

# Package ‘rmdcev’

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**Type** Package

**Title** Kuhn-Tucker and Multiple Discrete-Continuous Extreme Value Models

**Version** 1.2.4

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## Description

Estimates and simulates Kuhn-Tucker demand models with individual heterogeneity. The package implements the multiple-discrete continuous extreme value (MDCEV) model and the Kuhn-Tucker specification common in the environmental economics literature on recreation demand. Latent class and random parameters specifications can be implemented and the models are fit using maximum likelihood estimation or Bayesian estimation. All models are implemented in Stan, which is a C++ package for performing full Bayesian inference (see Stan Development Team, 2019) <<https://mc-stan.org/>>. The package also implements demand forecasting (Pinjari and Bhat (2011) <<https://repositories.lib.utexas.edu/handle/2152/23880>>) and welfare calculation (Lloyd-Smith (2018) <[doi:10.1016/j.jocm.2017.12.002](https://doi.org/10.1016/j.jocm.2017.12.002)>) for policy simulation.

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**Depends** R (>= 4.0.0), Rcpp (>= 1.0.5), methods

**Imports** rstan (>= 2.21.0), rstantools (>= 2.1.1), RcppParallel (>= 5.0.1), dplyr (>= 0.7.8), purrr, tibble, tidyr, utils, stats, Formula

**LinkingTo** BH (>= 1.72.0), Rcpp, RcppEigen (>= 0.3.3.3.0), RcppParallel (>= 5.0.1), rstan (>= 2.21.0), StanHeaders (>= 2.21.0)

**Encoding** UTF-8

**LazyData** true

**Repository** CRAN

**SystemRequirements** GNU make

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**Biarch** true

**Suggests** knitr, rmarkdown, testthat

**URL** <https://github.com/plloydsmith/rmdcev>

**BugReports** <https://github.com/plloydsmith/rmdcev/issues>

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CreateBlankPolicies    *CreateBlankPolicies*

---

### Description

Create 'zero effect' policies that can be modified

### Usage

```
CreateBlankPolicies(npols, model, price_change_only = TRUE)
```

### Arguments

npols            Number of policies to simulate

model            Estimated model from mdcev

price\_change\_only    Logical value for whether to include policy changes to dat\_psi. Defaults to TRUE. TRUE implies that only price changes are used in simulation.

### Examples

```
data_rec <- mdcev.data(data_rec, subset = id <= 500, id.var = "id",
  alt.var = "alt", choice = "quant")
```

```
mdcev_est <- mdcev(~ 0, data = data_rec,
  model = "hybrid0", algorithm = "MLE",
  std_errors = "mvn")
```

```
CreateBlankPolicies(npols = 2, mdcev_est)
```

---

data\_rec

*Recreation data from Value of Nature to Canadians Survey*

---

### Description

Data from 2000 individuals from the Value of Nature to Canadians (VNC) survey. The travel costs are calculated using the approach described in Lloyd-Smith (2020)

### Usage

```
data(data_rec)
```

### Format

A tibble with 34000 rows and 8 variables

### Source

[Canadian Nature Survey 2012](#)

### References

Federal, Provincial, and Territorial Governments of Canada. 2014. "2012 Canadian Nature Survey: Awareness, Participation, and Expenditures in Nature-Based Recreation, Conservation, and Subsistence Activities." Ottawa, ON: Canadian Councils of Resource Ministers. ([PubMed](#))

Lloyd-Smith, P (2020). "The Economic Benefits of Recreation in Canada". University of Saskatchewan Working Paper

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GenerateMDCEVData

*GenerateMDCEVData*

---

### Description

Simulate data for KT models

**Usage**

```

GenerateMDCEVData(
  model,
  nobs = 1000,
  nalts = 10,
  income = stats::runif(nobs, 1e+05, 150000),
  price = matrix(stats::runif(nobs * nalts, 100, 500), nobs, nalts),
  alpha_parms = 0.5,
  scale_parms = 1,
  gamma_parms = stats::runif(nalts, 1, 10),
  psi_i_parms = c(-1.5, 2, -1),
  psi_j_parms = c(-5, 0.5, 2),
  phi_parms = c(-5, 0.5, 2),
  dat_psi_i = matrix(2 * stats::runif(nobs * length(psi_i_parms)), nobs,
    length(psi_i_parms)),
  dat_psi_j = cbind(matrix(stats::runif(nalts * (length(psi_j_parms))), 0, 1), nrow =
    nalts),
  dat_phi = cbind(matrix(stats::runif(nalts * (length(phi_parms))), 0, 1), nrow =
    nalts),
  nerrs = 1,
  tol = 1e-20,
  max_loop = 999
)

```

**Arguments**

|             |   |
|-------------|---|
| model       | A string indicating which model specification is estimated. The options are "alpha", "gamma", "hybrid" and "hybrid0" for the MDCEV model and "kt_ee" for the environmental economics Kuhn-Tucker specification. |
| nobs        | Number of individuals   |
| nalts       | Number of non-numeraire alts  |
| income      | Vector of individual income   |
| price       | Matrix of prices for non-numeraire alternatives.  |
| alpha_parms | Parameter value for alpha term  |
| scale_parms | Parameter value for scale term  |
| gamma_parms | Parameter value for gamma terms   |
| psi_i_parms | Parameter value for psi terms that vary by individual   |
| psi_j_parms | Parameter value for psi terms that vary by alt (all models except kt_ee)  |
| phi_parms   | Parameter value for phi terms that vary by alt (kt_ee model only)   |
| dat_psi_i   | (nobs X # psi_i_parms) matrix with individual-specific characteristics  |
| dat_psi_j   | (nalts X # psi_j_parms) matrix with alternative-specific variables (all models except kt_ee)  |
| dat_phi     | (nalts X # phi_parms) matrix with alternative-specific variables (kt_ee model only)   |

|          |   |
|----------|---|
| nerrs    | Number of error draws for demand simulation                       |
| tol      | Tolerance level for simulations if using general approach         |
| max_loop | maximum number of loops for simulations if using general approach |

**Value**

A 'mdcev.data' object, which is a 'data.frame' in long format. Also includes parms\_true with parameter values

**Examples**

```
data <- GenerateMDCEVData(model = "gamma")
```

---

|       |              |
|-------|--------------|
| mdcev | <i>mdcev</i> |
|-------|--------------|

---

**Description**

Fit a MDCEV model using MLE or Bayes

**Usage**

```
mdcev(
  formula = NULL,
  data,
  weights = NULL,
  model = c("alpha", "gamma", "hybrid", "hybrid0", "kt_ee"),
  n_classes = 1,
  fixed_scale1 = 0,
  single_scale = 0,
  trunc_data = 0,
  psi_ascs = NULL,
  gamma_ascs = 1,
  seed = "123",
  max_iterations = 2000,
  jacobian_analytical_grad = 1,
  initial.parameters = "random",
  hessian = TRUE,
  algorithm = c("MLE", "Bayes"),
  flat_priors = NULL,
  print_iterations = TRUE,
  prior_psi_sd = 10,
  prior_gamma_sd = 10,
  prior_phi_sd = 10,
  prior_alpha_shape = 1,
```

```

prior_scale_sd = 1,
prior_delta_sd = 10,
gamma_nonrandom = 0,
alpha_nonrandom = 0,
std_errors = "deltamethod",
n_draws = 50,
keep_loglik = 0,
random_parameters = "fixed",
show_stan_warnings = TRUE,
n_iterations = 200,
n_chains = 4,
n_cores = 4,
max_tree_depth = 10,
adapt_delta = 0.8,
lkj_shape_prior = 4,
...
)

## S3 method for class 'mdcev'
print(
  x,
  digits = max(3, getOption("digits") - 3),
  width = getOption("width"),
  ...
)

## S3 method for class 'mdcev'
summary(object, printCI = FALSE, ...)

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

```

## Arguments

|         |   |
|---------|---|
| formula | Formula for the model to be estimated. The formula is divided in three parts, separated by the symbol  . The first part is reserved for alternative-specific and individual-specific variables in the $\psi$ parameters. Note that alternative-specific constants are handled by the <code>psi_asc</code> argument. The second part corresponds for individual-specific variables that enter in the probability assignment in models with latent classes. The third part is reserved for the $q_k$ variables included in the $\phi_k$ parameters in the KT model specification used in environmental economics <code>model = "kt_ee"</code> . |
| data    | The (I x J) data to be passed to Stan of class <code>mdcev.data</code> including 1) id, 2) alt, 3) choice, 4) price, 5) income, and columns for alternative-specific and individual specific variables. Note: I is number of individuals and J is number of non-numeraire alternatives.   |
| weights | an optional vector of weights. Default to 1.  |
| model   | A string indicating which model specification is estimated. The options are   |

|                          |  |
|--------------------------|--|
|                          | "alpha", "gamma", "hybrid" and "hybrid0" for the MDCEV model and "kt_ee" for the environmental economics Kuhn-Tucker specification.  |
| n_classes                | The number of latent classes. Note that the LC model is automatically estimated as long as the prespecified number of classes is set greater than 1.   |
| fixed_scale1             | Whether to fix scale at 1.   |
| single_scale             | For lc models, whether to estimate a single scale parameter  |
| trunc_data               | Whether the estimation should be adjusted for truncation of non-numeraire alternatives. This option is useful if the data only includes individuals with positive non-numeraire consumption levels such as recreation data collected on-site. To account for the truncation of consumption, the likelihood is normalized by one minus the likelihood of observing zero consumption for all non-numeraire alternatives (i.e. likelihood of positive consumption) following Englin, Boxall and Watson (1998) and von Haefen (2003).  |
| psi_ascs                 | Whether to include alternative-specific psi parameters. The first alternative is used as the reference category. Only specify to 1 for MDCEV models.   |
| gamma_ascs               | Indicator to include alternative-specific gammas parameters.   |
| seed                     | Random seed.   |
| max_iterations           | Maximum number of iterations in MLE.   |
| jacobian_analytical_grad | indicator whether to use analytical gradient method for Jacobian (=1) or numerical gradient method (=0). For "kt_ee" model only,   |
| initial.parameters       | The default for fixed and random parameter specifications is to use random starting values. (except for the scale parameter which a starting value set to 1). For LC models, the default is to use slightly adjusted MLE point estimates from the single class model. Initial parameter values should be included in a named list. For example, the LC "hybrid" specification initial parameters can be specified as: <code>initial.parameters = list(psi = array(0, dim = c(K, num_psi)), gamma = array(1, dim = c(K, num_alt)), alpha = array(0.5, dim = c(K, 0)), scale = array(1, dim = c(K)))</code> where K is the number of classes (i.e. K = 1 is used for single class models), num_psi is number of psi parameters, and num_alt is number of non-numeraire alternatives. |
| hessian                  | Whether to keep the Hessian matrix   |
| algorithm                | Either "Bayes" for Bayes or "MLE" for maximum likelihood estimation.   |
| flat_priors              | indicator if completely uninformative priors should be specified. Defaults to 1 if MLE used and 0 if Bayes used. If using MLE and set flat_priors = 0, penalized MLE is used and the optimizing objective is augmented with the priors.  |
| print_iterations         | Whether to print iteration information   |
| prior_psi_sd             | standard deviation for normal prior with mean 0.   |
| prior_gamma_sd           | standard deviation for half-normal prior with mean 1.  |
| prior_phi_sd             | standard deviation for normal prior with mean 0.   |
| prior_alpha_shape        | shape parameter for beta distribution.   |

|                    |   |
|--------------------|---|
| prior_scale_sd     | standard deviation for half-normal prior with mean 0.   |
| prior_delta_sd     | standard deviation for normal prior with mean 0.  |
| gamma_nonrandom    | indicator set to 1 if gamma parameters should not be random (i.e. no standard deviation).   |
| alpha_nonrandom    | indicator set to 1 if alpha parameters should not be random (i.e. no standard deviation).   |
| std_errors         | Compute standard errors using the delta method ("deltamethod") or multivariate normal draws ("mvn"). The default is "deltamethod". Note that mvn parameter draws should be used to incorporate parameter uncertainty for demand and welfare simulation. For maximum likelihood estimation only. |
| n_draws            | The number of multivariate normal draws for standard error calculations if "mvn" is specified.  |
| keep_loglik        | Whether to keep the log_lik calculations  |
| random_parameters  | The form of the covariance matrix for Bayes. Can be 'fixed' for no random parameters, 'uncorr' for uncorrelated random parameters, or 'corr' for correlated random parameters.  |
| show_stan_warnings | Whether to show warnings from Stan.   |
| n_iterations       | The number of iterations to use in Bayesian estimation. The default is for the number of iterations to be split evenly between warmup and posterior draws. The number of warmup draws can be directly controlled using the warmup argument (see <code>rstan::sampling</code> ).                 |
| n_chains           | The number of independent Markov chains in Bayesian estimation.   |
| n_cores            | The number of cores used to execute the Markov chains in parallel in Bayesian estimation. Can set using <code>options(mc.cores = parallel::detectCores())</code> .  |
| max_tree_depth     | <a href="https://mc-stan.org/misc/warnings.html#maximum-treedepth-exceeded">https://mc-stan.org/misc/warnings.html#maximum-treedepth-exceeded</a>   |
| adapt_delta        | <a href="https://mc-stan.org/misc/warnings.html#divergent-transitions-after-warmup">https://mc-stan.org/misc/warnings.html#divergent-transitions-after-warmup</a>   |
| lkj_shape_prior    | Prior for Cholesky matrix   |
| ...                | Additional parameters to pass on to <code>rstan::stan</code> and <code>rstan::sampling</code> .   |
| x, object          | an object of class 'mdcev'  |
| digits             | the number of digits,   |
| width              | the width of the printing,  |
| printCI            | set to TRUE to print 95% confidence intervals   |

### Value

A object of class `mdcev`



## Examples

```
data(data_rec, package = "rmdcev")

data_rec <- mdcev.data(data_rec, subset = id <= 500, id.var = "id",
                      alt.var = "alt", choice = "quant")

mdcev_est <- mdcev( ~ 0,
                  data = data_rec,
                  model = "hybrid0",
                  algorithm = "MLE")
```

---

|            |                                   |
|------------|-----------------------------------|
| mdcev.data | <i>data.frame for mdcev model</i> |
|------------|-----------------------------------|

---

## Description

shape a 'data.frame' in a suitable form for the use of the 'mdcev' function and complete some data checks

## Usage

```
mdcev.data(
  data,
  id.var = "id",
  alt.var = NULL,
  choice = "choice",
  price = "price",
  income = "income",
  alt.levels = NULL,
  drop.index = FALSE,
  subset = NULL,
  ...
)
```

## Arguments

|         |  |
|---------|--|
| data    | a 'data.frame',  |
| id.var  | the name of the variable that contains the individual index.   |
| alt.var | the name of the variable that contains the alternative index or the name under which the alternative index will be stored (the default name is 'alt'), |
| choice  | the variable indicating the consumption of non-numeraire alternatives that is made: it has to be a numerical vector. Default is "choice".              |
| price   | the variable indicating the price of the non-numeraire alternatives. Default is "price"  |

|            |  |
|------------|--|
| income     | the variable indicating the income of the individual. Default is "income".           |
| alt.levels | the name of the alternatives: if null, they are guessed from the 'alt.var' argument, |
| drop.index | should the index variables be dropped from the 'data.frame',                         |
| subset     | a logical expression which defines the subset of observations to be selected,        |
| ...        | further arguments.   |

### Value

A 'mdcev.data' object, which is a 'data.frame' in long format, \*i.e.\* one line for each alternative. It has a 'index' attribute, which is a 'data.frame' that contains the index of the individual ('id') and the index of the alternative ('alt').

---

mdcev.sim

*mdcev.sim*

---

### Description

Simulate welfare or demand for MDCEV model

### Usage

```
mdcev.sim(
  df_indiv,
  df_common,
  sim_options,
  sim_type = c("welfare", "demand"),
  nerrs = 30,
  cond_error = 1,
  draw_mlhs = 1,
  algo_gen = NULL,
  tol = 1e-20,
  max_loop = 999,
  suppressTime = FALSE,
  stan_seed = 3,
  ...
)

## S3 method for class 'mdcev.sim'
print(
  x,
  digits = max(3, getOption("digits") - 3),
  width = getOption("width"),
  ...
)

## S3 method for class 'mdcev.sim'
```

```
summary(object, ci = 0.95, ...)

## S3 method for class 'summary.mdcev.sim'
print(
  x,
  digits = max(3, getOption("digits") - 2),
  width = getOption("width"),
  ...
)
```

### Arguments

|                           |  |
|---------------------------|--|
| <code>df_indiv</code>     | Prepared individual level data from <code>PrepareSimulationData</code>   |
| <code>df_common</code>    | Prepared common data from <code>PrepareSimulationData</code>   |
| <code>sim_options</code>  | Prepared simulation options from <code>PrepareSimulationData</code>  |
| <code>sim_type</code>     | Either "welfare" or "demand"   |
| <code>nerrs</code>        | Number of error draws for welfare analysis   |
| <code>cond_error</code>   | Choose whether to draw errors conditional on actual demand or not. Conditional error draws (=1) or unconditional error draws.  |
| <code>draw_mlhc</code>    | Generate draws using Modified Latin Hypercube Sampling algorithm (=1) or uniform (=0)  |
| <code>algo_gen</code>     | Type of algorithm for simulation. <code>algo_gen = 0</code> for Hybrid Approach (i.e. constant alphas, only hybrid models) <code>algo_gen = 1</code> for General approach (i.e. heterogeneous alpha's, all models) |
| <code>tol</code>          | Tolerance level for simulations if using general approach  |
| <code>max_loop</code>     | maximum number of loops for simulations if using general approach  |
| <code>suppressTime</code> | Suppress simulation time calculation   |
| <code>stan_seed</code>    | Seed for pseudo-random number generator <code>get_rng</code> see <code>help(get_rng, package = "rstan")</code>   |
| <code>...</code>          | Additional parameters to pass to <code>mdcev.sim</code>  |
| <code>x, object</code>    | an object of class 'mdcev.sim'   |
| <code>digits</code>       | the number of digits,  |
| <code>width</code>        | the width of the printing,   |
| <code>ci</code>           | choose confidence interval for simulations. Default is 95 percent.   |

### Value

a object of class `mdcev.sim` which contains a list for each individual holding either 1) `nsims x npols` matrix of welfare changes if welfare is being simulated or 2) `nsims` number of lists of `npols x #` alternatives matrix of Marshallian demands is demand is being simulated.

### See Also

[`mdcev()`] for the estimation of `mdcev` models.

**Examples**

```

data(data_rec, package = "rmdcev")

data_rec <- mdcev.data(data_rec, subset = id <= 500, id.var = "id",
  alt.var = "alt", choice = "quant")

mdcev_est <- mdcev(~ 0, data = data_rec,
  model = "hybrid0", algorithm = "MLE",
  std_errors = "mvn")

policies <- CreateBlankPolicies(npols = 2,
  mdcev_est,
  price_change_only = TRUE)

df_sim <- PrepareSimulationData(mdcev_est, policies)

wtp <- mdcev.sim(df_sim$df_indiv,
  df_common = df_sim$df_common,
  sim_options = df_sim$sim_options,
  cond_err = 1, nerrs = 5, sim_type = "welfare")

```

---

PrepareSimulationData *PrepareSimulationData*

---

**Description**

Prepare Data for WTP simulation

**Usage**

```
PrepareSimulationData(object, policies, nsims = 30, class = "class1")
```

**Arguments**

|                       |  |
|-----------------------|--|
| <code>object</code>   | an object of class ‘mdcev’   |
| <code>policies</code> | list containing price_p with additive price increases, and dat_psi_p with new psi data |
| <code>nsims</code>    | Number of simulation draws to use for parameter uncertainty                            |
| <code>class</code>    | The class number for Latent Class models.  |

**Value**

A list with individual-specific data (`df_indiv`) and common data (`df_common`) and `n_classes` for number of classes and `model_num` for model type

## Examples

```
data(data_rec, package = "rmdcev")

data_rec <- mdcev.data(data_rec, subset = id <= 500, id.var = "id",
  alt.var = "alt", choice = "quant")

mdcev_est <- mdcev( ~ 0, data = data_rec,
  model = "hybrid0", algorithm = "MLE",
  std_errors = "mvn")

policies <- CreateBlankPolicies(npols = 2, mdcev_est,
  price_change_only = TRUE)

df_sim <- PrepareSimulationData(mdcev_est, policies)
```

---

|        |   |
|--------|---|
| rmdcev | <i>rmdcev: Estimating and simulating Kuhn-Tucker and multiple discrete-continuous extreme value (MDCEV) demand models</i> |
|--------|---|

---

## Description

The `rmdcev` R package estimate and simulates Kuhn-Tucker demand models with individual heterogeneity. The models supported by `rmdcev` are the multiple-discrete continuous extreme value (MDCEV) model and Kuhn-Tucker specification common in the environmental economics literature on recreation demand. Latent class and random parameters specifications can be implemented and the models are fit using maximum likelihood estimation or Bayesian estimation. All models are implemented in Stan, which is a C++ package for performing full Bayesian inference (see Stan Development Team, 2019) <<https://mc-stan.org/>>. The `rmdcev` package also implements demand forecasting (Pinjari and Bhat (2011) <<https://repositories.lib.utexas.edu/handle/2152/23880>>) and welfare calculation (Lloyd-Smith (2018) <[doi.org/10.1016/j.jocm.2017.12.002](https://doi.org/10.1016/j.jocm.2017.12.002)>) for policy simulation.

## Author(s)

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## References

Bhat, CR (2008). The multiple discrete-continuous extreme value (MDCEV) model: Role of utility function parameters, identification considerations, and model extensions. *Transportation Research Part B: Methodological*, 42(3): 274-303. [\(link\)](#) Lloyd-Smith, P (2018). A new approach to calculating welfare measures in Kuhn-Tucker demand models. *Journal of Choice Modeling* 26:19–27. [\(link\)](#) Pinjari, AR, Bhat, CR (2011). Computationally Efficient Forecasting Procedures for Kuhn-Tucker Consumer Demand Model Systems: Application to Residential Energy Consumption Analysis. Department of Civil and Environmental Engineering, University of South Florida. [\(link\)](#) Stan Development Team (2019). RStan: the R interface to Stan. R package version 2.19.2. [\(link\)](#)

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