

# Package: dcm2 (via r-universe)

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**Type** Package

**Title** Calculating the M2 Model Fit Statistic for Diagnostic Classification Models

**Version** 1.0.2.9000

**Description** A collection of functions for calculating the M2 model fit statistic for diagnostic classification models as described by Liu et al. (2016) <[DOI:10.3102/1076998615621293](https://doi.org/10.3102/1076998615621293)>. These functions provide multiple sources of information for model fit according to the M2 statistic, including the M2 statistic, the \*p\* value for that M2 statistic, and the Root Mean Square Error of Approximation based on the M2 statistic.

**License** GPL-3

**URL** <https://github.com/atlas-aai/dcm2>

**BugReports** <https://github.com/atlas-aai/dcm2/issues>

**Depends** R (>= 3.6)

**Imports** dplyr (>= 0.8.4), glue (>= 1.4.2), magrittr (>= 1.5), methods (>= 4.1.0), modelr (>= 0.1.8), purrr (>= 0.3.3), Rcpp, rlang (>= 0.4.11), stringr (>= 1.4.0), tibble (>= 2.1.3), tidyr (>= 1.1.4)

**Suggests** covr, GDINA, knitr, rmarkdown, roxygen2, spelling, testthat (>= 2.1.0)

**LinkingTo** Rcpp, RcppArmadillo (>= 0.9.800.1.0)

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**LazyData** true

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**RoxygenNote** 7.2.3

**Language** en-US

**NeedsCompilation** yes

**VignetteBuilder** knitr

**Repository** <https://atlas-aai.r-universe.dev>

**RemoteUrl** <https://github.com/atlas-aai/dcm2>

**RemoteRef** HEAD

**RemoteSha** b156764ddda3995b1945c671a67f6645268e39eb

## Contents

as_binary . . . . .	2
calc_m2 . . . . .	3
data_att1 . . . . .	4
fit_m2 . . . . .	5
log_odds . . . . .	6
sample_data . . . . .	7

<b>Index</b>	<b>8</b>
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as_binary	<i>Make Binary Profiles</i>
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### Description

Given a number of attributes, as\_binary will create all possible binary mastery profiles.

### Usage

```
as_binary(x)
```

### Arguments

x	The number of attributes
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### Value

A  $2^x$  by x matrix

### Examples

```
as_binary(3)
as_binary(4)
```

---

 calc\_m2

*Calculate the M2*


---

### Description

Calculate the M2

### Usage

```
calc_m2(
  data,
  struc_params,
  pi_matrix,
  qmatrix,
  ci = 0.9,
  link = "logit",
  model_type = c("LCDM", "GDINA", "ACDM", "LLM", "RRUM", "DINO", "DINA", "BUGDINO")
)
```

### Arguments

data	A data frame containing the raw data, where there is one row per respondent and one column per item
struc_params	A vector containing the structural parameters of the estimated model
pi_matrix	An item-by-class matrix containing the probability of a correct response by members of each latent class
qmatrix	A data frame containing the Q-matrix
ci	The confidence interval for the RMSEA, computed from the M2
link	A character containing the link function.
model_type	A character containing the model type (e.g., LCDM) that was estimated.

### Value

A data frame containing:

- m2: The M2 statistic
- df: Degrees of freedom for the M2 statistic
- pval:  $p$ -value for the M2 statistic
- rmsea: Root mean square error of approximation
- ci\_lower: Lower end of ci interval for RMSEA
- ci\_upper: Upper end of ci interval for RMSEA
- srmsr: Standardized root mean square residual

## Examples

```
possible_prof <- dcm2::as_binary(ncol(sample_data$q_matrix))

fit_dat <- sample_data$data %>%
  tidyr::pivot_wider(names_from = "item_id",
                    values_from = "score") %>%
  dplyr::select(-"resp_id") %>%
  as.matrix() %>%
  unname()
gdina_mod <- GDINA::GDINA(dat = fit_dat,
                        Q = data.frame(sample_data$q_matrix),
                        model = "logitGDINA",
                        control = list(conv.type = "neg2LL"))
struc_params <- gdina_mod$struc.parm
pi_matrix <- gdina_mod$LC.prob %>%
  as.matrix() %>%
  unname()
calc_m2(data = fit_dat, struc_params, pi_matrix,
        qmatrix = data.frame(sample_data$q_matrix), ci = 0.9, link = "logit",
        model_type = "LCDM")
```

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data\_att1

*Simulated Data for a Single Attribute Assessment*

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## Description

A list containing data from a randomly simulated single-attribute assessment.

## Usage

```
data_att1
```

## Format

A list frame containing 4 tibble objects:

- `resp_profiles`: A tibble with 1000 rows and 2 columns. The first column indicates `resp_id` (i.e., the respondent identification number) and the second column indicates `att_1` (i.e., a binary indicator for whether the respondent mastered the first attribute).
- `q_matrix`: A tibble with 2 rows and 1 column. Each row corresponds to an assessment item, and the column entries provide a binary indicator for whether the item assessed the attribute.
- `item_params`: A tibble with 2 rows and 3 columns. Each row corresponds to an item. The first column indicates `item_id` (i.e., the item identification number). The second column indicates `intercept` (i.e., the true item intercept parameter for the item). The third column indicates `att_1` (i.e., the true item main effect parameter for the item).

- data: A tibble with 2000 rows and 3 columns. The first column indicates resp\_id (i.e., the respondent identification number). The second column indicates item\_id (i.e., the item identification number). The third column indicates score (i.e., the dichotomously scored item response).

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fit\_m2

---

*Model Fit M2 Calculations*


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### Description

Estimate the M2 statistic as described by Liu et al. (2016).

### Usage

```
fit_m2(model, ci = 0.9, ...)
```

### Arguments

model	An estimated diagnostic classification model.
ci	The confidence interval for the RMSEA.
...	Unused, for extensibility.

### Value

A data frame containing:

- m2: The M2 statistic
- df: Degrees of freedom for the M2 statistic
- pval:  $p$ -value for the M2 statistic
- rmsea: Root mean square error of approximation
- ci\_lower: Lower end of ci interval for RMSEA
- ci\_upper: Upper end of ci interval for RMSEA
- srmsr: Standardized root mean square residual

### References

Liu, Y., Tian, W., & Xin, T. (2016). An application of  $M_2$  statistic to evaluate the fit of cognitive diagnostic models. *Journal of Educational and Behavioral Statistics*, 41, 3-26. doi:10.3102/1076998615621293

**Examples**

```
possible_prof <- dcm2::as_binary(ncol(sample_data$q_matrix))

fit_dat <- sample_data$data %>%
  tidyr::pivot_wider(names_from = "item_id",
                    values_from = "score") %>%
  dplyr::select(-"resp_id") %>%
  as.matrix() %>%
  unname()
gdina_mod <- GDINA::GDINA(dat = fit_dat,
                        Q = data.frame(sample_data$q_matrix),
                        model = "logitGDINA",
                        control = list(conv.type = "neg2LL"))
fit_m2(gdina_mod, ci = 0.9)
```

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log\_odds

*Log-odds Transformation*


---

**Description**

These functions implement the log-odds (or logit) transformation. This is a common transformation for psychometric models that is used to put probabilities on a continuous scale.

**Usage**

```
logit(x)

inv_logit(x)
```

**Arguments**

x                    A number to be transformed

**Value**

A transformed double

**Examples**

```
logit(0.6)
logit(0.5)

inv_logit(3.5)
inv_logit(0)
```

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`sample_data`*Simulated Data for Testing Functions*

---

**Description**

A matrix with randomly simulated data to test the package functions.

**Usage**`sample_data`**Format**

A list frame containing 4 tibble objects:

- `resp_profiles`: A tibble with 1000 rows and 3 columns. The first column indicates `resp_id` (i.e., the respondent identification number). The second column indicates `att_1` (i.e., a binary indicator for whether the respondent mastered the first attribute). The third column indicates `att_2` (i.e., a binary indicator for whether the respondent mastered the second attribute).
- `q_matrix`: A tibble with 8 rows and 2 columns. Each row corresponds to an assessment item, and the column entries provide a binary indicator for whether the item assessed each of the attribute.
- `item_params`: A tibble with 8 rows and 5 columns. Each row corresponds to an item. The first column indicates `item_id` (i.e., the item identification number). The second column indicates `intercept` (i.e., the true item intercept parameter for the item). The third column indicates `att_1` (i.e., the true item main effect parameter for the first attribute for the item). The fourth column indicates `att_2` (i.e., the true item main effect parameter for the second attribute for the item). The fifth column indicates `att_1__att_2` (i.e., the true item interaction effect parameter for the first and second attributes).
- `data`: A tibble with 8000 rows and 3 columns. The first column indicates `resp_id` (i.e., the respondent identification number). The second column indicates `item_id` (i.e., the item identification number). The third column indicates `score` (i.e., the dichotomously scored item response).

# Index

## \* datasets

data\_att1, 4

sample\_data, 7

as\_binary, 2

calc\_m2, 3

data\_att1, 4

fit\_m2, 5

inv\_logit(log\_odds), 6

log\_odds, 6

logit(log\_odds), 6

sample\_data, 7