

Package ‘DMCfun’

January 10, 2021

Type Package

Title Diffusion Model of Conflict (DMC) in Reaction Time Tasks

Version 1.3.0

Date 2020-12-27

Description DMC model simulation detailed in Ulrich, R., Schroeter, H., Leuthold, H., & Birngruber, T. (2015). Automatic and controlled stimulus processing in conflict tasks: Superimposed diffusion processes and delta functions. *Cognitive Psychology*, 78, 148-174. Ulrich et al. (2015) <doi:10.1016/j.cogpsych.2015.02.005>. Decision processes within choice reaction-time (CRT) tasks are often modelled using evidence accumulation models (EAMs), a variation of which is the Diffusion Decision Model (DDM, for a review, see Ratcliff & McKoon, 2008). Ulrich et al. (2015) introduced a Diffusion Model for Conflict tasks (DMC). The DMC model combines common features from within standard diffusion models with the addition of superimposed controlled and automatic activation. The DMC model is used to explain distributional reaction time (and error rate) patterns in common behavioural conflict-like tasks (e.g., Flanker task, Simon task). This R-package implements the DMC model and provides functionality to fit the model to observed data.

URL <https://github.com/igmmgi/DMCfun>,
<https://CRAN.R-project.org/package=DMCfun>

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Encoding UTF-8

Depends R (>= 3.5.0)

Imports DEoptim, Rcpp (>= 0.12.16), dplyr (>= 1.0.0), optimr,
parallel, pbapply, tidy

Suggests testthat

LinkingTo Rcpp, BH

RoxygenNote 7.1.1

LazyData true

SystemRequirements C++11

NeedsCompilation yes

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Repository CRAN

Date/Publication 2021-01-10 17:50:10 UTC

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addDataDF	<i>addDataDF: Add data to a simulated dataframe</i>
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Description

Add simulated ex-gaussian reaction-time (RT) data and binary error (Error = 1, Correct = 0) data to an R DataFrame. This function can be used to create simulated data sets.

Usage

```
addDataDF(dat, RT = NULL, Error = NULL)
```

Arguments

dat	DataFrame (see createDF)
RT	RT parameters (see rtDist)
Error	Error parameters (see errDist)

Value

DataFrame with RT (ms) and Error (bool) columns

Examples

```
# Example 1: default dataframe
dat <- createDF()
dat <- addDataDF(dat)
head(dat)
hist(dat$RT, 100)
table(dat$error)

# Example 2: defined overall RT parameters
dat <- createDF(nSubjects = 50, nTrl = 50, design = list("Comp" = c("comp", "incomp")))
dat <- addDataDF(dat, RT = c(500, 150, 100))
boxplot(dat$RT ~ dat$Comp)
table(dat$Comp, dat$error)

# Example 3: defined RT + Error parameters across conditions
dat <- createDF(nSubjects = 50, nTrl = 50, design = list("Comp" = c("comp", "incomp")))
dat <- addDataDF(dat,
  RT = list("Comp_comp" = c(500, 80, 100),
           "Comp_incomp" = c(600, 80, 140)),
  Error = list("Comp_comp" = 5,
              "Comp_incomp" = 15))
boxplot(dat$RT ~ dat$Comp)
table(dat$Comp, dat$error)

# Example 4:
# create dataframe with defined RT + Error parameters across different conditions
```

```

dat <- createDF(nSubjects = 50, nTrl = 50, design = list("Comp" = c("comp", "incomp", "neutral")))
dat <- addDataDF(dat,
  RT = list("Comp_comp"      = c(500, 150, 100),
            "Comp_neutral"   = c(550, 150, 100),
            "Comp_incomp"    = c(600, 150, 100)),
  Error = list("Comp_comp"   = 5,
              "Comp_neutral" = 10,
              "Comp_incomp"  = 15))

boxplot(dat$RT ~ dat$Comp)
table(dat$Comp, dat$Error)

# Example 5:
# create dataframe with defined RT + Error parameters across different conditions
dat <- createDF(nSubjects = 50, nTrl = 50,
  design = list("Hand" = c("left", "right"),
              "Side" = c("left", "right")))
dat <- addDataDF(dat,
  RT = list("Hand:Side_left:left" = c(400, 150, 100),
            "Hand:Side_left:right" = c(500, 150, 100),
            "Hand:Side_right:left" = c(500, 150, 100),
            "Hand:Side_right:right" = c(400, 150, 100)),
  Error = list("Hand:Side_left:left" = c(5,4,2,2,1),
              "Hand:Side_left:right" = c(15,4,2,2,1),
              "Hand:Side_right:left" = c(15,7,4,2,1),
              "Hand:Side_right:right" = c(5,8,5,3,1)))

boxplot(dat$RT ~ dat$Hand + dat$Side)
table(dat$Error, dat$Hand, dat$Side)

```

addErrorBars

addErrorBars: Add errorbars to plot.

Description

Add error bars to current plot (uses base arrows function).

Usage

```
addErrorBars(xpos, ypos, errorSize, arrowSize = 0.1)
```

Arguments

xpos	x-position of data-points
ypos	y-position of data-points
errorSize	+ - size of error bars
arrowSize	Width of the errorbar arrow

Value

Plot

Examples

```
# Example 1
plot(c(1, 2), c(450, 500), xlim = c(0.5, 2.5), ylim = c(400, 600), type = "o")
addErrorBars(c(1, 2), c(450, 500), errorSize = c(20, 20))

# Example 2
plot(c(1, 2), c(450, 500), xlim = c(0.5, 2.5), ylim = c(400, 600), type = "o")
addErrorBars(c(1, 2), c(450, 500), errorSize = c(20, 40), arrowSize = 0.2)
```

calculateCAF

*calculateCAF: Calculate conditional accuracy function (CAF).***Description**

Calculate conditional accuracy function (CAF). The DataFrame should contain columns defining the participant, compatibility condition, RT and error (Default column names: "Subject", "Comp", "RT", "Error"). The "Comp" column should define compatibility condition (Default: c("comp", "incomp")) and the "Error" column should define if the trial was an error or not (Default: c(0, 1)).

Usage

```
calculateCAF(
  dat,
  nCAF = 5,
  columns = c("Subject", "Comp", "RT", "Error"),
  compCoding = c("comp", "incomp"),
  errorCoding = c(0, 1)
)
```

Arguments

dat	DataFrame with columns containing the participant number, condition compatibility, RT data (in ms) and an Error column.
nCAF	Number of CAF bins.
columns	Name of required columns Default: c("Subject", "Comp", "RT", "Error")
compCoding	Coding for compatibility Default: c("comp", "incomp")
errorCoding	Coding for errors Default: c(0, 1)

Value

DataFrame

Examples

```

# Example 1
dat <- createDF(nSubjects = 1, nTrl = 100, design = list("Comp" = c("comp", "incomp")))
dat <- addDataDF(dat,
  RT = list("Comp_comp" = c(500, 80, 100),
    "Comp_incomp" = c(600, 80, 140)),
  Error = list("Comp_comp" = c(5, 4, 3, 2, 1),
    "Comp_incomp" = c(20, 8, 6, 4, 2)))
caf <- calculateCAF(dat)

# Example 2
dat <- createDF(nSubjects = 1, nTrl = 100, design = list("Congruency" = c("cong", "incong")))
dat <- addDataDF(dat,
  RT = list("Congruency_cong" = c(500, 80, 100),
    "Congruency_incong" = c(600, 80, 140)),
  Error = list("Congruency_cong" = c(5, 4, 3, 2, 1),
    "Congruency_incong" = c(20, 8, 6, 4, 2)))
head(dat)
caf <- calculateCAF(dat, columns = c("Subject", "Congruency", "RT", "Error"),
  compCoding = c("cong", "incong"))

```

calculateCostValue *calculateCostValue: Calculate RMSE from RT and error data*

Description

Calculate cost value (fit) from combination of RT and error rate.

Usage

```
calculateCostValue(resTh, resOb)
```

Arguments

resTh	list containing caf values for comp/incomp conditions (nbins*2*3) and delta values for comp/incomp conditions (nbins*5). See output from dmcSim (.Scaf).
resOb	list containing caf values for comp/incomp conditions (n*2*3) and delta values for comp/incomp conditions (nbins*5). See output from dmcSim (\$delta).

Value

cost value (RMSE)

Examples

```
# Example 1:
resTh <- dmcSim()
resOb <- dmcSim()
cost <- calculateCostValue(resTh, resOb)

# Example 2:
resTh <- dmcSim()
resOb <- dmcSim(tau = 150)
cost <- calculateCostValue(resTh, resOb)
```

calculateDelta	<i>calculateDelta: Calculate delta function</i>
----------------	---

Description

Calculate delta plot. Here RTs are split into n bins (Default: 5) for compatible and incompatible trials separately. Mean RT is calculated for each condition in each bin then subtracted (incompatible - compatible) to give a compatibility effect (delta) at each bin.

Usage

```
calculateDelta(
  dat,
  nDelta = 19,
  columns = c("Subject", "Comp", "RT"),
  compCoding = c("comp", "incomp"),
  quantileType = 5
)
```

Arguments

dat	DataFrame with columns containing the participant number, condition compatibility, and RT data (in ms).
nDelta	Number of delta bins.
columns	Name of required columns Default: c("Subject", "Comp", "RT")
compCoding	Coding for compatibility Default: c("comp", "incomp")
quantileType	Argument (1-9) from R function quantile specifying the algorithm (?quantile)

Value

DataFrame

Examples

```

# Example 1
dat <- createDF(nSubjects = 50, nTrl = 100, design = list("Comp" = c("comp", "incomp")))
dat <- addDataDF(dat,
  RT = list("Comp_comp" = c(500, 80, 100),
    "Comp_incomp" = c(600, 80, 140)))
delta <- calculateDelta(dat)

# Example 2
dat <- createDF(nSubject = 50, nTrl = 100, design = list("Congruency" = c("cong", "incong")))
dat <- addDataDF(dat,
  RT = list("Congruency_cong" = c(500, 80, 100),
    "Congruency_incong" = c(600, 80, 140)))
head(dat)
delta <- calculateDelta(dat, columns = c("Subject", "Congruency", "RT"),
  compCoding = c("cong", "incong"))

```

createDF

createDF: Create a simulated dataframe

Description

Create dataframe (see also addDataDF)

Usage

```

createDF(
  nSubjects = 20,
  nTrl = 50,
  design = list(A = c("A1", "A2"), B = c("B1", "B2"))
)

```

Arguments

nSubjects	Number of subjects
nTrl	Number of trials per factor/level for each participant
design	Factors and levels

Value

dataframe

Examples

```
# Example 1
dat <- createDF()

# Example 2
dat <- createDF(nSubjects = 50, nTrl = 50, design = list("Comp" = c("comp", "incomp")))

# Example 3
dat <- createDF(nSubjects = 50, nTrl = 50, design = list("Comp" = c("comp", "incomp"),
                                                       "Side" = c("left", "right", "middle")))
```

dmcCombineObservedData

dmcCombineObservedData: Combine results from dmcObservedData

Description

Combine observed datasets

Usage

```
dmcCombineObservedData(...)
```

Arguments

... Any number of outputs from dmcObservedData

Value

dmcobs

Examples

```
# Example 1
datFlanker <- dmcObservedData(flankerDataRow, nDelta = 9)
datSimon <- dmcObservedData(simonDataRow, nDelta = 9)
dat <- dmcCombineObservedData(datFlanker, datSimon) # combine flanker/simon data
plot(dat, figType = "delta", xlimDelta = c(200, 700),
     cols = c("black", "darkgrey"), pchs = c(1, 2), resetPar = FALSE)
legend(200, 0, legend = c("Flanker Task", "Simon Task"),
     col = c("black", "darkgrey"), lty = c(1, 1))
```

dmcCppR

dmcCppR

Description

dmcCppR

dmcFit

dmcFit: Fit DMC to aggregated data using R-package optimr (Nelder-Mead)

Description

Fit theoretical data generated from dmcSim to observed data by minimizing the root-mean-square error (RMSE) between a weighted combination of the CAF and CDF functions using the R-package optimr (Nelder-Mead).

Usage

```
dmcFit(
  resOb,
  nTrl = 1e+05,
  startVals = list(),
  minVals = list(),
  maxVals = list(),
  fixedFit = list(),
  fitInitialGrid = TRUE,
  fitInitialGridN = 10,
  fixedGrid = list(),
  nCAF = 5,
  nDelta = 19,
  pDelta = vector(),
  varSP = TRUE,
  rtMax = 5000,
  printInputArgs = TRUE,
  printResults = FALSE,
  maxit = 500
)
```

Arguments

resOb Observed data (see flankerData and simonTask for data format)
nTrl Number of trials to use within dmcSim.

startVals	Starting values for to-be estimated parameters. This is a list with values specified individually for amp, tau, drc, bnds, resMean, resSD, aaShape, spShape, sigm (e.g., startVals = list(amp = 20, tau = 200, drc = 0.5, bnds = 75, resMean = 300, resSD = 30, aaShape = 2, spShape = 3, sigm = 4)).
minVals	Minimum values for the to-be estimated parameters. This is a list with values specified individually for amp, tau, drc, bnds, resMean, resSD, aaShape, spShape, sigm (e.g., minVals = list(amp = 10, tau = 5, drc = 0.1, bnds = 20, resMean = 200, resSD = 5, aaShape = 1, spShape = 2, sigm = 1)).
maxVals	Maximum values for the to-be estimated parameters. This is a list with values specified individually for amp, tau, drc, bnds, resMean, resSD, aaShape, spShape, sigm (e.g., maxVals = list(amp = 40, tau = 300, drc = 1.0, bnds = 150, resMean = 800, resSD = 100, aaShape = 3, spShape = 4, sigm = 10))
fixedFit	Fix parameter to starting value. This is a list with bool values specified individually for amp, tau, drc, bnds, resMean, resSD, aaShape, spShape, sigm (e.g., fixedFit = list(amp = F, tau = F, drc = F, bnds = F, resMean = F, resSD = F, aaShape = F, spShape = F, sigm = T))
fitInitialGrid	TRUE/FALSE
fitInitialGridN	10 reduce if searching more than 1 initial parameter
fixedGrid	Fix parameter for initial grid search. This is a list with bool values specified individually for amp, tau, drc, bnds, resMean, resSD, aaShape, spShape, sigm (e.g., fixedGrid = list(amp = T, tau = F, drc = T, bnds = T, resMean = T, resSD = T, aaShape = T, spShape = T, sigm = T))
nCAF	Number of CAF bins.
nDelta	Number of delta bins.
pDelta	Alternative to nDelta by directly specifying required percentile values
varSP	Variable starting point TRUE/FALSE
rtMax	limit on simulated RT (decision + non-decisional component)
printInputArgs	TRUE/FALSE
printResults	TRUE/FALSE
maxit	The maximum number of iterations (Default: 500)

Value

dmcfit

The function returns a list with the relevant results from the fitting procedure. The list is accessed with obj\$name with the the following:

obj\$means	Condition means for reaction time and error rate
obj\$caf	Accuracy per bin for compatible and incompatible trials
obj\$delta	Mean RT and compatibility effect per bin
obj\$sim	Individual trial data points (reaction times/error) and activation vectors from simulation
obj\$par	The fitted model parameters + final RMSE of the fit

Examples

```
# Example 1: Flanker data from Ulrich et al. (2015)
fit <- dmcFit(flankerData) # only initial search tau
plot(fit, flankerData)
summary(fit)

# Example 2: Simon data from Ulrich et al. (2015)
fit <- dmcFit(simonData) # only initial search tau
plot(fit, simonData)
summary(fit)

# Example 3: Flanker data from Ulrich et al. (2015) with non-default
# start vals and some fixed values
fit <- dmcFit(flankerData,
              startVals = list(drc = 0.6, aaShape = 2.5),
              fixedFit = list(drc = TRUE, aaShape = TRUE))

# Example 4: Simulated Data (+ve going delta function)
dat <- createDF(nSubjects = 20, nTrl = 500, design = list("Comp" = c("comp", "incomp")))
dat <- addDataDF(dat,
                 RT = list("Comp_comp" = c(510, 100, 100),
                           "Comp_incomp" = c(540, 130, 85)),
                 Error = list("Comp_comp" = c(4, 3, 2, 1, 1),
                              "Comp_incomp" = c(20, 4, 3, 1, 1)))
datOb <- dmcObservedData(dat, columns = c("Subject", "Comp", "RT", "Error"))
plot(datOb)
fit <- dmcFit(datOb, nTrl = 5000)
plot(fit, datOb)
summary(fit)
```

dmcFitDE

dmcFitDE: Fit DMC to aggregated data using R-package DEoptim

Description

Fit theoretical data generated from `dmcSim` to observed data by minimizing the root-mean-square error (RMSE) between a weighted combination of the CAF and CDF functions using the R-package `DEoptim`.

Usage

```
dmcFitDE(
  resOb,
  nTrl = 1e+05,
  minVals = list(),
  maxVals = list(),
```

```

    fixedFit = list(),
    nCAF = 5,
    nDelta = 19,
    pDelta = vector(),
    varSP = TRUE,
    rtMax = 5000,
    control = list()
  )

```

Arguments

resOb	Observed data (see flankerData and simonTask for data format)
nTr1	Number of trials to use within dmcSim.
minVals	Minimum values for the to-be estimated parameters. This is a list with values specified individually for amp, tau, drc, bnds, resMean, resSD, aaShape, spShape, sigm (e.g., minVals = list(amp = 10, tau = 5, drc = 0.1, bnds = 20, resMean = 200, resSD = 5, aaShape = 1, spShape = 2, sigm = 1)).
maxVals	Maximum values for the to-be estimated parameters. This is a list with values specified individually for amp, tau, drc, bnds, resMean, resSD, aaShape, spShape, sigm (e.g., maxVals = list(amp = 40, tau = 300, drc = 1.0, bnds = 150, resMean = 800, resSD = 100, aaShape = 3, spShape = 4, sigm = 10))
fixedFit	Fix parameter to starting value. This is a list with bool values specified individually for amp, tau, drc, bnds, resMean, resSD, aaShape, spShape, sigm (e.g., fixedFit = list(amp = F, tau = F, drc = F, bnds = F, resMean = F, resSD = F, aaShape = F, spShape = F, sigm = T))
nCAF	Number of CAF bins.
nDelta	Number of delta bins.
pDelta	Alternative to nDelta by directly specifying required percentile values
varSP	Variable starting point TRUE/FALSE
rtMax	limit on simulated RT (decision + non-decisional component)
control	Additional control parameters passes to DEoptim

Value

dmcfit

The function returns a list with the relevant results from the fitting procedure. The list is accessed with obj\$name with the the following:

obj\$means	Condition means for reaction time and error rate
obj\$caf	Accuracy per bin for compatible and incompatible trials
obj\$delta	Mean RT and compatibility effect per bin
obj\$sim	Individual trial data points (reaction times/error) and activation vectors from simulation
obj\$par	The fitted model parameters + final RMSE of the fit

Examples

```
# Example 1: Flanker data from Ulrich et al. (2015)
fit <- dmcFitDE(flankerData)
plot(fit, flankerData)
summary(fit)

# Example 2: Simon data from Ulrich et al. (2015)
fit <- dmcFitDE(simonData, nTr1 = 20000)
plot(fit, simonData)
summary(fit)
```

dmcFitSubject

dmcFitSubject: Fit DMC to aggregated data using R-package optimr (Nelder-Mead)

Description

Fit theoretical data generated from dmcSim to observed data by minimizing the root-mean-square error (RMSE) between a weighted combination of the CAF and CDF functions.

Usage

```
dmcFitSubject(
  resOb,
  nTr1 = 1e+05,
  startVals = list(),
  minVals = list(),
  maxVals = list(),
  fixedFit = list(),
  fitInitialGrid = TRUE,
  fitInitialGridN = 10,
  fixedGrid = list(),
  nCAF = 5,
  nDelta = 19,
  pDelta = vector(),
  varSP = TRUE,
  rtMax = 5000,
  subjects = c(),
  printInputArgs = TRUE,
  printResults = FALSE,
  maxit = 500
)
```

Arguments

resOb	Observed data (see flankerData, simonData for data format)
nTrl	Number of trials to use within dmcSim
startVals	Starting values for to-be estimated parameters
minVals	Minimum values for the to-be estimated parameters
maxVals	Maximum values for the to-be estimated parameters
fixedFit	Fix parameter to starting value
fitInitialGrid	TRUE/FALSE (NB. overrides fitInitialTau)
fitInitialGridN	10
fixedGrid	Fix parameter for initial grid search
nCAF	Number of CAF bins.
nDelta	Number of delta bins.
pDelta	Alternative to nDelta by directly specifying required percentile values
varSP	Variable starting point TRUE/FALSE
rtMax	limit on simulated RT (decision + non-decisional component)
subjects	NULL (aggregated data across all subjects) or integer for subject number
printInputArgs	TRUE/FALSE
printResults	TRUE/FALSE
maxit	The maximum number of iterations (Default: 500)

Value

dmcfit_subject List of dmcfit per subject fitted (see dmcFit)

Examples

```
# Example 1: Flanker data from Ulrich et al. (2015)
fit <- dmcFitSubject(flankerData, nTrl = 1000, subjects = c(1, 2))
plot(fit, flankerData, subject = 1)
plot(fit, flankerData, subject = 2)
summary(fit)

# Example 2: Simon data from Ulrich et al. (2015)
fit <- dmcFitSubject(simonData, nTrl = 1000, subject = c(1, 2))
plot(fit, simonData, subject = 1)
plot(fit, simonData, subject = 2)
summary(fit)
```

dmcFitSubjectDE	<i>dmcFitSubjectDE: Fit DMC to aggregated data using R-package DE-optim.</i>
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Description

Fit theoretical data generated from dmcSim to observed data by minimizing the root-mean-square error (RMSE) between a weighted combination of the CAF and CDF functions.

Usage

```
dmcFitSubjectDE(
  resOb,
  nTr1 = 1e+05,
  minVals = list(),
  maxVals = list(),
  fixedFit = list(),
  nCAF = 5,
  nDelta = 19,
  pDelta = vector(),
  varSP = TRUE,
  rtMax = 5000,
  subjects = c(),
  control = list()
)
```

Arguments

resOb	Observed data (see flankerData, simonData for data format)
nTr1	Number of trials to use within dmcSim
minVals	Minimum values for the to-be estimated parameters
maxVals	Maximum values for the to-be estimated parameters
fixedFit	Fix parameter to starting value
nCAF	Number of CAF bins.
nDelta	Number of delta bins.
pDelta	Alternative to nDelta by directly specifying required percentile values
varSP	Variable starting point TRUE/FALSE
rtMax	limit on simulated RT (decision + non-decisional component)
subjects	NULL (aggregated data across all subjects) or integer for subject number
control	Additional control parameters passes to DEoptim

Value

dmcfit_subject List of dmcfit per subject fitted (see dmcFitDM)

Examples

```
# Example 1: Flanker data from Ulrich et al. (2015)
fit <- dmcFitSubjectDE(flankerData, nTrl = 1000, subjects = c(1, 2))
plot(fit, flankerData, subject = 1)
plot(fit, flankerData, subject = 2)
summary(fit)

# Example 2: Simon data from Ulrich et al. (2015)
fit <- dmcFitSubjectDE(simonData, nTrl = 1000, subject = c(1, 2))
plot(fit, simonData, subject = 1)
plot(fit, simonData, subject = 2)
summary(fit)
```

dmcObservedData

dmcObservedData: Run standard analyses on observed data

Description

Basic example analysis script to create data object required for observed data. Example raw *.txt files are flankerData.txt and simonData.txt. There are four critical columns: A column containing subject number A column coding for compatible or incompatible A column with RT (in ms) A column indicating of the response was correct

Usage

```
dmcObservedData(
  dat,
  nCAF = 5,
  nDelta = 19,
  outlier = c(200, 1200),
  columns = c("Subject", "Comp", "RT", "Error"),
  compCoding = c("comp", "incomp"),
  errorCoding = c(0, 1),
  quantileType = 5,
  delim = "\t",
  skip = 0
)
```

Arguments

dat	Text file(s) containing the observed data or an R DataFrame (see createDF/addDataDF)
nCAF	Number of CAF bins.
nDelta	Number of delta bins.
outlier	Outlier limits in ms (e.g., c(200, 1200))

columns	Name of required columns DEFAULT = c("Subject", "Comp", "RT", "Error")
compCoding	Coding for compatibility DEFAULT = c("comp", "incomp")
errorCoding	Coding for errors DEFAULT = c(0, 1)
quantileType	Argument (1-9) from R function quantile specifying the algorithm (?quantile)
delim	Single character used to separate fields within a record if reading from external text file.
skip	Number of lines to skip before reading data if reading from external text file.

Value

DataFrame

Examples

```
# Example 1
plot(flankerData) # flanker data from Ulrich et al. (2015)

# Example 2
plot(simonData) # simon data from Ulrich et al. (2015)

# Example 3 (Basic behavioural analysis from Ulrich et al. 2015)
flankerDat <- cbind(Task = "flanker", flankerData$summarySubject)
simonDat <- cbind(Task = "simon", simonData$summarySubject)
datAgg <- rbind(flankerDat, simonDat)

datAgg$Subject <- factor(datAgg$Subject)
datAgg$Task <- factor(datAgg$Task)
datAgg$Comp <- factor(datAgg$Comp)

aovErr <- aov(perErr ~ Comp*Task + Error(Subject/(Comp*Task)), datAgg)
summary(aovErr)
model.tables(aovErr, type = "mean")

aovRt <- aov(rtCor ~ Comp*Task + Error(Subject/(Comp*Task)), datAgg)
summary(aovRt)
model.tables(aovRt, type = "mean")

# Example 4
dat <- createDF(nSubjects = 50, nTrl = 500, design = list("Comp" = c("comp", "incomp")))
dat <- addDataDF(dat,
  RT = list("Comp_comp" = c(500, 75, 120),
    "Comp_incomp" = c(530, 75, 100)),
  Error = list("Comp_comp" = c(3, 2, 2, 1, 1),
    "Comp_incomp" = c(21, 3, 2, 1, 1)))
datOb <- dmcObservedData(dat)
plot(datOb)
plot(datOb, subject = 1)

# Example 5
dat <- createDF(nSubjects = 50, nTrl = 500, design = list("Congruency" = c("cong", "incong")))
dat <- addDataDF(dat,
```

```

RT = list("Congruency_cong" = c(500, 75, 100),
         "Congruency_incong" = c(530, 100, 110)),
Error = list("Congruency_cong" = c(3, 2, 2, 1, 1),
            "Congruency_incong" = c(21, 3, 2, 1, 1))
datOb <- dmcObservedData(dat, nCAF = 5, nDelta = 9,
                       columns = c("Subject", "Congruency", "RT", "Error"),
                       compCoding = c("cong", "incong"))
plot(datOb, labels = c("Congruent", "Incongruent"))
plot(datOb, subject = 1)

```

dmcSim

dmcSim: Run dmc simulation

Description

DMC model simulation detailed in Ulrich, R., Schroeter, H., Leuthold, H., & Birngruber, T. (2015). Automatic and controlled stimulus processing in conflict tasks: Superimposed diffusion processes and delta functions. *Cognitive Psychology*, 78, 148-174. This function is essentially a wrapper around the c++ function runDMC

Usage

```

dmcSim(
  amp = 20,
  tau = 30,
  drc = 0.5,
  bnds = 75,
  resMean = 300,
  resSD = 30,
  aaShape = 2,
  spShape = 3,
  sigm = 4,
  nTrl = 1e+05,
  tmax = 1000,
  varSP = FALSE,
  spLim = c(-75, 75),
  varDR = FALSE,
  drShape = 3,
  drLim = c(0.1, 0.7),
  rtMax = 5000,
  fullData = FALSE,
  nTrlData = 5,
  nDelta = 9,
  pDelta = vector(),
  nCAF = 5,
  printInputArgs = TRUE,
  printResults = TRUE,

```

```

    setSeed = FALSE
  )

```

Arguments

amp	amplitude of automatic activation
tau	time to peak automatic activation
drc	drift rate of controlled processes
bnds	+/- response criterion
resMean	mean of non-decisional component
resSD	standard deviation of non-decisional component
aaShape	shape parameter of automatic activation
spShape	shape parameter of starting point
sigm	diffusion constant
nTrl	number of trials
tmax	number of time points per trial
varSP	true/false variable starting point
spLim	limit range of distribution of starting point
varDR	true/false variable drift rate NB. In DMC, drift rate across trials is always constant.
drShape	shape parameter of drift rate
drLim	limit range of distribution of drift rate
rtMax	limit on simulated RT (decision + non-decisional component)
fullData	TRUE/FALSE (Default: FALSE)
nTrlData	Number of trials to plot
nDelta	Number of delta bins
pDelta	Alternative to nDelta by directly specifying required percentile values
nCAF	Number of CAF bins
printInputArgs	TRUE/FALSE
printResults	TRUE/FALSE
setSeed	TRUE/FALSE

Value

dmcSim

The function returns a list with the relevant results from the simulation. The list is accessed with `obj$name` with the the following:

<code>obj\$means</code>	Condition means for reaction time and error rate
<code>obj\$caf</code>	Accuracy per bin for compatible and incompatible trials
<code>obj\$delta</code>	Mean RT and compatibility effect per bin

obj\$sim	Individual trial data points (reaction times/error) and activation vectors from simulation
obj\$trials	Example individual trial timecourse for n compatible and incompatible trials
obj\$prms	The input parameters used in the simulation

Examples

```
# Example 1
dmc <- dmcSim(fullData = TRUE) # full data only required for activation plot (top left)
plot(dmc)
dmc <- dmcSim() # faster
plot(dmc)

# Example 2
dmc <- dmcSim(tau = 130)
plot(dmc)

# Example 3
dmc <- dmcSim(tau = 90)
plot(dmc)

# Example 4
dmc <- dmcSim(varSP = TRUE)
plot(dmc, "delta")

# Example 5
dmc <- dmcSim(tau = 130, varDR = TRUE)
plot(dmc, "caf")

# Example 6
dmc <- dmcSim(nDelta = 10, nCAF = 10)
plot(dmc)
```

dmcSims

dmcSims: Run multiple dmc simulations

Description

Run dmcSim with range of input parameters.

Usage

```
dmcSims(params, printInputArgs = FALSE, printResults = FALSE)
```

Arguments

params (list of parameters to dmcSim)
 printInputArgs Print DMC input arguments to console
 printResults Print DMC output to console

Value

list of dmcsim

Examples

```

# Example 1
params <- list(amp = seq(10, 20, 5), tau = c(50, 100, 150), nTr1 = 50000)
dmc <- dmcSims(params)
plot(dmc[[1]]) # full combination 1
plot(dmc) # delta plots for all combinations
plot(dmc[c(1:3)]) # delta plots for specific combinations

# Example 2
params <- list(amp = seq(10, 20, 5), tau = seq(20, 40, 20), bnds = seq(50, 100, 25))
dmc <- dmcSims(params)
plot(dmc[[1]]) # combination 1
plot(dmc, ncol = 2) # delta plots for all combinations
plot(dmc[c(1:3)]) # delta plots for specific combinations

```

errDist

errDist: Create random binary vector

Description

Returns a random vector of 0's (correct) and 1's (incorrect) with defined proportions (default = 10% errors).

Usage

```
errDist(n = 10000, proportion = 10)
```

Arguments

n Number
 proportion Approximate proportion of errors in percentage

Value

double

Examples

```
# Example 1
x <- errDist(1000, 10)
table(x)
```

flankerData	<i>A summarised dataset: see raw data file flankerDataRaw and dmcObservedData.R This is the summarised Flanker Task data from Ulrich et al. (2015)</i>
-------------	--

Description

- \$summary → Reaction time correct, standard deviation correct, percentage error, reaction time incorrect, and standard deviation for incorrect trials for both compatible and incompatible trials
- \$scf → Proportion correct for compatible and incompatible trials across 5 bins
- \$delta → Compatible reaction times, incompatible mean reaction times, mean reaction times, incompatible - compatible reaction times (delta), and standard deviation + standard error of this difference across 10 bins.

Usage

```
flankerData
```

Format

```
dmcob
```

flankerDataRaw	<i>Raw flanker data from Ulrich et al. (2015)</i>
----------------	---

Description

- Subject Subject number
- Comp comp vs. incomp
- RT
- Error 0 = correct, 1 = error

Usage

```
flankerDataRaw
```

mean.dmcfit	<i>mean.dmcfit: Return mean simulation results from dmcFitSubject</i>
-------------	---

Description

Aggregated simulation results from dmcFitSubject.

Usage

```
## S3 method for class 'dmcfit'  
mean(x, ...)
```

Arguments

x	Output from dmcFitSubject
...	pars

Value

dmcfit

The function returns a list with the relevant aggregated results dmcFitSubject. The list is accessed with obj\$name and so on with the the following:

obj\$means	means
obj\$delta	delta
obj\$caf	caf
obj\$prms	par

Examples

```
# Example 1: Fit individual data then aggregate  
fitSubjects <- dmcFitSubject(flankerData, nTrl = 1000, subjects = c(1, 2))  
plot(fitSubjects, flankerData, subject = 1)  
summary(fitSubjects)  
fitAgg <- mean(fitSubjects)  
plot(fitAgg, flankerData)
```

plot.dmcfit *plot.dmcfit: Plot observed + fitted data*

Description

Plot the simulation results from the output of dmcFit. The plot can be an overall summary, or individual plots (activation, trials, pdf, cdf, caf, delta, all). Plot type summary1 contains an activation plot, example individual trials, the probability distribution function (PDF), the cumulative distribution function (CDF), the conditional accuracy function (CAF) and delta plots. This required that dmcSim is run with fullData = TRUE. Plot type summary2 contains only the PDF, CDF, CAF and delta plots and does not require that dmcSim is run with fullData = TRUE.

Usage

```
## S3 method for class 'dmcfit'
plot(
  x,
  y,
  subject = NULL,
  figType = "summary",
  legend = TRUE,
  labels = c("Compatible", "Incompatible", "Observed", "Predicted"),
  cols = c("black", "green", "red"),
  ylimRt = c(200, 800),
  ylimEr = c(0, 20),
  ylimCAF = c(0, 1),
  cafBinLabels = FALSE,
  ylimDelta = c(-50, 100),
  xlimDelta = c(200, 1000),
  xlabs = TRUE,
  ylabs = TRUE,
  xaxts = TRUE,
  yaxts = TRUE,
  resetPar = TRUE,
  ...
)
```

Arguments

x	Output from dmcFit
y	Observed data
subject	NULL (aggregated data across all subjects) or integer for subject number
figType	summary, rtCorrect, errorRate, rtErrors, cdf, caf, delta, all
legend	TRUE/FALSE (or FUNCTION) plot legend on each plot
labels	Condition labels c("Compatible", "Incompatible", "Observed", "Predicted") default

cols	Condition colours c("green", "red") default
ylimRt	ylim for Rt plots
ylimEr	ylim for error rate plots
ylimCAF	ylim for CAF plot
cafBinLabels	TRUE/FALSE
ylimDelta	ylim for delta plot
xlimDelta	xlim for delta plot
xlabs	TRUE/FALSE
ylabs	TRUE/FALSE
xaxts	TRUE/FALSE
yaxts	TRUE/FALSE
resetPar	TRUE/FALSE Reset graphical parameters
...	additional plot pars

Examples

```
# Example 1
resTh <- dmcFit(flankerData, nTr1 = 5000)
plot(resTh, flankerData)

# Example 2
resTh <- dmcFit(flankerData, nTr1 = 5000)
plot(resTh, flankerData)
plot(resTh, flankerData, figType = "all")

# Example 3
resTh <- dmcFit(simonData, nTr1 = 5000)
plot(resTh, simonData)
```

plot.dmclist

plot.dmclist: Plot delta plots from multiple dmc simulations.

Description

Plot delta function from multiple dmc simulations (i.e., dmcSims).

Usage

```
## S3 method for class 'dmclist'
plot(
  x,
  ylim = c(-50, 150),
  xlim = NULL,
  col = c("black", "lightgrey"),
  lineType = "l",
  legendPos = "topleft",
  ncol = 1,
  ...
)
```

Arguments

x	Output from dmcSims
ylim	ylimit for delta plot
xlim	xlimit for delta plot
col	# color range start/end color
lineType	line type ("l", "b", "o") for delta plot
legendPos	legend position
ncol	number of legend columns
...	pars for legend

Examples

```
# Example 1
params <- list(amp = seq(20, 30, 2))
dmc <- dmcSims(params)
plot(dmc, ncol = 2, xlim = c(0, 1300), ylim = c(-100, 200))

# Example 2
params <- list(amp=c(10, 20), tau = seq(20, 80, 40), drc = seq(0.2, 0.6, 0.2), nTr1 = 50000)
dmc <- dmcSims(params)
plot(dmc, ncol = 2, col=c("green", "blue"), lineType = "l")
```

plot.dmcob

plot.dmcob: Plot observed data

Description

Plot results from the output of dmcObservedData. The plot can be an overall summary, or individual plots (rtCorrect, errorRate, rtErrors, cdf, caf, delta, all).

Usage

```
## S3 method for class 'dmcob'
plot(
  x,
  figType = "summary",
  subject = NULL,
  legend = TRUE,
  labels = c("Compatible", "Incompatible"),
  cols = c("black", "green", "red"),
  errorBars = FALSE,
  errorBarType = "sd",
  ylimRt = c(200, 800),
  ylimEr = c(0, 20),
  ylimCAF = c(0, 1),
  cafBinLabels = FALSE,
  ylimDelta = c(-50, 100),
  xlimDelta = c(200, 1000),
  xlabs = TRUE,
  ylabs = TRUE,
  xaxts = TRUE,
  yaxts = TRUE,
  resetPar = TRUE,
  ...
)
```

Arguments

x	Output from dmcObservedData
figType	summary, rtCorrect, errorRate, rtErrors, cdf, caf, delta, all
subject	NULL (aggregated data across all subjects) or integer for subject number
legend	TRUE/FALSE (or FUNCTION) plot legend on each plot
labels	Condition labels c("Compatible", "Incompatible") default
cols	Condition colours c("green", "red") default
errorBars	TRUE(default)/FALSE Plot errorbars
errorBarType	sd(default), or se
ylimRt	ylimit for Rt plots
ylimEr	ylimit for error rate plots
ylimCAF	ylimit for CAF plot
cafBinLabels	TRUE/FALSE
ylimDelta	ylimit for delta plot
xlimDelta	xlimit for delta plot
xlabs	TRUE/FALSE
ylabs	TRUE/FALSE

xaxts	TRUE/FALSE
yaxts	TRUE/FALSE
resetPar	TRUE/FALSE Reset graphical parameters
...	additional plot pars

Examples

```
# Example 1 (real dataset)
plot(flankerData)
plot(flankerData, errorBars = TRUE, errorBarType = "se")
plot(flankerData, figType = "delta")
plot(flankerData, figType = "caf")

# Example 2 (real dataset)
plot(simonData)
plot(simonData, errorBars = TRUE, errorBarType = "se")
plot(simonData, figType = "delta", errorBars = TRUE, errorBarType = "sd")

# Example 3 (simulated dataset)
dat <- createDF(nSubjects = 50, nTr1 = 50,
               design = list("Comp" = c("comp", "incomp")))
dat <- addDataDF(dat,
                RT = list("Comp_comp" = c(420, 100, 80),
                          "Comp_incomp" = c(470, 100, 95)),
                Error = list("Comp_comp" = c(5, 3, 2, 1, 2),
                              "Comp_incomp" = c(15, 8, 4, 2, 2)))
datOb <- dmcObservedData(dat)
plot(datOb, errorBars = TRUE, errorBarType = "sd")

# Example 4 (simulated dataset)
dat <- createDF(nSubjects = 50, nTr1 = 50,
               design = list("Comp" = c("comp", "incomp")))
dat <- addDataDF(dat,
                RT = list("Comp_comp" = c(420, 100, 150),
                          "Comp_incomp" = c(470, 100, 120)),
                Error = list("Comp_comp" = c(5, 3, 2, 1),
                              "Comp_incomp" = c(15, 8, 4, 2)))
datOb <- dmcObservedData(dat, nCAF = 4)
plot(datOb)
```

plot.dmcobs

plot.dmcobs: Plot combined observed data

Description

Plot results from the output of `dmcObservedData`. The plot can be an overall `rtCorrect`, `errorRate`, `rtErrors`, `cdf`, `caf`, `delta`, or all of the previous plots.

Usage

```
## S3 method for class 'dmcobs'
plot(
  x,
  figType = "all",
  subject = NULL,
  legend = TRUE,
  labels = c("Compatible", "Incompatible"),
  cols = c("black", "green", "red"),
  ltys = c(1, 1),
  pchs = c(1, 1),
  errorBars = FALSE,
  errorBarType = "sd",
  ylimRt = c(200, 800),
  ylimEr = c(0, 20),
  ylimCAF = c(0, 1),
  cafBinLabels = FALSE,
  ylimDelta = c(-50, 100),
  xlimDelta = c(200, 1000),
  xlabs = TRUE,
  ylabs = TRUE,
  xaxts = TRUE,
  yaxts = TRUE,
  resetPar = TRUE,
  ...
)
```

Arguments

x	Output from dmcObservedData
figType	rtCorrect, errorRate, rtErrors, cdf, caf, delta, all
subject	NULL (aggregated data across all subjects) or integer for subject number
legend	TRUE/FALSE (or FUNCTION) plot legend on each plot
labels	Condition labels c("Compatible", "Incompatible") default
cols	Condition colours c("green", "red") default
ltys	Linetype see par
pchs	Symbols see par
errorBars	TRUE(default)/FALSE Plot errorbars
errorBarType	sd(default), or se
ylimRt	ylimit for Rt plots
ylimEr	ylimit for error rate plots
ylimCAF	ylimit for CAF plot
cafBinLabels	TRUE/FALSE
ylimDelta	ylimit for delta plot

xlimDelta	xlimit for delta plot
xlabs	TRUE/FALSE
ylabs	TRUE/FALSE
xaxts	TRUE/FALSE
yaxts	TRUE/FALSE
resetPar	TRUE/FALSE Reset graphical parameters
...	additional plot pars

Examples

```
# Example 1
datFlanker <- dmcObservedData(flankerDataRow, nDelta = 9)
datSimon <- dmcObservedData(simonDataRow, nDelta = 9)
dat <- dmcCombineObservedData(datFlanker, datSimon) # combine flanker/simon data
plot(dat, figType = "delta", xlimDelta = c(200, 700),
      cols = c("black", "darkgrey"), pchs = c(1, 2), resetPar = FALSE)
legend(200, 0, legend = c("Flanker Task", "Simon Task"),
      col = c("black", "darkgrey"), lty = c(1, 1))
```

plot.dmcsim

plot.dmcsim: Plot dmc simulation

Description

Plot the simulation results from the output of dmcSim. The plot can be an overall summary, or individual plots (activation, trials, pdf, cdf, caf, delta, all). Plot type summary1 contains an activation plot, example individual trials, the probability distribution function (PDF), the cumulative distribution function (CDF), the conditional accuracy function (CAF) and delta plot. This requires that dmcSim is run with fullData = TRUE. Plot type summary2 contains only the PDF, CDF, CAF and delta plots and does not require that dmcSim is run with fullData = TRUE.

Usage

```
## S3 method for class 'dmcsim'
plot(
  x,
  figType = "summary1",
  ylimCAF = c(0, 1),
  cafBinLabels = FALSE,
  ylimDelta = c(-50, 150),
  xlimDelta = NULL,
  ylimRt = c(200, 800),
  ylimErr = c(0, 20),
```

```

legend = TRUE,
labels = c("Compatible", "Incompatible"),
cols = c("black", "green", "red"),
errorBars = FALSE,
xlabs = TRUE,
ylabs = TRUE,
xaxts = TRUE,
yaxts = TRUE,
resetPar = TRUE,
...
)

```

Arguments

x	Output from dmcSim
figType	summary1, summary2, summary3, activation, trials, pdf, cdf, caf, delta, rtCorrect, rtErrors, errorRate, all
ylimCAF	ylimit for CAF plot
cafBinLabels	TRUE/FALSE
ylimDelta	ylimit for delta plot
xlimDelta	xlimit for delta plot (Default is 0 to tmax)
ylimRt	ylimit for rt plot
ylimErr	ylimit for er plot
legend	TRUE/FALSE (or FUNCTION) plot legend on each plot
labels	Condition labels c("Compatible", "Incompatible") default
cols	Condition colours c("green", "red") default
errorBars	TRUE/FALSE
xlabs	TRUE/FALSE
ylabs	TRUE/FALSE
xaxts	TRUE/FALSE
yaxts	TRUE/FALSE
resetPar	TRUE/FALSE Reset graphical parameters
...	additional plot pars

Examples

```

# Example 1
dmc = dmcSim(fullData = TRUE)
plot(dmc)

# Example 2
dmc = dmcSim()
plot(dmc)

```



```
# Example 3
dmc = dmcSim()
plot(dmc, figType = "all")

# Example 4
dmc = dmcSim()
plot(dmc, figType = "summary3")
```

rtDist

rtDist: Create random RT distribution

Description

Returns value(s) from a distribution appropriate to simulate reaction times. The distribution is a combined exponential and gaussian distribution called an exponentially modified Gaussian (EMG) distribution or ex-gaussian distribution.

Usage

```
rtDist(n = 10000, gaussMean = 600, gaussSD = 50, expRate = 200)
```

Arguments

n	Number of observations
gaussMean	Mean of the gaussian distribution
gaussSD	SD of the gaussian distribution
expRate	Rate of the exponential function

Value

double

Examples

```
# Example 1
x <- rtDist()
hist(x, 100)

# Example 2
x <- rtDist(n=20000, gaussMean=800, gaussSD=50, expRate=100)
hist(x, 100)
```

simonData	<i>A summarised dataset: see raw data file simonDataRaw and dmcObservedData.R This is the summarised Simon Task data from Ulrich et al. (2015)</i>
-----------	--

Description

- \$summary → Reaction time correct, standard deviation correct, percentage error, reaction time incorrect, and standard deviation for incorrect trials for both compatible and incompatible trials
- \$scf → Proportion correct for compatible and incompatible trials across 5 bins
- \$delta → Compatible reaction times, incompatible mean reaction times, mean reaction times, incompatible - compatible reaction times (delta), and standard deviation + standard error of this difference across 10 bins.

Usage

```
simonData
```

Format

```
dmcob
```

simonDataRaw	<i>Raw simon data from Ulrich et al. (2015)</i>
--------------	---

Description

- Subject Subject number
- Comp comp vs. incomp
- RT
- Error 0 = correct, 1 = error

Usage

```
simonDataRaw
```

summary.dmcfit	<i>summary.dmcfit: dmc fit aggregate summary</i>
----------------	--

Description

Summary of the simulation results from dmcFitAgg

Usage

```
## S3 method for class 'dmcfit'  
summary(object, digits = 2, ...)
```

Arguments

object	Output from dmcFitAgg
digits	Number of digits in the output
...	pars

Value

DataFrame

Examples

```
# Example 1  
fitAgg <- dmcFit(flankerData, nTr1 = 1000)  
summary(fitAgg)
```

summary.dmcsim	<i>summary.dmcsim: dmc simulation summary</i>
----------------	---

Description

Summary of the overall results from dmcSim

Usage

```
## S3 method for class 'dmcsim'  
summary(object, digits = 1, ...)
```

Arguments

object	Output from dmcSim
digits	Number of digits in the output
...	pars

Value

list

Examples

```
# Example 1
dmc <- dmcSim()
summary(dmc)

# Example 2
dmc <- dmcSim(tau = 90)
summary(dmc)
```

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