package="survey")
z.out1 <- zelig(enroll ~ api99 + yr.rnd ,
model = "poisson.survey", data = apistrat)
summary(z.out1)
x.low <- setx(z.out1, api99= quantile(apistrat$api99, 0.2))
x.high <- setx(z.out1, api99= quantile(apistrat$api99, 0.8))
s.out1 <- sim(z.out1, x=x.low, x1=x.high)
summary(s.out1)
plot(s.out1)
```

---

Zelig-quantile-class    *Quantile Regression for Continuous Dependent Variables*

---

**Description**

Quantile Regression for Continuous Dependent Variables

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to zelig, relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

## Details

In addition to the standard inputs, `zelig` takes the following additional options for quantile regression:

- `tau`: defaults to 0.5. Specifies the conditional quantile(s) that will be estimated. 0.5 corresponds to estimating the conditional median, 0.25 and 0.75 correspond to the conditional quartiles, etc. `tau` vectors with length greater than 1 are not currently supported. If `tau` is set outside of the interval [0,1], `zelig` returns the solution for all possible conditional quantiles given the data, but does not support inference on this fit (`setx` and `sim` will fail).
- `se`: a string value that defaults to "nid". Specifies the method by which the covariance matrix of coefficients is estimated during the `sim` stage of analysis. `se` can take the following values, which are passed to the `summary.rq` function from the `quantreg` package. These descriptions are copied from the `summary.rq` documentation.
  - "iid" which presumes that the errors are iid and computes an estimate of the asymptotic covariance matrix as in KB(1978).
  - "nid" which presumes local (in `tau`) linearity (in `x`) of the the conditional quantile functions and computes a Huber sandwich estimate using a local estimate of the sparsity.
  - "ker" which uses a kernel estimate of the sandwich as proposed by Powell(1990).
- `...`: additional options passed to `rq` when fitting the model. See documentation for `rq` in the `quantreg` package for more information.

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

## Methods

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

## See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_quantile.html](http://docs.zeligproject.org/articles/zelig_quantile.html)



**Examples**

```
library(Zelig)
data(stackloss)
z.out1 <- zelig(stack.loss ~ Air.Flow + Water.Temp + Acid.Conc.,
model = "rq", data = stackloss, tau = 0.5)
summary(z.out1)
```

---

Zelig-relogit-class     *Rare Events Logistic Regression for Dichotomous Dependent Variables*

---

**Description**

Rare Events Logistic Regression for Dichotomous Dependent Variables

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

**Details**

The `relogit` procedure supports four optional arguments in addition to the standard arguments for `zelig()`. You may additionally use:

- `tau`: a vector containing either one or two values for `tau`, the true population fraction of ones. Use, for example, `tau = c(0.05, 0.1)` to specify that the lower bound on `tau` is 0.05 and the upper bound is 0.1. If left unspecified, only finite-sample bias correction is performed, not case-control correction.

- `case.control`: if `tau` is specified, choose a method to correct for case-control sampling design: "prior" (default) or "weighting".
- `bias.correct`: a logical value of TRUE (default) or FALSE indicating whether the intercept should be corrected for finite sample (rare events) bias.

Additional parameters available to many models include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

## Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

## Methods

`modcall_formula_transformer()` Transform model call formula.

`show(signif.stars = FALSE, subset = NULL, bagging = FALSE)` Display a Zelig object

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

## See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_relogit.html](http://docs.zeligproject.org/articles/zelig_relogit.html)

## Examples

```
library(Zelig)
data(mid)
z.out1 <- zelig(conflict ~ major + contig + power + maxdem + mindem + years,
               data = mid, model = "relogit", tau = 1042/303772)
summary(z.out1)
```

---

Zelig-survey-class      *Survey models in Zelig for weights for complex sampling designs*

---

**Description**

Survey models in Zelig for weights for complex sampling designs

**Methods**

zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE) The zelig function estimates a variety of statistical models

---

Zelig-timeseries-class  
*Time-series models in Zelig*

---

**Description**

Time-series models in Zelig

**Methods**

packagename() Automatically retrieve wrapped package name

sim(num = NULL) Generic Method for Computing and Organizing Simulated Quantities of Interest

zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE) The zelig function estimates a variety of statistical models

---

Zelig-tobit-bayes-class  
*Bayesian Tobit Regression*

---

**Description**

Bayesian Tobit Regression

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by <i>Amelia</i> or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <i>zelig</i> , relevant for the model to be estimated.
by	a factor variable contained in <i>data</i> . If supplied, <i>zelig</i> will subset the data frame based on the levels in the <i>by</i> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <i>by</i> to run models using <i>MatchIt</i> subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
below:	point at which the dependent variable is censored from below. If the dependent variable is only censored from above, set <code>below = -Inf</code> . The default value is 0.
above:	point at which the dependent variable is censored from above. If the dependent variable is only censored from below, set <code>above = Inf</code> . The default value is <code>Inf</code> .
below:	point at which the dependent variable is censored from below. If the dependent variable is only censored from above, set <code>below = -Inf</code> . The default value is 0.
above:	point at which the dependent variable is censored from above. If the dependent variable is only censored from below, set <code>above = Inf</code> . The default value is <code>Inf</code> .

**Details**

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- `burnin`: number of the initial MCMC iterations to be discarded (defaults to 1,000).
- `mcmc`: number of the MCMC iterations after burnin (defaults to 10,000).
- `thin`: thinning interval for the Markov chain. Only every `thin`-th draw from the Markov chain is kept. The value of `mcmc` must be divisible by this value. The default value is 1.
- `verbose`: defaults to FALSE. If TRUE, the progress of the sampler (every 10%) is printed to the screen.
- `seed`: seed for the random number generator. The default is NA which corresponds to a random seed of 12345.

- `beta.start`: starting values for the Markov chain, either a scalar or vector with length equal to the number of estimated coefficients. The default is NA, such that the maximum likelihood estimates are used as the starting values.

Use the following parameters to specify the model's priors:

- `b0`: prior mean for the coefficients, either a numeric vector or a scalar. If a scalar value, that value will be the prior mean for all the coefficients. The default is 0.
- `B0`: prior precision parameter for the coefficients, either a square matrix (with the dimensions equal to the number of the coefficients) or a scalar. If a scalar value, that value times an identity matrix will be the prior precision parameter. The default is 0, which leads to an improper prior.
- `c0`:  $c0/2$  is the shape parameter for the Inverse Gamma prior on the variance of the disturbance terms.
- `d0`:  $d0/2$  is the scale parameter for the Inverse Gamma prior on the variance of the disturbance terms.

### Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_tobitbayes.html](http://docs.zeligproject.org/articles/zelig_tobitbayes.html)

### Examples

```
data(turnout)
z.out <- zelig(vote ~ race + educate, model = "tobit.bayes", data = turnout, verbose = FALSE)
```

### Description

Linear Regression for a Left-Censored Dependent Variable

**Arguments**

formula	a symbolic representation of the model to be estimated, in the form $y \sim x1 + x2$ , where $y$ is the dependent variable and $x1$ and $x2$ are the explanatory variables, and $y$ , $x1$ , and $x2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x1*x2$ without computing them in prior steps; $I(x1*x2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x1^2)$ .
model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to <code>zelig</code> , relevant for the model to be estimated.
by	a factor variable contained in <code>data</code> . If supplied, <code>zelig</code> will subset the data frame based on the levels in the <code>by</code> variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use <code>by</code> to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.
below	(defaults to 0) The point at which the dependent variable is censored from below. If any values in the dependent variable are observed to be less than the censoring point, it is assumed that that particular observation is censored from below at the observed value.
above	(defaults to 0) The point at which the dependent variable is censored from above. If any values in the dependent variable are observed to be more than the censoring point, it is assumed that that particular observation is censored from above at the observed value.
robust	defaults to FALSE. If TRUE, <code>zelig()</code> computes robust standard errors based on sandwich estimators and the options selected in <code>cluster</code> .
cluster	if <code>robust = TRUE</code> , you may select a variable to define groups of correlated observations. Let $x3$ be a variable that consists of either discrete numeric values, character strings, or factors that define strata. Then <code>z.out &lt;- zelig(y ~ x1 + x2, robust = TRUE, cluster = "x3", model = "tobit", data = mydata)</code> means that the observations can be correlated within the strata defined by the variable $x3$ , and that robust standard errors should be calculated according to those clusters. If <code>robust = TRUE</code> but <code>cluster</code> is not specified, <code>zelig()</code> assumes that each observation falls into its own cluster.

**Details**

Additional parameters available to this model include:

- `weights`: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.

- `bootstrap`: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

### Value

Depending on the class of model selected, `zelig` will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using `from_zelig_model`.

### Methods

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The `zelig` function estimates a variety of statistical models

### See Also

Vignette: [http://docs.zeligproject.org/articles/zelig\\_tobit.html](http://docs.zeligproject.org/articles/zelig_tobit.html)

### Examples

```
library(Zelig)
data(tobin)
z.out <- zelig(durable ~ age + quant, model = "tobit", data = tobin)
summary(z.out)
```

---

Zelig-weibull-class      *Weibull Regression for Duration Dependent Variables*

---

### Description

Weibull Regression for Duration Dependent Variables

### Arguments

<code>formula</code>	a symbolic representation of the model to be estimated, in the form $y \sim x_1 + x_2$ , where $y$ is the dependent variable and $x_1$ and $x_2$ are the explanatory variables, and $y$ , $x_1$ , and $x_2$ are contained in the same dataset. (You may include more than two explanatory variables, of course.) The $+$ symbol means “inclusion” not “addition.” You may also include interaction terms and main effects in the form $x_1*x_2$ without computing them in prior steps; $I(x_1*x_2)$ to include only the interaction term and exclude the main effects; and quadratic terms in the form $I(x_1^2)$ .
----------------------	---

model	the name of a statistical model to estimate. For a list of other supported models and their documentation see: <a href="http://docs.zeligproject.org/articles/">http://docs.zeligproject.org/articles/</a> .
data	the name of a data frame containing the variables referenced in the formula or a list of multiply imputed data frames each having the same variable names and row numbers (created by Amelia or <a href="#">to_zelig_mi</a> ).
...	additional arguments passed to zelig, relevant for the model to be estimated.
by	a factor variable contained in data. If supplied, zelig will subset the data frame based on the levels in the by variable, and estimate a model for each subset. This can save a considerable amount of effort. You may also use by to run models using MatchIt subclasses.
cite	If is set to 'TRUE' (default), the model citation will be printed to the console.

### Details

In addition to the standard inputs, zelig() takes the following additional options for weibull regression:

- **robust**: defaults to FALSE. If TRUE, zelig() computes robust standard errors based on sandwich estimators based on the options in cluster.
- **cluster**: if robust = TRUE, you may select a variable to define groups of correlated observations. Let x3 be a variable that consists of either discrete numeric values, character strings, or factors that define strata. Then `z.out <- zelig(y ~ x1 + x2, robust = TRUE, cluster = "x3"`, means that the observations can be correlated within the strata defined by the variable x3, and that robust standard errors should be calculated according to those clusters. If `robust=TRUE` but cluster is not specified, zelig() assumes that each observation falls into its own cluster.

Additional parameters available to this model include:

- **weights**: vector of weight values or a name of a variable in the dataset by which to weight the model. For more information see: <http://docs.zeligproject.org/articles/weights.html>.
- **bootstrap**: logical or numeric. If FALSE don't use bootstraps to robustly estimate uncertainty around model parameters due to sampling error. If an integer is supplied, the number of bootstraps to run. For more information see: <http://docs.zeligproject.org/articles/bootstraps.html>.

### Value

Depending on the class of model selected, zelig will return an object with elements including coefficients, residuals, and formula which may be summarized using `summary(z.out)` or individually extracted using, for example, `coef(z.out)`. See <http://docs.zeligproject.org/articles/getters.html> for a list of functions to extract model components. You can also extract whole fitted model objects using [from\\_zelig\\_model](#).

### Methods

`zelig(formula, data, model = NULL, ..., weights = NULL, by, bootstrap = FALSE)` The zelig function estimates a variety of statistical models



**See Also**

Vignette: [http://docs.zeligproject.org/articles/zelig\\_weibull.html](http://docs.zeligproject.org/articles/zelig_weibull.html)

**Examples**

```
data(coalition)
z.out <- zelig(Surv(duration, ciep12) ~ fract + numst2, model = "weibull", data = coalition)
```

---

Zelig.url	<i>Table of links for Zelig</i>
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---

**Description**

Table of links for `help.zelig` for the core Zelig package.

---

<code>zelig_qi_to_df</code>	<i>Extract simulated quantities of interest from a zelig object</i>
-----------------------------	---

---

**Description**

Extract simulated quantities of interest from a zelig object

**Usage**

```
zelig_qi_to_df(obj)
```

**Arguments**

`obj` a zelig object with simulated quantities of interest

**Details**

A simulated quantities of interest in a tidy data formatted `data.frame`. This can be useful for creating custom plots.

Each row contains a simulated value and each column contains:

- `setx_value` whether the simulations are from the base `x` `setx` or the contrasting `x1` for finding first differences.
- The fitted values specified in `setx` including a `by` column if `by` was used in the `zelig` call.
- `expected_value`
- `predicted_value`

For multinomial reponse models, a separate column is given for the expected probability of each outcome in the form `expected_*`. Additionally, there is a column of the predicted outcomes (`predicted_value`).

**Author(s)**

Christopher Gandrud

**Source**

For a discussion of tidy data see <https://www.jstatsoft.org/article/view/v059i10>.

**See Also**

[qi\\_slimmer](#)

**Examples**

```
##### QIs without first difference or range, from covariates fitted at
## central tendencies
z.1 <- zelig(Petal.Width ~ Petal.Length + Species, data = iris,
             model = "ls")
z.1 <- setx(z.1)
z.1 <- sim(z.1)
head(zelig_qi_to_df(z.1))

##### QIs for first differences
z.2 <- zelig(Petal.Width ~ Petal.Length + Species, data = iris,
             model = "ls")
z.2a <- setx(z.2, Petal.Length = 2)
z.2b <- setx(z.2, Petal.Length = 4.4)
z.2 <- sim(z.2, x = z.2a, x1 = z.2a)
head(zelig_qi_to_df(z.2))

##### QIs for first differences, estimated by Species
z.3 <- zelig(Petal.Width ~ Petal.Length, by = "Species", data = iris,
             model = "ls")
z.3a <- setx(z.3, Petal.Length = 2)
z.3b <- setx(z.3, Petal.Length = 4.4)
z.3 <- sim(z.3, x = z.3a, x1 = z.3a)
head(zelig_qi_to_df(z.3))

##### QIs for a range of fitted values
z.4 <- zelig(Petal.Width ~ Petal.Length + Species, data = iris,
             model = "ls")
z.4 <- setx(z.4, Petal.Length = 2:4)
z.4 <- sim(z.4)
head(zelig_qi_to_df(z.4))

##### QIs for a range of fitted values, estimated by Species
z.5 <- zelig(Petal.Width ~ Petal.Length, by = "Species", data = iris,
             model = "ls")
z.5 <- setx(z.5, Petal.Length = 2:4)
z.5 <- sim(z.5)
head(zelig_qi_to_df(z.5))

##### QIs for two ranges of fitted values
```

```

z.6 <- zelig(Petal.Width ~ Petal.Length + Species, data = iris,
             model = "ls")
z.6a <- setx(z.6, Petal.Length = 2:4, Species = "setosa")
z.6b <- setx(z.6, Petal.Length = 2:4, Species = "virginica")
z.6 <- sim(z.6, x = z.6a, x1 = z.6b)

head(zelig_qi_to_df(z.6))

```

---

zelig\_setx\_to\_df      *Extracted fitted values from a Zelig object with setx values*

---

### Description

Extracted fitted values from a Zelig object with setx values

### Usage

```
zelig_setx_to_df(obj)
```

### Arguments

obj                    a zelig object with simulated quantities of interest

### Details

Fitted (setx) values in a tidy data formatted data.frame. This was designed to enable the WhatIf package's whatif function to extract "counterfactuals".

### Author(s)

Christopher Gandrud

### Examples

```

#### QIs without first difference or range, from covariates fitted at
## central tendencies
z.1 <- zelig(Petal.Width ~ Petal.Length + Species, data = iris,
             model = "ls")
z.1 <- setx(z.1)
zelig_setx_to_df(z.1)

#### QIs for first differences
z.2 <- zelig(Petal.Width ~ Petal.Length + Species, data = iris,
             model = "ls")
z.2 <- setx(z.2, Petal.Length = 2)
z.2 <- setx1(z.2, Petal.Length = 4.4)
zelig_setx_to_df(z.2)

#### QIs for first differences, estimated by Species

```

```
z.3 <- zelig(Petal.Width ~ Petal.Length, by = "Species", data = iris,
            model = "ls")
z.3 <- setx(z.3, Petal.Length = 2)
z.3 <- setx1(z.3, Petal.Length = 4.4)
zelig_setx_to_df(z.3)

#### QIs for a range of fitted values
z.4 <- zelig(Petal.Width ~ Petal.Length + Species, data = iris,
            model = "ls")
z.4 <- setx(z.4, Petal.Length = 2:4)
zelig_setx_to_df(z.4)

#### QIs for a range of fitted values, estimated by Species
z.5 <- zelig(Petal.Width ~ Petal.Length, by = "Species", data = iris,
            model = "ls")
z.5 <- setx(z.5, Petal.Length = 2:4)
zelig_setx_to_df(z.5)

#### QIs for two ranges of fitted values
z.6 <- zelig(Petal.Width ~ Petal.Length + Species, data = iris,
            model = "ls")
z.6 <- setx(z.6, Petal.Length = 2:4, Species = "setosa")
z.6 <- setx1(z.6, Petal.Length = 2:4, Species = "virginica")
zelig_setx_to_df(z.6)
```

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