# Package 'prediction'

June 11, 2024

Type Package **Title** Tidy, Type-Safe 'prediction()' Methods **Description** A one-function package containing prediction(), a type-safe alternative to predict() that always returns a data frame. The summary() method provides a data frame with average predictions, possibly over counterfactual versions of the data (à la the margins command in 'Stata'). Marginal effect estimation is provided by the related package, 'margins' <https://cran.r-project.org/package=margins>. The package currently supports common model types (e.g., lm, glm) from the 'stats' package, as well as numerous other model classes from other add-on packages. See the README file or main package documentation page for a complete listing. License MIT + file LICENSE **Version** 0.3.18 URL https://github.com/bbolker/prediction BugReports https://github.com/bbolker/prediction/issues **Depends** R (>= 3.5.0) Imports utils, stats, data.table Suggests datasets, methods, testthat Enhances AER, and, betareg, biglm, brglm, caret, crch, e1071, earth, ff, gam (>= 1.15), gee, glmnet, glmx, kernlab, lme4, MASS, mclogit, mda, mlogit, MNP, nlme, nnet, ordinal, plm, pscl, quantreg, rpart, sampleSelection, speedglm, survey (>= 3.31-5), survival, truncreg, VGAM ByteCompile true **Encoding UTF-8** RoxygenNote 7.3.1 **NeedsCompilation** no **Author** Thomas J. Leeper [aut] (<a href="https://orcid.org/0000-0003-4097-6326">https://orcid.org/0000-0003-4097-6326</a>), Carl Ganz [ctb], Vincent Arel-Bundock [ctb] (<a href="https://orcid.org/0000-0003-2042-7063">https://orcid.org/0000-0003-2042-7063</a>), Ben Bolker [ctb, cre] (<a href="https://orcid.org/0000-0002-2127-0443">https://orcid.org/0000-0002-2127-0443</a>)

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prediction-package

Extract Predictions from a Model Object

# Description

Extract predicted values via predict from a model object, conditional on data, and return a data frame.

#### Usage

```
prediction(model, ...)
## Default S3 method:
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = "response",
  vcov = stats::vcov(model),
  calculate_se = TRUE,
)
## S3 method for class 'Arima'
prediction(model, calculate_se = TRUE, ...)
## S3 method for class 'ar'
prediction(model, data, at = NULL, calculate_se = TRUE, ...)
## S3 method for class 'arima0'
prediction(model, data, at = NULL, calculate_se = TRUE, ...)
```

```
## S3 method for class 'betareg'
prediction(
 model,
  data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link", "precision", "variance", "quantile"),
  calculate_se = FALSE,
)
## S3 method for class 'bigLm'
prediction(model, data = NULL, calculate_se = FALSE, ...)
## S3 method for class 'bigglm'
prediction(
 model,
  data = find_data(model, parent.frame()),
  at = NULL,
  type = "response",
 calculate_se = TRUE,
)
## S3 method for class 'biglm'
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
  type = "response",
  calculate_se = TRUE,
)
## S3 method for class 'bruto'
prediction(
 model,
 data = NULL,
  at = NULL,
  type = "fitted",
  calculate_se = FALSE,
## S3 method for class 'clm'
prediction(
 model,
 data = find_data(model, parent.frame()),
```

```
at = NULL,
  type = NULL,
  calculate_se = TRUE,
  category,
)
## S3 method for class 'coxph'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = c("risk", "expected", "lp"),
  calculate_se = TRUE,
)
## S3 method for class 'crch'
prediction(
 model,
 data = find_data(model),
  at = NULL,
  type = c("response", "location", "scale", "quantile"),
  calculate_se = FALSE,
)
## S3 method for class 'earth'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link"),
  calculate_se = TRUE,
  category,
)
## S3 method for class 'fda'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = NULL,
  calculate_se = FALSE,
  category,
)
```

```
## S3 method for class 'Gam'
prediction(
 model,
  data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link", "terms"),
  calculate_se = TRUE,
)
## S3 method for class 'gausspr'
prediction(
 model,
 data,
  at = NULL,
  type = NULL,
  calculate_se = TRUE,
  category,
)
## S3 method for class 'gee'
prediction(model, calculate_se = FALSE, ...)
## S3 method for class 'glimML'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link"),
  calculate_se = TRUE,
)
## S3 method for class 'glimQL'
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
  type = c("response", "link"),
  calculate_se = TRUE,
)
## S3 method for class 'glm'
prediction(
 model,
```

```
data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link"),
  vcov = stats::vcov(model),
  calculate_se = TRUE,
)
## S3 method for class 'glmnet'
prediction(
 model,
  data,
  lambda = model[["lambda"]][1L],
  at = NULL,
  type = c("response", "link"),
  calculate_se = FALSE,
)
## S3 method for class 'glmx'
prediction(
 model,
  data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link"),
  calculate_se = FALSE,
)
## S3 method for class 'gls'
prediction(
 model,
 data = find_data(model),
  at = NULL,
  calculate_se = FALSE,
)
## S3 method for class 'hetglm'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link", "scale"),
  calculate_se = FALSE,
)
```

```
## S3 method for class 'hurdle'
prediction(
 model,
  data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "count", "prob", "zero"),
  calculate_se = FALSE,
)
## S3 method for class 'hxlr'
prediction(
 model,
 data = find_data(model),
  at = NULL,
  type = c("class", "probability", "cumprob", "location", "scale"),
  calculate_se = FALSE,
)
## S3 method for class 'ivreg'
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
 calculate_se = FALSE,
)
## S3 method for class 'knnreg'
prediction(model, data, at = NULL, calculate_se = FALSE, ...)
## S3 method for class 'kqr'
prediction(model, data, at = NULL, calculate_se = FALSE, ...)
## S3 method for class 'ksvm'
prediction(
 model,
 data,
 at = NULL,
  type = NULL,
  calculate_se = TRUE,
  category,
)
## S3 method for class 'lda'
prediction(
```

```
model,
 data = find_data(model, parent.frame()),
  at = NULL,
  calculate_se = FALSE,
 category,
)
## S3 method for class 'lm'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = "response",
  vcov = stats::vcov(model),
  calculate_se = TRUE,
)
## S3 method for class 'lme'
prediction(
 model,
 data = find_data(model),
 at = NULL,
 calculate_se = FALSE,
)
## S3 method for class 'loess'
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
  type = "response",
  calculate_se = TRUE,
)
## S3 method for class 'lqs'
prediction(
 model,
 data = find_data(model),
 at = NULL,
 calculate_se = FALSE,
)
## S3 method for class 'mars'
```

```
prediction(
 model,
 data = NULL,
 at = NULL,
  type = "fitted",
  calculate_se = FALSE,
)
## S3 method for class 'mca'
prediction(
 model,
 data = find_data(model),
 at = NULL,
 calculate_se = FALSE,
)
## S3 method for class 'mclogit'
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
  type = "response",
 vcov = stats::vcov(model),
 calculate_se = TRUE,
)
## S3 method for class 'merMod'
prediction(
 model,
  data = find_data(model),
  at = NULL,
  type = c("response", "link"),
  re.form = NULL,
  calculate_se = FALSE,
)
## S3 method for class 'mlogit'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  calculate_se = FALSE,
  category,
  . . .
```

```
)
## S3 method for class 'mnlogit'
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
 calculate_se = FALSE,
 category,
)
## S3 method for class 'mnp'
prediction(
 model,
  data = find_data(model, parent.frame()),
  at = NULL,
  type = NULL,
  calculate_se = FALSE,
  category,
)
## S3 method for class 'multinom'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = NULL,
  calculate_se = FALSE,
  category,
)
## S3 method for class 'naiveBayes'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
 type = NULL,
 calculate_se = FALSE,
  category,
)
## S3 method for class 'nls'
prediction(
 model,
```

```
data = find_data(model, parent.frame()),
 at = NULL,
  calculate_se = FALSE,
)
## S3 method for class 'nnet'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = NULL,
  calculate_se = FALSE,
 category,
)
## S3 method for class 'plm'
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
 calculate_se = FALSE,
)
## S3 method for class 'polr'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = NULL,
  calculate_se = FALSE,
  category,
  . . .
)
## S3 method for class 'polyreg'
prediction(
 model,
 data = NULL,
 at = NULL,
  type = "fitted",
  calculate_se = FALSE,
)
## S3 method for class 'ppr'
```

```
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
 calculate_se = FALSE,
)
## S3 method for class 'princomp'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
 calculate_se = FALSE,
)
## S3 method for class 'qda'
prediction(
 model,
 data = find_data(model, parent.frame()),
 at = NULL,
 calculate_se = FALSE,
 category,
)
## S3 method for class 'rlm'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = "response",
  vcov = stats::vcov(model),
  calculate_se = TRUE,
)
## S3 method for class 'rpart'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = NULL,
  calculate_se = FALSE,
  category,
)
```

```
## S3 method for class 'rq'
prediction(
 model,
  data = find_data(model, parent.frame()),
 at = NULL,
 calculate_se = TRUE,
)
## S3 method for class 'selection'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = "response",
  calculate_se = FALSE,
)
## S3 method for class 'speedglm'
prediction(
 model,
  data = find_data(model, parent.frame()),
 at = NULL,
  type = c("response", "link"),
  calculate_se = FALSE,
)
## S3 method for class 'speedlm'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
 calculate_se = FALSE,
)
## S3 method for class 'survreg'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "lp", "quantile", "uquantile"),
  calculate_se = TRUE,
)
```

```
## S3 method for class 'svm'
prediction(model, data = NULL, at = NULL, calculate_se = TRUE, category, ...)
## S3 method for class 'svyglm'
prediction(
 model,
  data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link"),
  calculate_se = TRUE,
)
## S3 method for class 'train'
prediction(
 model,
  data = find_data(model),
 at = NULL,
  type = c("raw", "prob"),
)
## S3 method for class 'tree'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = NULL,
  calculate_se = FALSE,
  category,
)
## S3 method for class 'truncreg'
prediction(model, data, at = NULL, calculate_se = FALSE, ...)
## S3 method for class 'vgam'
prediction(
 model,
  data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link"),
  calculate_se = FALSE,
 category,
)
```

```
## S3 method for class 'vglm'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "link"),
  calculate_se = TRUE,
  category,
)
## S3 method for class 'zeroinfl'
prediction(
 model,
 data = find_data(model, parent.frame()),
  at = NULL,
  type = c("response", "count", "prob", "zero"),
  calculate_se = FALSE,
)
prediction_summary(model, ..., level = 0.95)
```

# Arguments

model	A model object, perhaps returned by 1m or g1m.	
	Additional arguments passed to predict methods.	
data	A data frame over which to calculate marginal effects. If missing, find_data is used to specify the data frame.	
at	A list of one or more named vectors, specifically values at which to calculate the predictions. These are used to modify the value of data (see build_datalist for details on use).	
type	A character string indicating the type of marginal effects to estimate. Mostly relevant for non-linear models, where the reasonable options are "response" (the default) or "link" (i.e., on the scale of the linear predictor in a GLM). For models of class "polr" (from polr), possible values are "class" or "probs"; both are returned.	
vcov	A matrix containing the variance-covariance matrix for estimated model coefficients, or a function to perform the estimation with model as its only argument.	
calculate_se	A logical indicating whether to calculate standard errors for observation-specific predictions and average predictions (if possible). The output will always contain a "calculate_se" column regardless of this value; this only controls the calculation of standard errors. Setting it to FALSE may improve speed.	
category	For multi-level or multi-category outcome models (e.g., ordered probit, multinomial logit, etc.), a value specifying which of the outcome levels should be used	

for the "fitted" column. If missing, some default is chosen automatically.

lambda For models of class "glmnet", a value of the penalty parameter at which predic-

tions are required.

re.form An argument passed forward to predict.merMod.

level A numeric value specifying the confidence level for calculating p-values and

confidence intervals.

#### **Details**

This function is simply a wrapper around predict that returns a data frame containing the value of data and the predicted values with respect to all variables specified in data.

Methods are currently implemented for the following object classes:

- "lm", see lm
- "glm", see glm, glm.nb, glmx, hetglm, brglm
- "ar", see ar
- "Arima", see arima
- "arima0", see arima0
- "bigglm", see bigglm
- "betareg", see betareg
- "bruto", see bruto
- "clm", see clm
- "coxph", see coxph
- "crch", see crch
- "earth", see earth
- "fda", see fda
- "Gam", see gam
- "gausspr", see gausspr
- "gee", see gee
- "glmnet", see glmnet
- "gls", see gls
- "glimML", see betabin, negbin
- "glimQL", see quasibin, quasipois
- "hurdle", see hurdle
- "hxlr", see hxlr
- "ivreg", see ivreg
- "knnreg", see knnreg
- "kqr", see kqr
- "ksvm", see ksvm
- "lda", see lda
- "lme", see 1me

- "loess", see loess
- "lqs", see lqs
- "mars", see mars
- "mca", see mca
- "mclogit", see mclogit
- "mda", see mda
- "merMod", see lmer, glmer
- "mnp", see mnp
- "naiveBayes", see naiveBayes
- "nlme", see nlme
- "nls", see nls
- "nnet", see nnet
- "plm", see plm
- "polr", see polr
- "polyreg", see polyreg
- "ppr", see ppr
- "princomp", see princomp
- "qda", see qda
- "rlm", see rlm
- "rpart", see rpart
- "rq", see rq
- "selection", see selection
- "speedglm", see speedglm
- "speedlm", see speedlm
- "survreg", see survreg
- "svm", see svm
- "svyglm", see svyglm
- "tobit", see tobit
- "train", see train
- "truncreg", see truncreg
- "zeroinfl", see zeroinfl

Where implemented, prediction also returns average predictions (and the variances thereof). Variances are implemented using the delta method, as described by Xu and Long 2005 doi:10.1177/1536867X0500500405.

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#### Value

A data frame with class "prediction" that has a number of rows equal to number of rows in data, or a multiple thereof, if !is.null(at). The return value contains data (possibly modified by at using build\_datalist), plus a column containing fitted/predicted values ("fitted") and a column containing the standard errors thereof ("calculate\_se"). Additional columns may be reported depending on the object class. The data frame also carries attributes used by print and summary, which will be lost during subsetting.

#### See Also

```
find_data, build_datalist, mean_or_mode, seq_range
```

# **Examples**

```
require("datasets")
x <- lm(Petal.Width ~ Sepal.Length * Sepal.Width * Species, data = iris)
# prediction for every case
prediction(x)
# prediction for first case
prediction(x, iris[1,])
# basic use of 'at' argument
summary(prediction(x, at = list(Species = c("setosa", "virginica"))))
# basic use of 'at' argument
prediction(x, at = list(Sepal.Length = seq_range(iris$Sepal.Length, 5)))
# prediction at means/modes of input variables
prediction(x, at = lapply(iris, mean_or_mode))
# prediction with multi-category outcome
## Not run:
 library("mlogit")
 data("Fishing", package = "mlogit")
 Fish <- mlogit.data(Fishing, varying = c(2:9), shape = "wide", choice = "mode")
 mod <- mlogit(mode ~ price + catch, data = Fish)</pre>
 prediction(mod)
 prediction(mod, category = 3)
## End(Not run)
```

build\_datalist

Build list of data.frames

# Description

Construct a list of data frames based upon an input data frame and a list of one or more at values

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#### Usage

```
build_datalist(data, at = NULL, as.data.frame = FALSE, ...)
```

#### **Arguments**

data A data.frame containing the original data.

A list of one or more named vectors of values, which will be used to specify val-

ues of variables in data. All possible combinations are generated. Alternatively, this can be a data frame of combination levels if only a subset of combinations

are desired. See examples.

as.data.frame A logical indicating whether to return a single stacked data frame rather than a

list of data frames

... Ignored.

#### Value

A list of data.frames, unless as.data.frame = TRUE in which case a single, stacked data frame is returned.

#### Author(s)

Thomas J. Leeper

#### See Also

```
find_data, mean_or_mode, seq_range
```

#### **Examples**

```
# basic examples
require("datasets")
build_datalist(head(mtcars), at = list(cyl = c(4, 6)))
str(build_datalist(head(mtcars), at = list(cyl = c(4,6), wt = c(2.75,3,3.25))), 1)
str(build_datalist(head(mtcars), at = data.frame(cyl = c(4,4), wt = c(2.75,3))))
```

find\_data

Extract data from a model object

### Description

Attempt to reconstruct the data used to create a model object

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#### Usage

```
find_data(model, ...)
## Default S3 method:
find_data(model, env = parent.frame(), ...)
## S3 method for class 'data.frame'
find_data(model, ...)
## S3 method for class 'crch'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'glimML'
find_data(model, ...)
## S3 method for class 'glimQL'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'glm'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'hxlr'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'lm'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'mca'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'merMod'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'svyglm'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'train'
find_data(model, ...)
## S3 method for class 'vgam'
find_data(model, env = parent.frame(), ...)
## S3 method for class 'vglm'
find_data(model, env = parent.frame(), ...)
```

### **Arguments**

model The model object.

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... Additional arguments passed to methods.

env An environment in which to look for the data argument to the modelling call.

#### **Details**

This is a convenience function and, as such, carries no guarantees. To behave well, it typically requires that a model object be specified using a formula interface and an explicit data argument. Models that can be specified using variables from the .GlobalEnv or with a non-formula interface (e.g., a matrix of data) will tend to generate errors. find\_data is an S3 generic so it is possible to expand it with new methods.

#### Value

A data frame containing the original data used in a modelling call, modified according to the original model's 'subset' and 'na.action' arguments, if appropriate.

#### See Also

```
prediction, build_datalist, mean_or_mode, seq_range
```

#### **Examples**

```
require("datasets")
x <- lm(mpg ~ cyl * hp + wt, data = head(mtcars))
find_data(x)</pre>
```

margex

Artificial data for margins, copied from Stata

#### **Description**

The dataset is identical to the one provided by Stata and available from webuse::webuse("margex") with categorical variables explicitly encoded as factors.

#### Usage

```
data("margex")
```

#### **Format**

A data frame with 3000 observations on the following 11 variables.

```
'y' A numeric vector
```

'outcome' A binary numeric vector with values (0,1)

'sex' A factor with two levels

'group' A factor with three levels

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```
'age' A numeric vector
'distance' A numeric vector
'ycn' A numeric vector
'yc' A numeric vector
'treatment' A factor with two levels
'agegroup' A factor with three levels
'arm' A factor with three levels
```

#### Source

```
https://www.stata-press.com/data/r14/margex.dta
```

#### See Also

prediction

#### **Examples**

```
# Examples from Stata's help files
# Also available from: webuse::webuse("margex")
data("margex")
# A simple case after regress
# . regress y i.sex i.group
# . margins sex
m1 <- lm(y ~ factor(sex) + factor(group), data = margex)
prediction(m1, at = list(sex = c("male", "female")))
# A simple case after logistic
# . logistic outcome i.sex i.group
# . margins sex
m2 <- glm(outcome ~ sex + group, binomial(), data = margex)</pre>
prediction(m2, at = list(sex = c("male", "female")))
# Average response versus response at average
# . margins sex
prediction(m2, at = list(sex = c("male", "female")))
# . margins sex, atmeans
## TODO
# Multiple margins from one margins command
# . margins sex group
prediction(m2, at = list(sex = c("male", "female")))
prediction(m2, at = list(group = c("1", "2", "3")))
# Margins with interaction terms
# . logistic outcome i.sex i.group sex#group
# . margins sex group
m3 <- glm(outcome ~ sex * group, binomial(), data = margex)</pre>
```

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```
prediction(m3, at = list(sex = c("male", "female")))
prediction(m3, at = list(group = c("1", "2", "3")))
# Margins with continuous variables
# . logistic outcome i.sex i.group sex#group age
# . margins sex group
m4 <- glm(outcome ~ sex * group + age, binomial(), data = margex)</pre>
prediction(m4, at = list(sex = c("male", "female")))
prediction(m4, at = list(group = c("1", "2", "3")))
# Margins of continuous variables
# . margins, at(age=40)
prediction(m4, at = list(age = 40))
# . margins, at(age=(30 35 40 45 50))
prediction(m4, at = list(age = c(30, 35, 40, 45, 50)))
# Margins of interactions
# . margins sex#group
prediction(m4, at = list(sex = c("male", "female"), group = c("1", "2", "3")))
```

mean\_or\_mode

Class-dependent variable aggregation

### Description

Summarize a vector/variable into a single number, either a mean (median) for numeric vectors or the mode for categorical (character, factor, ordered, or logical) vectors. Useful for aggregation.

#### Usage

```
mean_or_mode(x)

## Default S3 method:
mean_or_mode(x)

## S3 method for class 'numeric'
mean_or_mode(x)

## S3 method for class 'data.frame'
mean_or_mode(x)

median_or_mode(x)

## Default S3 method:
median_or_mode(x)

## S3 method for class 'numeric'
```

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```
median_or_mode(x)
## S3 method for class 'data.frame'
median_or_mode(x)
```

#### **Arguments**

Χ

A vector.

#### Value

A numeric or factor vector of length 1.

#### See Also

```
prediction, build_datalist, seq_range
```

# **Examples**

```
require("datasets")
# mean for numerics
mean_or_mode(iris)
mean_or_mode(iris[["Sepal.Length"]])
mean_or_mode(iris[["Species"]])
# median for numerics
median_or_mode(iris)
```

seq\_range

Create a sequence over the range of a vector

# Description

Define a sequence of evenly spaced values from the minimum to the maximum of a vector

## Usage

```
seq_range(x, n = 2)
```

#### **Arguments**

x A numeric vector

n An integer specifying the length of sequence (i.e., number of points across the range of x)

### Value

A vector of length n.

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# See Also

```
mean_or_mode, build_datalist
```

# Examples

```
identical(range(1:5), seq_range(1:5, n = 2)) seq_range(1:5, n = 3)
```

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