Package 'optimalcausalities'

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Title Optimal Stochastic Interventions in High-dimensional Data

Version 2.0

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Description Description here.

Depends R (>= 3.3.3)

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Encoding UTF-8

LazyData true

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Imports Rsolnp, keyATM

RoxygenNote 7.1.1

R topics documented:

analyze_fixedStrategy *analyze_fixedStrategy*

Description

Implements ...

Usage

```
analyze_fixedStrategy(
  specifiedAssignmentMechanism = NULL,
  hypotheticalAssignmentMechanism = NULL
)
```

Arguments

dfm	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke Imai, Rohit, Connor

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

computeQse_conjoint *computeQse_conjoint*

Description

Implements ...

Usage

```
computeQse_conjoint(
  FactorsMat,
  Yobs,
  hypotheticalProbList,
  assignmentProbList,
  log_pr_w = NULL,
  hajek = T,
  returnLog = T,
  log_treatment_combs
)
```

Arguments

dfm	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke, Rohit, Connor. Working Paper.

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

---

computeQse_lda *computeQse_lda*

Description

Implements ...

Usage

```
computeQse_lda(
  THETA__,
  INDICES_,
  DOC_INDICES_U,
  D_INDICES_U,
  PI_MAT_INPUT,
  MARGINAL_BOUNDS,
  DOC_LIST,
```

```
  MODAL_DOC_LEN,
  TERMS_MAT_INPUT,
  LOG_TREATCOMBS,
  YOBS,
  returnLog = T,
  LOG_PR_W = NULL
)
```

Arguments

dfm	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke, Rohit, Connor. Working Paper.

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

`computeQ_conjoint` *computeQ_conjoint*

Description

Implements ...

Usage

```
computeQ_conjoint(
  FactorsMat,
  Yobs,
  hypotheticalProbList,
  assignmentProbList,
  log_pr_w = NULL,
  hajek = T
)
```

Arguments

dfm	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke, Rohit, Connor. Working Paper.

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

`computeQ_lda` *computeQ_lda*

Description

Implements ...

Usage

```
computeQ_lda(
  theta = NULL,
  term_mat,
  Yobs,
  doc_words,
  dtm = NULL,
  pi_mat = NULL,
  alpha_mat = NULL,
  log_pr_w = NULL,
  computeSE = F,
  trim_q = 1,
  quiet = T,
  iters = 100,
  smoothWts = F,
  TreatFxn = NULL,
  maxWt = 1e+10,
  maxWt_hajek = NULL,
  term_mat_TRUE = NULL,
  doc_indices_u = NULL,
  d_indices_u = NULL,
  diagnostics = F
)
```

Arguments

`dfm`	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke, Rohit, Connor. Working Paper.

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

`find_optimalStrategy` *find_optimalStrategy*

Description

Implements ...

Usage

```
find_optimalStrategy(specifiedAssignmentMechanism = NULL)
```

Arguments

`dfm`	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke Imai, Rohit, Connor

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

optimizeQ_conjoint *computeQ_conjoint*

Description

Implements ...

Usage

```
optimizeQ_conjoint(
  FactorsMat,
  Yobs,
  assignmentProbList,
  se_ub,
  INDICES_SPLIT1,
  INDICES_SPLIT2 = NULL,
  computeSEs = F,
  hajek = T,
  doMax = T,
  quiet = T
)
```

Arguments

dfm	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke Imai, Rohit, Connor

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

optimizeQ_lda *computeQ_lda*

Description

Implements ...

Usage

```
optimizeQ_lda(
  INDICES_SPLIT1 = NULL,
  INDICES_SPLIT2 = NULL,
  DTM_MAT,
  n_fold = 3,
  YOBS,
  PI_MAT,
  DOC_LIST,
  TERMS_MAT,
  SE_UB = sd(YOBS)/10,
  nboot = 10,
  trim_q = 1,
  maxWt = 1e+10,
  maxWt_hajek = NULL,
  computeSEs = T,
  doMax = T,
  alphaLevel = 0.05,
  openBrowser = F
)
```

Arguments

dfm	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke Imai, Rohit, Connor

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

`specify_treatmentMechanism` *specify*

Description

Implements ...

Usage

```
specify_treatmentMechanism(Yobs, W, PrW_parameters = list())
```

Arguments

`dfm`	'document-feature matrix'. A list ...

Value

A list consiting of

- Items.

References

- Kosuke Imai, Rohit, Connor

Examples

```
#set seed
set.seed(1)

#Geneate data
x <- rnorm(100)
```

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