

# Package ‘EMLI’

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**Type** Package

**Title** Efficient Maximum Likelihood Identification

**Version** 0.1.0

**Description** Provides implementations of computationally efficient maximum likelihood estimation algorithms for system identification tasks. Currently, one such algorithm is implemented which identifies the one-dimensional cumulative structural equation model with normality assumptions. The corresponding scientific paper is yet to be published, therefore the relevant reference is not available yet.

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**Imports** stats

**Encoding** UTF-8

**RoxygenNote** 7.1.2

**NeedsCompilation** no

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**evaluate\_identification**  
*evaluate\_identification*

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## Description

Calculates a discrepancy function based metric of estimation accuracy for the one-dimensional cumulative structural equation model with normality assumptions. Suitable when there are no contradictions in the factuals/estimates.

## Usage

```
evaluate_identification(f, e, n)
```

## Arguments

- f A list consisting of 3 elements: 1) the factual Sigma ((m + 1) x (m + 1) matrix of finite numeric elements); 2) the factual sigma\_y^2 (vector of length 1, finite numeric element); 3) the factual mu ((m + 1) x 1 matrix of finite numeric elements).
- e Analogous to parameter f but with estimates instead of factuals.
- n The number of time moments used for obtaining parameter e (vector of length 1, finite positive integer).

## Value

Calculated metric value (vector of length 1, numeric element). The lower the value, the better the accuracy, with 0 indicating perfect accuracy.

## Examples

```
set.seed(1)

m <- 4
k <- 2

L <- matrix(runif((m + 1) * k, min = -10, max = 10), nrow = m + 1)
sigma <- matrix(runif(m + 2, min = 0, max = 10), nrow = m + 2)
mu <- matrix(runif(m + 1, min = -10, max = 10), nrow = m + 1)

n <- 100
data <- generate_data(n, L, sigma, mu)

Sigma <- L %*% t(L) + diag(sigma[1:(m + 1), ] ^ 2)
sigma_y_squared <- sigma[m + 2, ] ^ 2
Sigma[m + 1, m + 1] <- Sigma[m + 1, m + 1] + 2 * sigma_y_squared

factual_parameters <- list(Sigma, sigma_y_squared, mu)
```

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```
estimated_parameters <- identify_model(data[[1]], data[[2]], 0.00001)
evaluate_identification(factual_parameters, estimated_parameters, n)
```

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**generate\_data***generate\_data***Description**

Generates data according to the one-dimensional cumulative structural equation model with normality assumptions with given model parameter values.

**Usage**

```
generate_data(n, L, sigma, mu)
```

**Arguments**

<b>n</b>	The number of time moments to generate the data for (vector of length 1, finite positive integer).
<b>L</b>	Factor loadings ((m + 1) x k matrix of finite numeric elements: the first m rows correspond to the input measurement equation; the last row corresponds to the transition equation).
<b>sigma</b>	Standard deviations of the error/noise terms ((m + 2) x 1 matrix of finite non-negative numeric elements: the first m rows correspond to the input measurement equation; the row before the last one corresponds to the transition equation; the last row corresponds to the output measurement equation).
<b>mu</b>	Intercept terms ((m + 1) x 1 matrix of finite numeric elements; the first m rows correspond to the input measurement equation; the last row corresponds to the transition equation).

**Value**

A list consisting of 2 elements: 1) observed input data (n x m matrix of numeric elements); 2) observed output differences data (n x 1 matrix of numeric elements).

**Examples**

```
set.seed(1)

m <- 4
k <- 2

L <- matrix(runif((m + 1) * k, min = -10, max = 10), nrow = m + 1)
sigma <- matrix(runif(m + 2, min = 0, max = 10), nrow = m + 2)
mu <- matrix(runif(m + 1, min = -10, max = 10), nrow = m + 1)
generate_data(10, L, sigma, mu)
```

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<code>identify_model</code>	<i>identify_model</i>
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## Description

Calculates maximum likelihood estimates of the statistical parameters of the one-dimensional cumulative structural equation model with normality assumptions.

## Usage

```
identify_model(x, dy, tol)
```

## Arguments

<code>x</code>	Observed input data (n x m matrix of finite numeric elements).
<code>dy</code>	Observed output differences data (n x 1 matrix of finite numeric elements).
<code>tol</code>	A tolerance parameter of the golden section search algorithm used for minimizing the one-dimensional likelihood function (vector of length 1, finite positive numeric element).

## Value

A list consisting of 3 elements: 1) estimate of the covariance of cbind(x, dy) at lag 0 (`Sigma`; (m + 1) x (m + 1) matrix of numeric elements); 2) estimate of the only non-zero element of the negative covariance of cbind(x, dy) at lag 1 (`sigma_y^2`; vector of length 1, numeric element); 3) estimate of the mean of cbind(x, dy) (`mu`; (m + 1) x 1 matrix of numeric elements).

## Examples

```
set.seed(1)

m <- 4
k <- 2

L <- matrix(runif((m + 1) * k, min = -10, max = 10), nrow = m + 1)
sigma <- matrix(runif(m + 2, min = 0, max = 10), nrow = m + 2)
mu <- matrix(runif(m + 1, min = -10, max = 10), nrow = m + 1)

data <- generate_data(100, L, sigma, mu)

identify_model(data[[1]], data[[2]], 0.00001)
```

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