

Package ‘dcTensor’

July 7, 2023

Type Package

Title Discrete Matrix/Tensor Decomposition

Version 1.2.0

Depends R (>= 3.4.0)

Imports methods, MASS, fields, rTensor, nnTensor

Suggests knitr, rmarkdown, testthat

Description Semi-Binary and Semi-Ternary Matrix Decomposition are performed based on Non-negative Matrix Factorization (NMF) and Singular Value Decomposition (SVD). For the details of the methods, see the reference section of GitHub README.md <<https://github.com/rikenbit/dcTensor>>.

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URL <https://github.com/rikenbit/dcTensor>

VignetteBuilder knitr

NeedsCompilation no

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Repository CRAN

Date/Publication 2023-07-07 08:00:02 UTC

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Description

Semi-Binary and Semi-Ternary Matrix Decomposition are performed based on Non-negative Matrix Factorization (NMF) and Singular Value Decomposition (SVD). For the details of the methods, see the reference section of GitHub README.md <<https://github.com/rikenbit/dcTensor>>.

Details

The DESCRIPTION file:

Package:	dcTensor
Type:	Package
Title:	Discrete Matrix/Tensor Decomposition
Version:	1.2.0
Authors@R:	c(person("Koki", "Tsuyuzaki", role = c("aut", "cre"), email = "k.t.the-answer@hotmail.co.jp"))
Depends:	R (>= 3.4.0)
Imports:	methods, MASS, fields, rTensor, nnTensor
Suggests:	knitr, rmarkdown, testthat
Description:	Semi-Binary and Semi-Ternary Matrix Decomposition are performed based on Non-negative Matrix Factorization (NMF) and Singular Value Decomposition (SVD).
License:	MIT + file LICENSE
URL:	https://github.com/rikenbit/dcTensor
VignetteBuilder:	knitr
Author:	Koki Tsuyuzaki [aut, cre]
Maintainer:	Koki Tsuyuzaki <k.t.the-answer@hotmail.co.jp>

Index of help topics:

dNMF	Discretized Non-negative Matrix Factorization Algorithms (dNMF)
dNTD	Discretized Non-negative Tucker Decomposition Algorithms (dNTD)
dNTF	Discretized Non-negative CP Decomposition Algorithms (dNTF)
dPLS	Discretized Partial Least Squares (dPLS)
dSVD	Discretized Gradient Descent Singular Value Decomposition (dSVD)
dcTensor-package	Discrete Matrix/Tensor Decomposition
djNMF	Discretized Joint Non-negative Matrix Factorization Algorithms (djNMF)
dsiNMF	Discretized Simultaneous Non-negative Matrix Factorization Algorithms (dsiNMF)
toyModel	Toy model data for using dNMF, dSVD, dsiNMF, djNMF, dPLS, dNTF, and dNTD

Author(s)

NA

Maintainer: NA

References

Z. Zhang, T. Li, C. Ding and X. Zhang, (2007). Binary Matrix Factorization with Applications, *Seventh IEEE International Conference on Data Mining (ICDM 2007)*, 391-400

See Also

[toyModel](#), [dNMF](#), [dSVD](#)

Examples

```
ls("package:dcTensor")
```

djNMF

*Discretized Joint Non-negative Matrix Factorization Algorithms
(djNMF)*

Description

The input data objects are assumed to be non-negative matrices. djNMF decompose the matrices to two low-dimensional factor matrices simultaneously.

Usage

```
djNMF(X, M=NULL, pseudocount=.Machine$double.eps,
       initW=NULL, initV=NULL, initH=NULL,
       fixW=FALSE, fixV=FALSE, fixH=FALSE,
       Bin_W=1e-10, Bin_V=rep(1e-10, length=length(X)), Bin_H=rep(1e-10, length=length(X)),
       Ter_W=1e-10, Ter_V=rep(1e-10, length=length(X)), Ter_H=rep(1e-10, length=length(X)),
       L1_W=1e-10, L1_V=rep(1e-10, length=length(X)), L1_H=rep(1e-10, length=length(X)),
       L2_W=1e-10, L2_V=rep(1e-10, length=length(X)), L2_H=rep(1e-10, length=length(X)),
       J = 3, w=NULL, algorithm = c("Frobenius", "KL", "IS", "PLTF"), p=1,
       thr = 1e-10, num.iter = 100,
       viz = FALSE, figdir = NULL, verbose = FALSE)
```

Arguments

X	A list containing input matrices ($X_k, <N*Mk>$, $k=1..K$).
M	A list containing the mask matrices ($X_k, <N*Mk>$, $k=1..K$). If the input matrix has missing values, specify the element as 0 (otherwise 1).
pseudocount	The pseudo count to avoid zero division, when the element is zero (Default: Machine Epsilon).

initW	The initial values of factor matrix W, which has N-rows and J-columns (Default: NULL).
initV	A list containing the initial values of multiple factor matrices (V_k, <N*J>, k=1..K, Default: NULL).
initH	A list containing the initial values of multiple factor matrices (H_k, <Mk*J>, k=1..K, Default: NULL).
fixW	Whether the factor matrix W is updated in each iteration step (Default: FALSE).
fixV	Whether the factor matrices V_k are updated in each iteration step (Default: FALSE).
fixH	Whether the factor matrices H_k are updated in each iteration step (Default: FALSE).
Bin_W	Paramter for binary (0,1) regulararitation (Default: 1e-10).
Bin_V	A K-length vector containing the paramters for binary (0,1) regulararitation (Default: rep(1e-10, length=length(dim(X)))).
Bin_H	A K-length vector containing the paramters for binary (0,1) regulararitation (Default: rep(1e-10, length=length(dim(X)))).
Ter_W	Paramter for terary (0,1,2) regulararitation (Default: 1e-10).
Ter_V	A K-length vector containing the paramters for terary (0,1,2) regulararitation (Default: rep(1e-10, length=length(dim(X)))).
Ter_H	A K-length vector containing the paramters for terary (0,1,2) regulararitation (Default: rep(1e-10, length=length(dim(X)))).
L1_W	Paramter for L1 regulararitation (Default: 1e-10). This also works as small positive constant to prevent division by zero, so should be set as 0.
L1_V	A K-length vector containing the paramters for L1 regulararitation (Default: rep(1e-10, length=length(dim(X)))). This also works as small positive constant to prevent division by zero, so should be set as 0.
L1_H	A K-length vector containing the paramters for L1 regulararitation (Default: rep(1e-10, length=length(dim(X)))). This also works as small positive constant to prevent division by zero, so should be set as 0.
L2_W	Paramter for L2 regulararitation (Default: 1e-10).
L2_V	A K-length vector containing the paramters for L2 regulararitation (Default: rep(1e-10, length=length(dim(X)))).
L2_H	A K-length vector containing the paramters for L2 regulararitation (Default: rep(1e-10, length=length(dim(X)))).
J	Number of low-dimension (J < N, Mk).
w	Weight vector (Default: NULL)
algorithm	Divergence between X and X_bar. "Frobenius", "KL", and "IS" are available (Default: "KL").
p	The parameter of Probabilistic Latent Tensor Factorization (p=0: Frobenius, p=1: KL, p=2: IS)
thr	When error change rate is lower than thr, the iteration is terminated (Default: 1E-10).

<code>num. iter</code>	The number of interation step (Default: 100).
<code>viz</code>	If <code>viz == TRUE</code> , internal reconstructed matrix can be visualized.
<code>figdir</code>	the directory for saving the figure, when <code>viz == TRUE</code> .
<code>verbose</code>	If <code>verbose == TRUE</code> , Error change rate is generated in console windos.

Value

`W` : A matrix which has `N`-rows and `J`-columns ($J < N, M_k$). `V` : A list which has multiple elements containing `N`-rows and `J`-columns ($J < N, M_k$). `H` : A list which has multiple elements containing `M_k`-rows and `J`-columns matrix ($J < N, M_k$). `RecError` : The reconstruction error between data matrix and reconstructed matrix from `W` and `H`. `TrainRecError` : The reconstruction error calculated by training set (observed values specified by `M`). `TestRecError` : The reconstruction error calculated by test set (missing values specified by `M`). `RelChange` : The relative change of the error.

Author(s)

Koki Tsuyuzaki

References

- Liviu Badea, (2008) Extracting Gene Expression Profiles Common to Colon and Pancreatic Adenocarcinoma using Simultaneous nonnegative matrix factorization. *Pacific Symposium on Biocomputing* 13:279-290
- Shihua Zhang, et al. (2012) Discovery of multi-dimensional modules by integrative analysis of cancer genomic data. *Nucleic Acids Research* 40(19), 9379-9391
- Zi Yang, et al. (2016) A non-negative matrix factorization method for detecting modules in heterogeneous omics multi-modal data, *Bioinformatics* 32(1), 1-8
- Y. Kenan Yilmaz et al., (2010) Probabilistic Latent Tensor Factorization, *International Conference on Latent Variable Analysis and Signal Separation* 346-353
- N. Fujita et al., (2018) Biomarker discovery by integrated joint non-negative matrix factorization and pathway signature analyses, *Scientific Report*

Examples

```
matdata <- toyModel(model = "dsiNMF_Hard")
out <- djNMF(matdata, J=2, num. iter=2)
```

Description

The input data is assumed to be a non-negative matrix. dNMF decompose the matrix to two low-dimensional factor matices.

Usage

```
dNMF(X, M=NULL, pseudocount=.Machine$double.eps,
      initU=NULL, initV=NULL, fixU=FALSE, fixV=FALSE,
      Bin_U=1e-10, Bin_V=1e-10, Ter_U=1e-10, Ter_V=1e-10,
      L1_U=1e-10, L1_V=1e-10, L2_U=1e-10, L2_V=1e-10, J = 3,
      algorithm = c("Frobenius", "KL", "IS", "Beta"), Beta = 2,
      thr = 1e-10, num.iter = 100,
      viz = FALSE, figdir = NULL, verbose = FALSE)
```

Arguments

X	The input matrix which has N-rows and M-columns.
M	The mask matrix which has N-rows and M-columns. If the input matrix has missing values, specify the element as 0 (otherwise 1).
pseudocount	The pseudo count to avoid zero division, when the element is zero (Default: Machine Epsilon).
initU	The initial values of factor matrix U, which has N-rows and J-columns (Default: NULL).
initV	The initial values of factor matrix V, which has M-rows and J-columns (Default: NULL).
fixU	Whether the factor matrix U is updated in each iteration step (Default: FALSE).
fixV	Whether the factor matrix V is updated in each iteration step (Default: FALSE).
Bin_U	Paramter for binary (0,1) regularitartion (Default: 1e-10).
Bin_V	Paramter for binary (0,1) regularitartion (Default: 1e-10).
Ter_U	Paramter for terary (0,1,2) regularitartion (Default: 1e-10).
Ter_V	Paramter for terary (0,1,2) regularitartion (Default: 1e-10).
L1_U	Paramter for L1 regularitartion (Default: 1e-10). This also works as small positive constant to prevent division by zero, so should be set as 0.
L1_V	Paramter for L1 regularitartion (Default: 1e-10). This also works as small positive constant to prevent division by zero, so should be set as 0.
L2_U	Paramter for L2 regularitartion (Default: 1e-10).
L2_V	Paramter for L2 regularitartion (Default: 1e-10).
J	The number of low-dimension (J < N, M, Default: 3)
algorithm	dNMF algorithms. "Frobenius", "KL", "IS", and "Beta" are available (Default: "Frobenius").
Beta	The parameter of Beta-divergence.
thr	When error change rate is lower than thr, the iteration is terminated (Default: 1E-10).
num.iter	The number of interation step (Default: 100).
viz	If viz == TRUE, internal reconstructed matrix can be visualized.
figdir	The directory for saving the figure, when viz == TRUE.
verbose	If verbose == TRUE, Error change rate is generated in console window.

Value

U : A matrix which has N-rows and J-columns ($J < N, M$). V : A matrix which has M-rows and J-columns ($J < N, M$). RecError : The reconstruction error between data tensor and reconstructed tensor from U and V. TrainRecError : The reconstruction error calculated by training set (observed values specified by M). TestRecError : The reconstruction error calculated by test set (missing values specified by M). RelChange : The relative change of the error.

Author(s)

Koki Tsuyuzaki

References

Z. Zhang, T. Li, C. Ding and X. Zhang, (2007). Binary Matrix Factorization with Applications, *Seventh IEEE International Conference on Data Mining (ICDM 2007)*, 391-400

Examples

```
# Test data
matdata <- toyModel(model = "dNMF")

# Simple usage
out <- dNMF(matdata, J=5)
```

Description

The input data is assumed to be non-negative tensor. dNTD decompose the tensor to the dense core tensor (S) and low-dimensional factor matices (A).

Usage

```
dNTD(X, M=NULL, pseudocount=.Machine$double.eps,
      initS=NULL, initA=NULL, fixS=FALSE, fixA=FALSE,
      Bin_A=rep(1e-10, length=length(dim(X))),
      Ter_A=rep(1e-10, length=length(dim(X))),
      L1_A=rep(1e-10, length=length(dim(X))),
      L2_A=rep(1e-10, length=length(dim(X))),
      rank = rep(3, length=length(dim(X))),
      modes = seq_along(dim(X)),
      algorithm = c("Frobenius", "KL", "IS", "Beta"),
      init = c("dNMF", "Random"),
      Beta = 2, thr = 1e-10, num.iter = 100,
      viz = FALSE,
      figdir = NULL, verbose = FALSE)
```

Arguments

X	K-order input tensor which has I_1, I_2, ..., and I_K dimensions.
M	K-order mask tensor which has I_1, I_2, ..., and I_K dimensions. If the mask tensor has missing values, specify the element as 0 (otherwise 1).
pseudocount	The pseudo count to avoid zero division, when the element is zero (Default: Machine Epsilon).
initS	The initial values of core tensor which has I_1, I_2, ..., and I_K dimensions (Default: NULL).
initA	A list containing the initial values of K factor matrices (A_k, <I_k*J_k>, k=1..K, Default: NULL).
fixS	Whether the core tensor S is updated in each iteration step (Default: FALSE).
fixA	Whether the factor matrices Ak are updated in each iteration step (Default: FALSE).
Bin_A	A K-length vector containing the paramters for binary (0,1) regulararitation (Default: rep(1e-10, length=length(dim(X)))).
Ter_A	A K-length vector containing the paramters for terary (0,1,2) regulararitation (Default: rep(1e-10, length=length(dim(X)))).
L1_A	A K-length vector containing the paramters for L1 regularization (Default: rep(1e-10, length=length(dim(X)))). This also works as small positive constant to prevent division by zero, so should be set as 0.
L2_A	A K-length vector containing the paramters for L2 regularization (Default: rep(1e-10, length=length(dim(X)))).
rank	The number of low-dimension in each mode (Default: 3 for each mode).
modes	The vector of the modes on which to perform the decomposition (Default: 1:K <all modes>).
algorithm	dNTD algorithms. "Frobenius", "KL", "IS", and "Beta" are available (Default: "Frobenius").
init	The initialization algorithms. "NMF", "ALS", and "Random" are available (Default: "NMF").
Beta	The parameter of Beta-divergence.
thr	When error change rate is lower than thr1, the iteration is terminated (Default: 1E-10).
num.iter	The number of interation step (Default: 100).
viz	If viz == TRUE, internal reconstructed tensor can be visualized.
figdir	the directory for saving the figure, when viz == TRUE (Default: NULL).
verbose	If verbose == TRUE, Error change rate is generated in console windos.

Value

S : K-order tensor object, which is defined as S4 class of rTensor package. A : A list containing K factor matrices. RecError : The reconstruction error between data tensor and reconstructed tensor from S and A. TrainRecError : The reconstruction error calculated by training set (observed values specified by M). TestRecError : The reconstruction error calculated by test set (missing values specified by M). RelChange : The relative change of the error.

Author(s)

Koki Tsuyuzaki

References

- Yong-Deok Kim et. al., (2007). Nonnegative Tucker Decomposition. *IEEE Conference on Computer Vision and Pattern Recognition*
- Yong-Deok Kim et. al., (2008). Nonnegative Tucker Decomposition With Alpha-Divergence. *IEEE International Conference on Acoustics, Speech and Signal Processing*
- Anh Huy Phan, (2008). Fast and efficient algorithms for nonnegative Tucker decomposition. *Advances in Neural Networks - ISNN2008*
- Anh Huy Phan et. al. (2011). Extended HALS algorithm for nonnegative Tucker decomposition and its applications for multiway analysis and classification. *Neurocomputing*

See Also

[plotTensor3D](#)

Examples

```
tensorData <- toyModel(model = "dNTD")
out <- dNTD(tensorData, rank=c(2,2,2), algorithm="Frobenius",
           init="Random", num.iter=2)
```

dNTF

Discretized Non-negative CP Decomposition Algorithms (dNTF)

Description

The input data is assumed to be non-negative tensor. dNTF decompose the tensor to the diagonal core tensor (S) and low-dimensional factor matrices (A).

Usage

```
dNTF(X, M=NULL, pseudocount=.Machine$double.eps,
      initA=NULL, fixA=FALSE,
      Bin_A=rep(1e-10, length=length(dim(X))),
      Ter_A=rep(1e-10, length=length(dim(X))),
      L1_A=rep(1e-10, length=length(dim(X))),
      L2_A=rep(1e-10, length=length(dim(X))),
      rank = 3,
      algorithm = c("Frobenius", "KL", "IS", "Beta"),
      init = c("dNMF", "Random"),
      Beta = 2, thr = 1e-10, num.iter = 100, viz = FALSE, figdir = NULL,
      verbose = FALSE)
```

Arguments

X	K-order input tensor which has I_1, I_2, ..., and I_K dimensions.
M	K-order mask tensor which has I_1, I_2, ..., and I_K dimensions. If the mask tensor has missing values, specify the element as 0 (otherwise 1).
pseudocount	The pseudo count to avoid zero division, when the element is zero (Default: Machine Epsilon).
initA	A list containing the initial values of K factor matrices (A_k, $\langle I_k^* J_k \rangle$, k=1..K, Default: NULL).
fixA	Whether the factor matrices Ak are updated in each iteration step (Default: FALSE).
Bin_A	A K-length vector containing the paramters for binary (0,1) regulararitation (Default: rep(1e-10, length=length(dim(X)))).
Ter_A	A K-length vector containing the paramters for terary (0,1,2) regulararitation (Default: rep(1e-10, length=length(dim(X)))).
L1_A	A K-length vector containing the paramters for L1 regulararitation (Default: rep(1e-10, length=length(dim(X)))). This also works as small positive constant to prevent division by zero, so should be set as 0.
L2_A	A K-length vector containing the paramters for L2 regulararitation (Default: rep(1e-10, length=length(dim(X)))).
rank	The number of low-dimension in each mode (Default: 3).
algorithm	dNTF algorithms. "Frobenius", "KL", "IS", and "Beta" are available (Default: "Frobenius").
init	The initialization algorithms. "dNMF", and "Random" are available (Default: "dNMF").
Beta	The parameter of Beta-divergence.
thr	When error change rate is lower than thr1, the iteration is terminated (Default: 1E-10).
num. iter	The number of interation step (Default: 100).
viz	If viz == TRUE, internal reconstructed tensor can be visualized.
figdir	the directory for saving the figure, when viz == TRUE (Default: NULL).
verbose	If verbose == TRUE, Error change rate is generated in console windos.

Value

S : K-order tensor object, which is defined as S4 class of rTensor package. A : A list containing K factor matrices. RecError : The reconstruction error between data tensor and reconstructed tensor from S and A. TrainRecError : The reconstruction error calculated by training set (observed values specified by M). TestRecError : The reconstruction error calculated by test set (missing values specified by M). RelChange : The relative change of the error.

Author(s)

Koki Tsuyuzaki

References

- Andrzej CICHOCKI et. al., (2007). Non-negative Tensor Factorization using Alpha and Beta Divergence. *IEEE ICASSP 2007*
- Anh Huy PHAN et. al., (2008). Multi-way Nonnegative Tensor Factorization Using Fast Hierarchical Alternating Least Squares Algorithm (HALS). *NOLTA2008*
- Andrzej CICHOCKI et. al., (2008). Fast Local Algorithms for Large Scale Nonnegative Matrix and Tensor Factorizations. *IEICE Transactions on Fundamentals of Electronics, Communications and Computer Sciences*

See Also

[plotTensor3D](#)

Examples

```
tensordata <- toyModel(model = "dNTF")
out <- dNTF(tensordata, rank=3, num.ite=2)
```

dPLS

Discretized Partial Least Squares (dPLS)

Description

The input data is assumed to be a matrix. dPLS decompose the matrix to two low-dimensional factor matices.

Usage

```
dPLS(X, M=NULL, pseudocount=.Machine$double.eps,
      initV=NULL, fixV=FALSE, Ter_V=1e-10,
      L1_V=1e-10, L2_V=1e-10, eta=1e+10, J = 3,
      thr = 1e-10, num.ite = 100,
      viz = FALSE, figdir = NULL, verbose = FALSE)
```

Arguments

X	The input matrix which has N-rows and M-columns.
M	The mask matrix which has N-rows and M-columns. If the input matrix has missing values, specify the element as 0 (otherwise 1).
pseudocount	The pseudo count to avoid zero division, when the element is zero (Default: Machine Epsilon).
initV	The initial values of factor matrix V, which has M-rows and J-columns (Default: NULL).
fixV	Whether the factor matrix V is updated in each iteration step (Default: FALSE).
Ter_V	Paramter for terary (0,1,2) regularitation (Default: 1e-10).

L1_V	Paramter for L1 regularitation (Default: 1e-10). This also works as small positive constant to prevent division by zero, so should be set as 0.
L2_V	Paramter for L2 regularitation (Default: 1e-10).
eta	Stepsize of gradient descent algorithm (Default: 1e+10).
J	The number of low-dimension (J < N, M, Default: 3)
thr	When error change rate is lower than thr, the iteration is terminated (Default: 1E-10).
num.iter	The number of interation step (Default: 100).
viz	If viz == TRUE, internal reconstructed matrix can be visualized.
figdir	The directory for saving the figure, when viz == TRUE.
verbose	If verbose == TRUE, Error change rate is generated in console window.

Value

U : A matrix which has N-rows and J-columns (J < N, M). V : A matrix which has M-rows and J-columns (J < N, M). RecError : The reconstruction error between data tensor and reconstructed tensor from U and V. TrainRecError : The reconstruction error calculated by training set (observed values specified by M). TestRecError : The reconstruction error calculated by test set (missing values specified by M). RelChange : The relative change of the error.

Author(s)

Koki Tsuyuzaki

Examples

```
# Test data
matdata <- toyModel(model = "dPLS_Easy")

# Simple usage
out <- dPLS(matdata, J=2, num.iter=2)
```

Description

The input data objects are assumed to be non-negative matrices. dsiNMF decompose the matrices to two low-dimensional factor matices simultaneously.

Usage

```
dsiNMF(X, M=NULL, pseudocount=.Machine$double.eps,
        initW=NULL, initH=NULL,
        fixW=FALSE, fixH=FALSE,
        Bin_W=1e-10, Bin_H=rep(1e-10, length=length(X)),
        Ter_W=1e-10, Ter_H=rep(1e-10, length=length(X)),
        L1_W=1e-10, L1_H=rep(1e-10, length=length(X)),
        L2_W=1e-10, L2_H=rep(1e-10, length=length(X)),
        J = 3, w=NULL, algorithm = c("Frobenius", "KL", "IS", "PLTF"), p=1,
        thr = 1e-10, num.iter = 100,
        viz = FALSE, figdir = NULL, verbose = FALSE)
```

Arguments

X	A list containing the input matrices ($X_k, <N*Mk>$, $k=1..K$).
M	A list containing the mask matrices ($X_k, <N*Mk>$, $k=1..K$). If the input matrix has missing values, specify the element as 0 (otherwise 1).
pseudocount	The pseudo count to avoid zero division, when the element is zero (Default: Machine Epsilon).
initW	The initial values of factor matrix W, which has N-rows and J-columns (Default: NULL).
initH	A list containing the initial values of multiple factor matrices ($H_k, <Mk*J>$, $k=1..K$, Default: NULL).
fixW	Whether the factor matrix W is updated in each iteration step (Default: FALSE).
fixH	Whether the factor matrices H_k are updated in each iteration step (Default: FALSE).
Bin_W	Paramter for binary (0,1) regularitation (Default: 1e-10).
Bin_H	A K-length vector containing the paramters for binary (0,1) regularitation (Default: rep(1e-10, length=length(dim(X)))).
Ter_W	Paramter for terary (0,1,2) regularitation (Default: 1e-10).
Ter_H	A K-length vector containing the paramters for terary (0,1,2) regularitation (Default: rep(1e-10, length=length(dim(X)))).
L1_W	Paramter for L1 regularitation (Default: 1e-10). This also works as small positive constant to prevent division by zero, so should be set as 0.
L1_H	A K-length vector containing the paramters for L1 regularitation (Default: rep(1e-10, length=length(dim(X)))). This also works as small positive constant to prevent division by zero, so should be set as 0.
L2_W	Paramter for L2 regularitation (Default: 1e-10).
L2_H	A K-length vector containing the paramters for L2 regularitation (Default: rep(1e-10, length=length(dim(X)))).
J	Number of low-dimension ($J < N, Mk$).
w	Weight vector (Default: NULL)

algorithm	Divergence between X and X_bar. "Frobenius", "KL", and "IS" are available (Default: "KL").
p	The parameter of Probabilistic Latent Tensor Factorization (p=0: Frobenius, p=1: KL, p=2: IS)
thr	When error change rate is lower than thr, the iteration is terminated (Default: 1E-10).
num. iter	The number of interation step (Default: 100).
viz	If viz == TRUE, internal reconstructed matrix can be visualized.
figdir	the directory for saving the figure, when viz == TRUE.
verbose	If verbose == TRUE, Error change rate is generated in console windos.

Value

W : A matrix which has N-rows and J-columns ($J < N, M_k$). H : A list which has multiple elements containing M_k -rows and J-columns matrix ($J < N, M_k$). RecError : The reconstruction error between data matrix and reconstructed matrix from W and H. TrainRecError : The reconstruction error calculated by training set (observed values specified by M). TestRecError : The reconstruction error calculated by test set (missing values specified by M). RelChange : The relative change of the error.

Author(s)

Koki Tsuyuzaki

References

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Examples

```
matdata <- toyModel(model = "dsiNMF_Easy")
out <- dsiNMF(matdata, J=2, num.iter=2)
```

dSVD*Discretized Gradient Descent Singular Value Decomposition (dSVD)*

Description

The input data is assumed to be a matrix. dSVD decompose the matrix to two low-dimensional factor matices.

Usage

```
dSVD(X, M=NULL, pseudocount=.Machine$double.eps,
      initU=NULL, initV=NULL, fixU=FALSE, fixV=FALSE,
      Ter_U=1e-10, L1_U=1e-10, L2_U=1e-10, eta=1e+10, J = 3,
      thr = 1e-10, num.iter = 100,
      viz = FALSE, figdir = NULL, verbose = FALSE)
```

Arguments

X	The input matrix which has N-rows and M-columns.
M	The mask matrix which has N-rows and M-columns. If the input matrix has missing values, specify the element as 0 (otherwise 1).
pseudocount	The pseudo count to avoid zero division, when the element is zero (Default: Machine Epsilon).
initU	The initial values of factor matrix U, which has N-rows and J-columns (Default: NULL).
initV	The initial values of factor matrix V, which has M-rows and J-columns (Default: NULL).
fixU	Whether the factor matrix U is updated in each iteration step (Default: FALSE).
fixV	Whether the factor matrix V is updated in each iteration step (Default: FALSE).
Ter_U	Paramter for terary (0,1,2) regularitartion (Default: 1e-10).
L1_U	Paramter for L1 regularization (Default: 1e-10). This also works as small positive constant to prevent division by zero, so should be set as 0.
L2_U	Paramter for L2 regularization (Default: 1e-10).
eta	Stepsize of gradient descent algorithm (Default: 1e+10).
J	The number of low-dimension (J < N, M, Default: 3)
thr	When error change rate is lower than thr, the iteration is terminated (Default: 1E-10).
num.iter	The number of interation step (Default: 100).
viz	If viz == TRUE, internal reconstructed matrix can be visualized.
figdir	The directory for saving the figure, when viz == TRUE.
verbose	If verbose == TRUE, Error change rate is generated in console window.

Value

U : A matrix which has N-rows and J-columns ($J < N, M$). V : A matrix which has M-rows and J-columns ($J < N, M$). RecError : The reconstruction error between data tensor and reconstructed tensor from U and V. TrainRecError : The reconstruction error calculated by training set (observed values specified by M). TestRecError : The reconstruction error calculated by test set (missing values specified by M). RelChange : The relative change of the error.

Author(s)

Koki Tsuyuzaki

Examples

```
# Test data
matdata <- toyModel(model = "dSVD")

# Simple usage
out <- dSVD(matdata, J=2, num.iter=2)
```

toyModel

Toy model data for using dNMF, dSVD, dsiNMF, djNMF, dPLS, dNTF, and dNTD

Description

The data is used for confirming the algorithm are properly working.

Usage

```
toyModel(model = "dNMF", seeds=123)
```

Arguments

model	Single character string is specified. "dNMF", "dSVD", "dsiNMF_Easy", "dsiNMF_Hard", "dPLS_Easy", "dPLS_Hard", "dNTF", and "dNTD" are available (Default: "dNMF").
seeds	Random number for setting set.seeds in the function (Default: 123).

Value

If model is specified as "dNMF" or "dSVD" a matrix is generated. If model is specified as "dsiNMF_Easy", "dsiNMF_Hard", "dPLS_Easy", or "dPLS_Hard" three matrices are generated. Otherwise, a tensor is generated.

Author(s)

Koki Tsuyuzaki

See Also[dNMF,dSVD](#)**Examples**

```
matdata <- toyModel(model = "dNMF", seeds=123)
```

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